# Intracellular infections in the pathogenesis of vascular diseases; *in vitro* studies

**Bouwman, Johannes Jacobus Maria** 

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Bouwman, Johannes Jacobus Maria © 2009

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# Intracellular infections in the pathogenesis of vascular diseases; *in vitro* studies

Intracellulaire infecties en het ontstaan van vaatziekten; in vitro studies (met een samenvatting in het Nederlands)

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door

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Kordia

Bijna alles wat je doet is onbelangrijk, maar het is erg belangrijk dat je het doet.

(Gandhi)

Voor Annelies, Monique en Rianne Aan mijn lieve ouders en schoonouders

## Manuscripts based on the studies presented in this thesis

#### **Chapter 2**

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#### **Chapter 3**

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#### **Chapter 4**

Bouwman JJM, Visseren FLJ, Bouter KP, Diepersloot RJA.

Azithromycin inhibits interleukin-6 but not fibrinogen production in hepatocytes infected with Cytomegalovirus and *Chlamydia pneumoniae* 

J Lab Clin Med 2004; 144: 18-26.

#### Chapter 5

Bouwman JJ, Visseren FL, Bevers LM, van der Vlist WE, Bouter KP, Diepersloot RJ.

Azithromycin reduces Chlamydia pneumoniae-induced attenuation of eNOS and cGMP production by endothelial cells.

Eur J Clin Invest 2005; 35(9): 573-582.

#### Chapter 6

Kartikasari AE, Georgiou NA, de Geest M, van Kats-Renaud JH, Bouwman JJ, van Asbeck BS, Marx JJ, Visseren FL.

Iron enhances endothelial cell activation in response to Cytomegalovirus or Chlamydia pneumoniae infection.

Eur J Clin Invest 2006; 36(10):743-52.

#### **Chapter 7**

Bouwman JJM, Visseren FLJ, Bouter KP, Diepersloot RJA.

Infection induced inflammatory response of adipocytes in vitro.

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#### **Chapter 8**

Bouwman JJM, Visseren FLJ, Diepersloot RJA.

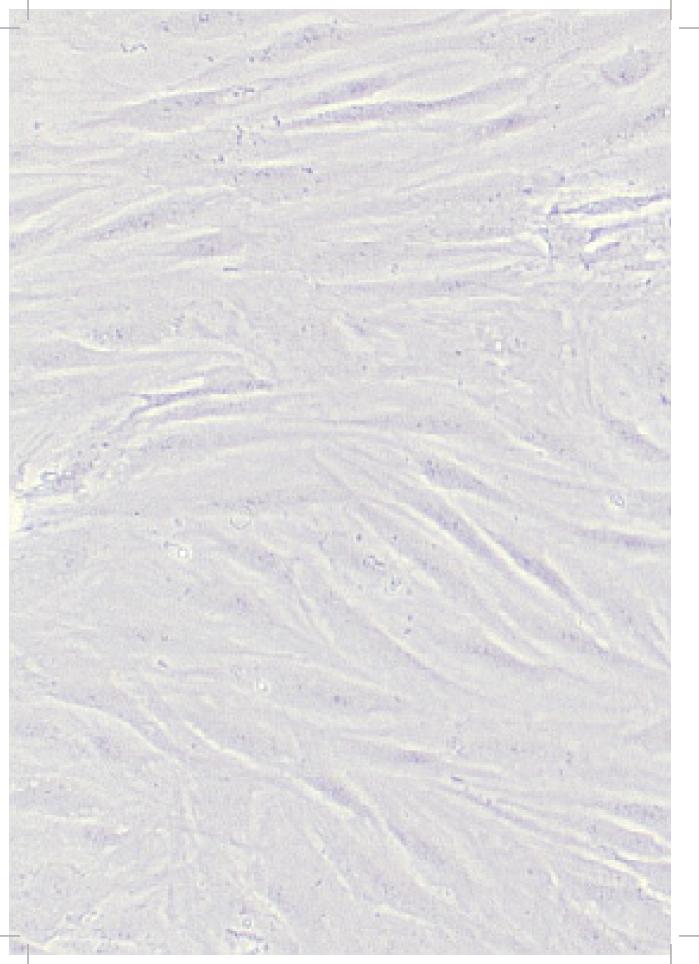
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### Inflammation

Inflammation is a process that develops in response to infection or injury. The inflammatory response usually protects tissues from infection or injury but may become harmful if misdirected. Inflammation is involved in destroying microorganisms and removing debris of damaged cells. Infiltration of leucocytes (acute inflammation) in the diseased tissue and production and release of inflammatory mediators by cells of key organs of the immune system are the first signs of inflammation. In chronic inflammation, predominantly mononuclear cells invade the infected or injured tissue. When an infectious agent has been able to penetrate the system the innate immune response is the next line of defense. Among neutrophils, dendritic cells and NK cells, macrophages play the most important role in this mechanism<sup>1,2</sup>. Should a pathogen still be able to escape the innate response, T cells and B cells activate the adaptive immune response. Adaptive immunity recognizes specific molecular structures (antigens) presented by major histocompatibility complex (MHC) classes II on antigen presenting cells such as macrophages. Once T cells recognize a presented antigen, an adaptive immune response against this specific antigen is initiated<sup>3</sup>. These responses include direct killing of antigen bearing cells by cytotoxic T lymphocytes, stimulation of B cells to produce antibodies against the antigen, and induction of an enhanced innate response in the area where the antigen is present<sup>4</sup>. CD8 expressing T cells are cytotoxic and kill the antigen bearing cells infected by virus or other intracellular organisms. Upon activation, CD4 positive cells may differentiate into T-helper (Th) cells or regulatory T-cells<sup>5</sup>. Th1 cells primarily secrete IFN-γ, IL-2 and TNF-α, which promote cellular immunity against intracellular bacteria and viruses. Th1 cytokines are generally referred to as pro-inflammatory and Th2 cytokines as anti-inflammatory. Common human diseases such as allergy, autoimmunity, chronic infections and sepsis are characterized by a deregulation of the pro- versus anti-inflammatory and Th1 versus Th2 cytokine balance<sup>6</sup>. The failure of these defensive measures suggests that the monocytic and/or T lymphocyte responses are either inadequate or that these inflammatory responses have been altered or subverted from performing their usual protective and reparative functions.

## Monocytes and macrophages

Monocytes play a prominent role in inflammation, coagulation and atherosclerosis by their ability to produce tissue factor (TF) and cytokines<sup>7-10</sup>. The production of cytokines by monocytes may accelerate the chronic process of atherosclerosis and may cause coronary syndromes by eliciting plaque instability. In atherosclerosis, monocytes infiltrate the endothelial layer of the vessel wall and inflammatory mediators enhance uptake of modified lipoprotein particles and formation of lipid-filled macrophages<sup>11,12</sup>. Macrophages can not only be found in endothelial layers. For example, monocytes can migrate into various tissues and organs where they undergo a series of changes and differentiate into mature macrophages. Basically, two types of macrophages can be distinguished: resident (normal) and exudate (inflammatory) macrophages. Each type of resident macrophage, determined by its location, has a specific name<sup>13</sup> e.g. alveolar macrophages in the lung, "Kupffer cells" (liver)<sup>14</sup>, Langerhans cells (skin), osteoclasts (bone) or ATM (adipose tissue macrophages). Macrophages are considered to be a heterogeneous cell population differing in their origin, development stage (differentiation) and local adaptation. Therefore macrophages, in their function and purpose, are responsible for numerous metabolic, immunological, and inflammatory processes in physiological and pathological conditions.

## Atherosclerosis: an inflammatory disease

Monocytes adhere to the endothelium and can infiltrate the sub endothelial space. Eventually leading to foam cell formation and atherosclerotic lesions<sup>15,16</sup>. Atherosclerotic plaques contain inflammatory cells, cholesterol and cellular debris. The migration and infiltration of chronic inflammatory cells, i.e. monocytes and T-cells, reflect an "active" inflammatory atherosclerotic disease<sup>17-19</sup>. In turn, chronically inflamed tissues may perpetuate and even amplify the inflammatory state by releasing vasoactive and proinflammatory substances into the circulation. Inflammatory cell infiltrates are found only within or overlying atherosclerotic plaques. Monocytes are not found in normal intima that has no evidence of atherosclerosis<sup>20,21</sup>. Atherosclerosis, depending on its severity and the location of the artery it affects, may result in vessel obstruction and can cause a rupture of inflamed plaques<sup>22</sup>. Interleukin-6 (IL-6), a cytokine that can be measured in the circulation, initiates and accelerates the process of atherosclerosis. IL-6 has been identified as a marker of inflammation in coronary atherosclerotic plaques. Serum levels of IL-6 increase in response to acute

myocardial infarction, unstable angina, percutaneous coronary intervention and in late restenosis. IL-6 stimulates platelet aggregation and the expression of tissue factor, macrophage LDL receptors, CRP, and fibrinogen. IL-6 also regulates the expression of other inflammatory cytokines, such as interleukin-1 (IL-1) and tumor necrosis factor-  $\alpha$  (TNF- $\alpha$ )<sup>23</sup>. It is conceivable that inflammatory cytokines play a causal role in the development of CHD. On the other hand it can be argued that these cytokines simply reflect an underlying disease process. C-reactive protein (CRP), one of many human acute-phase reactants, is produced in the liver in response to IL-6, IL-1ß and TNF-α. It activates the classic complement cascade, mediates phagocytosis, regulates inflammation, and is a nonspecific but sensitive marker of infection and tissue inflammation<sup>24</sup>. Because of their interrelationship, CRP could simply be a surrogate marker for IL-6 or some other factor<sup>25</sup>. Inflammatory cytokines directly stimulate the production of adhesion molecules. The expression of pro-inflammatory cytokines is decisive for the progression of atherosclerosis. Many pro-inflammatory cytokines, such as IL-12 and IFN-y, and various chemokines exert their pro-atherosclerotic effects during all stages of lesion development<sup>26,27</sup>. Continued release of cytokines in the lesion by macrophages and T-cells not only perpetuates inflammation within the lesion but also modulates smooth muscle cell activity<sup>28</sup>. For example, cytokines such as TNF-α and IFN-y can promote the uptake of modified lipoproteins that leads to smooth muscle cell derived foam cells in vitro and IL-10 inhibits smooth muscle cell accumulation in several animal models<sup>29-31</sup>. Also, many cytokines can influence tissue factor expression on monocytes and on endothelial cells in vitro. Tissue factor plays a central role in the initiation of coagulation activation. TNF-α, IL-1, IL-2, IL-6, and IL-12 are cytokines that have been found capable of activating the coagulation system in vivo, whereas IL-10 can act as an anticoagulant<sup>32</sup>. The presence of T-lymphocytes within an atherosclerotic plaque indicates that the immune system has been activated 18, 33-35. Now that inflammation is considered the underlying pathophysiological cause and key process in atherosclerosis, the driving force behind chronic low-grade inflammation becomes a matter of question. The importance of macrophages normal function in wound healing has been noted before<sup>7,36</sup>. The chronic inflammatory process involving the arterial endothelium that ultimately results in the complications of atherosclerosis, may be caused by a response to the oxidative components of modified low-density lipoprotein (LDL), or to chronic infection, free radicals, classical risk factors such as diabetes, smoking, hypercholesterolemia and hypertension or to other factors 15,16.

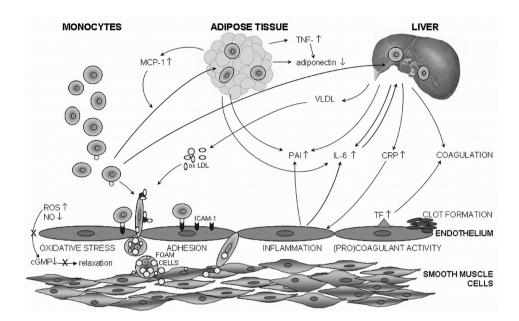


Figure 1. Mechansims involved in vascular injury and inflammation.

In atherosclerosis, monocytes infiltrate the endothelial layer of the vessel wall. Monocytes uptake low density lipoproteins (LDL) and differentiate into macrophages. Macrophages release cytokines that direct migration of monocytes from the blood to the sites of inflammation. The modified LDL particles are taken up by scavenger receptors of macrophages, which evolve into foam cells. Excess LDL infiltrates the artery and is retained in the intima. Oxidative and enzymatic modifications lead to the release of inflammatory lipids that induce endothelial cells to express leukocyte adhesion molecules. Endothelial activation leads to the release of oxygen and nitrogen radicals, increased expression of adhesion molecules and inflammatory cytokines. Ultimately, this results in inflammation and tissue damage. The inflammatory response of adipocytes to infection may stimulate the production of inflammatory cytokines (TNF-alpha, IL-6). Visceral fat, in particular, contributes to endothelial dysfunction through the direct effect of adipokines, mainly adiponectin and TNF-α, which are secreted by fat tissue after macrophage recruitment (through monocyte chemoattractant protein-1, MCP-1). Indirect effects of TNF-α and IL-6 might influence inflammation (CRP) and endothelial function. By releasing hormones and cytokines into the portal circulation, abdominal adipose tissue is able to directly influence liver metabolism, potentially leading to accelerated inflammatory response, e.g. C-Reactive Protein (CRP) production, IL-6 production and leading to changes in hemostasis by the unbalanced production of coagulant factors and proteins involved in fibrinolysis (PAI-1).

## Infection

The immune system is vital in defending the body against bacteria and viruses. Improper functioning can lead to a variety of diseases, ranging from immune deficiencies to chronic inflammation. In chronic infections, a limited number of cells (in the target organs) are infected. These infected cells may demonstrate a cytopathology effect, synthesize virus macromolecules, and release infectious virus. The spread of infection is limited by host factors such as humoral and cell-mediated immune responses. Viral affinity for specific body tissues (tropism) is determined by cell receptors for virus, cell transcription factors that recognize viral promoters and enhancer sequences, ability of the cell to support virus replication, physical barriers, local temperature, pH and digestive enzymes and bile in the gastrointestinal tract that may inactivate some viruses. Viruses enter the human body via the respiratory system, gastrointestinal tract, skin and genital routes although other routes can be used. The final outcome of infection may be determined by the dose and location of the virus as well as its infectivity and virulence. Most virus types spread among cells extracellular, but some may also spread intracellular. Establishment of local infection may lead to localized disease and localized shedding of virus. The most common route of systemic spread from the portal of entry is the circulation, which the virus reaches via the lymphatic system. Virus may enter the target organs from the capillaries by multiplying in endothelial cells or fixed macrophages, diffusing through gaps, and being carried in a migrating leukocyte. Dissemination via peripheral nerves usually occurs with rabies virus and sometimes with herpes virus and poliovirus infections. Depending on the balance between virus and host defenses, virus multiplication in the target organ may be sufficient to cause disease and death. Although the respiratory tract, alimentary tract, urogenital tract and blood are the most frequent sites of shedding, various viruses may be shed at virtually every site.

## Infection causing 'non-infectious diseases'

## **Chronic infection and oncogenesis**

Many chronic diseases are caused by pathogens or by the chronic inflammatory response of our own bodies to pathogens. Infections provoke an immune response from the body. It has recently been recognized that certain chronic infectious agents may contribute to carcinogenesis by inducing a state of persistent inflammation. Certain

persistent viral or bacterial infections can promote a chain of events by interfering with the regulation of cell proliferation or cell death or by inducing DNA damage. A feature common to all oncogenic infectious agents is their long term persistence in infected individuals. The infectious agents that have been recognized as human carcinogens include the viruses Epstein-Barr virus (EBV), Kaposi sarcoma herpes virus (KSHV), Human papillomavirus (HPV), Human T-cell lymphotropic virus (HTLV-I), Hepatitis B virus (HBV), Hepatitis C virus (HCV) and the Helicobacter pylori (HP) bacterium. EBV is associated with Non-Hodgkin's lymphoma in the lymph nodes<sup>37,38</sup> KSHV with Kaposi sarcomas<sup>39,40</sup> HPV with cervical cancer<sup>41,42</sup> and HBV and HCV with hepatocellular carcinoma<sup>43-45</sup>. The H. pylori bacterium is related to gastric cancer<sup>46</sup>.

#### **Chronic infection and atherosclerosis**

The concept that chronic infection may contribute to the pathogenesis of atherosclerosis was already proposed in the early 1980's. With respect to the importance of inflammation in atherosclerosis, many have hypothesized that infectious agents may be the cause of chronic vascular inflammation. Although it remains not fully elucidated whether pathogens are etiological factors in atherosclerosis, pathogens can aggravate the inflammatory process in atherosclerosis<sup>47-49</sup>. The major organisms that have been studied with respect to vascular diseases are Chlamydia pneumoniae (Cp), Cytomegalovirus (CMV), and Helicobacter pylori, but enterovirus, hepatitis A virus and herpes simplex virus type 1 and type 2 have also been implicated. An association is described between cytomegalovirus or Cp and an increased risk of vascular events50. These micro-organisms are able to infect endothelial cells and smooth muscle cells in vivo and in vitro, evoking pathophysiological reactions of these cells which may lead to atherosclerosis, arterial thrombosis and plaque rupture<sup>50-54</sup>. CMV, *Cp* and many other bacteria have been detected in human atherosclerotic lesions<sup>51,55</sup>. *Cp* in particular has been the focus of attention as a possible contributor to coronary atherosclerosis<sup>56</sup>. In an animal model, repeated infections with Cp resulted in endothelial dysfunction via impaired NO availability<sup>57</sup>. Elevated titers of antibodies against Chlamydia were found in patients with CAD<sup>47</sup> and it was speculated that this microbe causes atherosclerosis. Although *Cp*-infection did not cause atherosclerosis in animals it may stimulate disease progression and plaque activation<sup>58,59</sup>. This could be due either to a direct action in plagues or to remote signaling by inflammatory mediators<sup>60</sup>. Molecular mimicry between Cp-antigens and human molecules may contribute to the activation of inflammation<sup>61</sup>. However, several recent secondary-prevention trials, failed to prevent

acute coronary syndromes by administering antibiotics targeting Cp, suggesting that Cp-infection is not a predominant cause of these syndromes<sup>62-64</sup>. Herpes family viruses may also contribute to CAD. Infection with CMV and other herpes viruses, such as HSV, has been implicated in some, but not all studies<sup>65-67</sup>. Serum anti-CMV antibody levels are significantly elevated in patients with coronary disease when compared to control subjects<sup>65,68</sup> and in those with carotid intimal thickening<sup>69</sup>. In addition, CMV antigens, nucleic acid sequences, and DNA have been detected in smooth muscle cells of carotid artery plaques obtained from patients undergoing endarterectomy<sup>70</sup>. CMV is found in arterial lesions, can modulate immune-cell as well as vascular-cell activity, and increases experimental atherosclerosis<sup>71</sup>. Furthermore, in vitro infection of macrophages with CMV increases secretion of IL-1, TNF-α and M-CSF. Clinical data imply an important role for CMV intransplantation-related arteriosclerosis causing graft rejection<sup>72</sup>. There is an association between antibodies to CMV and elevated levels of markers of inflammation, including C-reactive protein (CRP) and interleukin (IL)-6. The combination of CMV-seropositivity and elevated serum CRP is a strong, independent predictor of mortality<sup>56</sup>; the data are similar for CMV-seropositivity and elevated serum IL-6 73. These findings suggest that CMV elicits a subclinical inflammatory response in certain individuals who are therefore susceptible to the development of atherosclerosis; those without an inflammatory response are resistant<sup>74</sup>. In contrast to CMV and Cp, influenza virus represents a micro-organism which causes more acute infections with severe (systemic) complications. Vaccination against influenza has been evaluated for a potential benefit in preventing cardiovascular disease<sup>75-77</sup>. Several studies have found a beneficial effect of influenza vaccination on cardiovascular events. While these studies are of interest and do document a beneficial effect of influenza vaccination in the elderly, they do not establish a causative relationship between influenza infection and the pathogenesis of atherosclerosis. It is possible that the observed beneficial effect of vaccination on the incidence of cardiac events relates to the prevention of clinical influenza, with consequent reduction in such complications as dehydration, hypoxemia, and demand ischemia. In a study of mice infected with influenza A, HDL lost its anti-inflammatory properties during the acute-phase response. *In vitro* analyses found that the ability of HDL to inhibit LDL oxidation and LDL-induced monocyte chemotactic activity in human arterial cell co-cultures decreased with time after infection. These changes were not associated with a direct effect of the virus on HDL, but were thought to result from a systemic response<sup>78</sup>. Regarding the possible role of infectious pathogens in atherosclerosis, it has been suggested that antimicrobial

therapy might reduce coronary risk. Results of clinical trials investigating antichlamydial antibiotics as an addition to standard therapy in patients with coronary artery disease have been inconsistent<sup>79-81</sup>. Although quinolone treatment seems to reduce adverse cardiac events in patients with acute coronary syndromes, the effect was independent of H. pylori or *Cp*-seropositivity<sup>82</sup>. The results of a meta-analysis of large randomized trials do not indicate that antibiotic therapy against *Cp* is beneficial for reducing cardiovascular coronary events<sup>83</sup>.

## Cellular dysfunction caused by infection

Chronic infection may contribute to the pathogenesis of cellular dysfunction by initiating and maintaining cellular injury leading to a local or systemic inflammatory response. In persistent viral infections, the host's immune system is challenged by the constant exposure to microbial antigens with no evidence of protein synthesis but still interference with cellular functions. Low-grade chronic inflammation that is persistent, as can be seen in obesity, diabetes, and the metabolic syndrome could result in acceleration of the cellular and humoral immunologic response<sup>84</sup>. An acute inflammatory response in normal subjects, as measured by elevated plasma concentrations of C-reactive protein, a marker of chronic inflammation, is an independent predictor of endothelial dysfunction. Even before clinical symptoms become evident, a complex of inflammatory and metabolic changes may have been initiated. In early stages of disease (pro-)inflammatory cytokines, adhesion molecules, oxygen radicals and impaired NO availability<sup>57</sup> can be detected as markers and predictors of cellular dysfunction. In other cases, the effect may simply be that of enhancing the pre-existing chronic inflammatory response of the body to standard risk factors such as hypercholesterolemia. Even though the infectious agent may not directly infect the arterial wall, it may perform its critical role from afar.

### **Endothelial dysfunction**

Atherosclerosis is a major cause of morbidity and mortality worldwide. Inflammatory processes and endothelial dysfunction appear to play important roles in the etiology of cardiovascular diseases. Nevertheless, the mechanisms underlying these conditions are not fully understood, yet. A large number of risk factors for atherosclerosis have been proposed and specifically hypercholesterolemia, systemic hypertension, smoking and diabetes have been associated with endothelial dysfunction<sup>85-87</sup>.

Endothelial dysfunction plays a pivotal role in atherogenesis and development of clinical evident vascular diseases. Under normal conditions the endothelium regulates vascular tone and has anti-atherogenic and anti-thrombotic properties. Dysfunctional endothelium may initiate and promote atherogenesis due to expression of adhesion molecules, tissue factor expression, platelet activation and release of pro-inflammatory cytokines. Enhanced expression of adhesion molecules leads to enhanced monocyte adhesion and subsequent migration across the endothelium into the vessel wall and accumulation of oxidized low density lipoproteins (LDL) leads to foam cell formation88. Reduced bioavailability of nitric oxide (NO) is the primary cause and/or result of endothelial dysfunction. Elevated plasma cholesterol concentrations and oxidized LDL in combination with high levels of radicals may inactivate NO and impair NOsynthesis by affecting the function of NO synthase<sup>89-92</sup>. A number of risk factors for atherogenesis, including infectious agents, have been shown to exert their influence via inflammatory actions93. Endothelial cell activation leads to expression of proinflammatory cytokines like IL-6, TNF-α and CRP. These inflammatory cytokines have local effects on neighboring endothelial cells and have systemic effects on the vessel wall and on insulin resistance. Infection and the resulting inflammatory response can produce endothelial dysfunction. In the last decade a role for infections in (chronic) inflammation in lesions and the pathogenesis of atherosclerosis has been proposed frequently<sup>50,53,57,94</sup>. Detection of microbial sequences in the vessel wall has proven the direct involvement of microorganisms in atherogenesis<sup>51,55</sup>. Direct infection of endothelial cells may cause acute and chronic activation of the endothelium and loss of its anti-atherogenic protection. In addition, there is now a growing recognition of non-traditional risk factors, like infections as potential modulators of the endothelial phenotype in obesity, including fat tissue production of pro-atherogenic adipokines, oxidative stress, and chronic inflammation<sup>95-97</sup>.

## **Macrophage dysfunction**

Macrophages are involved in all stages of the immune response and constitute an important cellular component of the immune responses against viruses. They serve as antigen presenting cells and also secrete inflammatory mediators to activate innate and adaptive immune cells. Resident tissue macrophages are relatively quiescent immunologically, but in response to invasive stimuli and inflammation they become activated and their numbers may increase dramatically. Mononuclear phagocytes (monocytes and macrophages) are central to inflammation, as they produce many

components that regulate the mediators of the inflammatory response. They are also actively phagocytic and are involved in microbial killing. Analysis of cytokine production during immune responses suggests that different macrophage populations participate at various stages, or that the changing conditions within the lesion differentially affect the functions of distinct macrophage populations. Recently the concept of macrophage polarization was described, indicating that macrophages are functionally polarized in response to microorganisms and host mediators. Polarized activation of cells of the monocyte-heterogeneous macrophage lineage into M1 and M2 cells reveals that M1 polarization can lead to tissue injury and contribute to pathogenesis. The so-called M2 macrophages play a critical role in the resolution of inflammation by producing anti-inflammatory mediators indicating a range of immune and inflammatory conditions<sup>98-100</sup>. In some conditions ineffective phagocytosis and macrophage dysfunction have been described<sup>101</sup> but activation of macrophages probably is a better reproduction of "macrophage dysfunction". In the pathophysiology of monocyte and macrophage dysfunction chronic inflammatory diseases play an important role. Dysfunctional macrophages feature oxidative stress, as reflected by increased levels of ROS, deregulated cytokine release and enhanced chemotaxis. Macrophages that have been modified by oxidized LDL release a variety of inflammatory substances, cytokines, and growth factors<sup>102</sup>. Among the many molecules that have been implicated are: monocyte chemotactic protein (MCP)-1, intercellular adhesion molecule (ICAM)-1)103, granulocyte-macrophage colony stimulating factors<sup>104</sup>, soluble CD40 ligand, interleukins<sup>105</sup> and TNF-α<sup>106</sup>. During infection with intracellular pathogens, macrophages serve a dual role as host and effector cells. Macrophage activation has to be tightly regulated to achieve effective anti-pathogen responses and at the same time avoid immunopathology. Down regulation of macrophages is mediated by multiple mechanisms, including inhibitory cytokines, induction of tolerance, cytokine antagonists, or inhibition of activating signaling pathways. Defects in these regulatory pathways can lead to autoimmunity and hyperinflammation; on the other hand, there is evidence that pathogens, especially those adapted to intracellular life within macrophages, exploit these endogenous inducers of deactivation<sup>107</sup>. Among numerous micro-organisms *Cp*, CMV and influenza are able to infect monocytes and macrophages. Cp-infection activates human vascular endothelium, smooth muscle cells and macrophages and affects TNF-α and MMPproduction<sup>108</sup>. Growth kinetics of human cytomegalovirus may be altered in monocytederived macrophages<sup>109</sup>. Impaired viral replication indicated by non-lytic growth of

human cytomegalovirus in macrophages indicates a chronic persistent CMV-infection still enabling up regulation of cytokine production contributing to inflammatory processes<sup>110</sup>, <sup>111</sup>. Influenza A virus-infected macrophages are primed for a high TNF- $\alpha$  release<sup>112</sup> and Influenza A infection in mice macrophages leads to increased arterial macrophage traffic and is associated with loss of anti-inflammatory properties of HDL <sup>113</sup>.

## **Hepatocyte dysfunction**

The liver is a key organ in inflammatory processes. Both local and systemic reactions are initiated. Hepatocytes respond by producing "acute phase proteins" like IL-6 and CRP. The intentional purpose of the acute phase reaction is protective aiming to limit tissue injury, infection or inflammation and restoration of homeostasis. Hepatocytes are responsible for a response to inflammation called the acute-phase reaction by production of "acute phase proteins". Positive acute-phase proteins serve different physiological functions for the immune system. These include pro-inflammatory factors like CRP and IL-6, coagulation factors like fibrinogen, prothrombin, factor VIII, von Willebrand factor and plasminogen but also complement factors, ferritin, ceruloplasmin, serum amyloid A and haptoglobin. The negative acute phase reactants like serpins (e.g. PAI-1) and alpha 2-macroglobulin give negative feedback on the inflammatory response. In response to injury, local inflammatory cells (neutrophil granulocytes and macrophages) secrete a number of cytokines into the bloodstream, most notable of which are the interleukins IL-1, IL-6 and IL-8, and TNF-α<sup>114-116</sup>. Under pathological conditions hepatocyte dysfunction can lead to lipid abnormalities and may elicit intensive production of inflammatory mediators and profound changes in the concentrations of proteins participating in coagulation and in fibrinolysis<sup>117</sup>. Infection and inflammation in the liver induce the acute-phase response (APR)118,119. This may lead to alterations in lipid and lipoprotein metabolism, oxidative stress, coaqulation disorders and production of (pro-) inflammatory cytokines Microorganisms may also initiate and maintain chronic low-grade inflammation by infecting liver cells<sup>120</sup>. The most investigated liver infection is hepatitis caused by Hepatitis B or -C viruses (HBV, HCV). One of the most remarkable aspects of HCV infection is that it causes low-grade infection in the liver over many years 121,122. Infection of the liver by CMV or Cp may induce a general inflammatory reaction contributing to accelerated atherogenesis. Infections with CMV and Cp increase fibringen production of hepatocytes in vitro. In addition, infections also increase IL-6 production. In addition to direct vascular wall infection by Cp and CMV, virus-related development of atherosclerosis might also be initiated by

chronic liver infection and subsequent production of inflammatory and procoagulant mediators released in the circulation. This may be another pathophysiological link for the observed relation between infections and the development of atherosclerosis<sup>120</sup>. On the other hand, intrahepatic cytokine production triggers the recruitment of mononuclear cells, which sustain acute and chronic liver damage<sup>123</sup>. A productive infection by *Cp* may take place in Kupffer cells and *Cp* might induce a local proinflammatory activity. *Cp* may therefore, be able to act as antigenic stimulus when localized in the liver. One could speculate that *Cp*-infection, involving cells of the innate immunity such as Kupffer cells, could also trigger pathological immune reactions involving the liver<sup>124</sup>.

## **Adipocyte dysfunction**

Adipose tissue is distributed over several locations in the body: subcutaneous, abdominal and visceral. Abdominal adipose tissue is correlated with cardiovascular diseases, hypertension and diabetes. Adipose tissue exists of adipocytes; lipidfilled cells which are embedded in an environment of collagen fibers. In addition to adipocytes, adipose tissue contains stromal-vascular cells including fibroblast-like cells, leukocytes, macrophages and pre-adipocytes. Adipose tissue thus appears as a complex tissue composed of different cell subsets that could vary according to the nature and the location of fat pads, or to the physiological or pathological status<sup>125</sup>. Obesity induces adipose tissue macrophage (ATM) infiltration in white adipose tissue in both humans and mice<sup>126,127</sup>. ATM content correlates with measures of adiposity and insulin resistance<sup>128,129</sup>, based on the evidence that obesity is associated with infiltration<sup>127</sup> and activation<sup>130</sup> of macrophages in adipose tissue. Obese patients also have an increased risk for the development of atherosclerosis and diabetes mellitus type 2 and myocardial infarction<sup>131</sup>. Although the exact pathophysiological mechanisms are not yet clear, insulin resistance, closely related with obesity, and inflammation are key components<sup>132,133</sup>. Inflammatory and prothrombotic markers are associated with an increased risk for type 2 diabetes and subsequent CVD and obesity and the metabolic syndrome have been proposed to be a state of chronic inflammation, associated with elevated levels of CRP, IL-6, and plasminogen activator inhibitor (PAI)-1<sup>134-136</sup>. Consequently, adipose tissue appears to be important in vascular injury and adipocyte dysfunction can be the cause but also the consequence of insulin resistance. By producing (adiponectin) and inflammatory cytokines (TNF-α, IL-6), adipose tissue is likely to play an important role in the development of diabetes

and atherosclerosis 134,135,137. Causes for adipocyte dysfunction are not well established but include abdominal obesity and elevated free fatty acid plasma concentration 138. By releasing hormones and cytokines into the portal circulation, abdominal adipose tissue is able to directly influence liver metabolism, potentially leading to accelerated inflammatory response, e.g. C-reactive protein (CRP) production, and leading to changes in hemostasis by the unbalanced production of coagulant factors and proteins involved in fibrinolysis. Several microorganisms, hormones and inflammatory factors have already been related to diseases such as obesity, atherosclerosis and diabetes mellitus type 2. Diabetics have an increased propensity to develop morbidity from infections. In addition, the types of infections observed in patients with diabetes mellitus (DM) also are complex. Neutrophil chemotaxis and adherence to vascular endothelium, phagocytosis, intracellular bactericidal activity, opsonization, and cell-mediated immunity are all depressed in diabetics with hyperglycemia<sup>139</sup>. Vascular disease related to diabetes may impair the local inflammatory response, the bactericidal functions of leukocytes and the absorption of antibiotics. Although obesity has multiple causes, an often overlooked possibility is that of obesity due to an infection. The expression "infectobesity" describes the possibility that obesity may be, at least in part, the result of viral infections<sup>140,141</sup>. Among several other micro-organisms the human Adenovirus subtype 36, but also subtypes 5, 31 and 37 have been associated with adipogenesis and obesity<sup>142,143</sup>. Data from animal models suggest that the role of viral disease in the etiology of human obesity must be considered 144-146. Ad-36 is associated with increased body weight and lower serum lipids in humans. Prospective studies need to indicate if Ad-36 plays a role in the etiology of human obesity<sup>147</sup>. The inflammatory response of adipocytes upon infection may also have paracrine effects by affecting the function of neighboring adipocytes and may attract macrophages into adipose tissue, increasing the capacity for the production of inflammatory mediators

## **Objectives of this thesis**

In this thesis we aimed to investigate the role for microorganisms in inducing and maintaining cellular dysfunction causing pathophysiological processes potentially involved in the development of atherosclerosis and diabetes mellitus.

#### **Outline of this thesis**

In **chapter 2** we investigate the ability of respiratory viruses (influenza A, influenza B, parainfluenza-1, respiratory syncytial virus, adenovirus and CMV) to infect lung fibroblasts and human umbilical vein endothelial cells and the effect of infection on the procoagulant activity of these cells *in vitro*. Production of cytokines by monocytes may accelerate the chronic process of atherosclerosis and may contribute to coronary syndromes by eliciting plaque instability. Whether virus-infected monocytes initiate coagulation and produce (pro-) inflammatory cytokines was investigated in **chapter 3**. In **chapter 4** we describe that human hepatocytes can be infected with the intracellular pathogens *Cp* and CMV. We measure the production of fibrinogen, IL-6, and PAI-1 after infection. Azithromycin will be added to establish whether biosynthesis of these parameters could be influenced by antibiotic treatment.

Nitric oxide (NO) is a key regulator of endothelial function. Under pathological conditions uncoupling of endothelial nitric oxide synthase (eNOS) leads to vessel damage as a result of production of oxygen radicals instead of NO, indicating endothelial dysfunction. In **chapter 5** we measured production of eNOS, cyclic guanosine monophosphate (cGMP) as a surrogate for NO and reactive oxygen species (ROS) by endothelial cells infected with *Cp* or CMV *in vitro*.

The labile forms of iron may play a role in the development of atherosclerotic vascular disease. Iron has been found accumulating in human atheroma and several studies have shown reduced formation of early atherosclerotic lesions by means of iron chelation or iron-deficient diets in experimental studies. In **chapter 6** we mimic a chronic low-grade vascular *Cp*-infection and observe the presumed modulating effects of iron on endothelial response towards *Cp*. Endothelial intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (VCAM-1) and endothelial selectin (E-selectin) expression are measured as an indication of endothelial activation. We also evaluate potential counteracting effects of iron chelation as well as radical scavenging on the effects of iron on endothelial cells.

In the study described in **chapter 7** we investigate the potential role for microorganisms in inducing and maintaining adipocyte dysfunction and investigate whether several viruses are able to infect adipocytes and pre-adipocytes *in vitro* and whether this leads to adipocyte dysfunction as measured by the altered production of the adipokines IL-6, TNF- $\alpha$ , adiponectin and PAI-1. The inflammatory response of adipocytes upon infection may also have paracrine effects by affecting the function of neighboring

adipocytes and may attract macrophages into adipose tissue, increasing the capacity for the production of inflammatory mediators. In **chapter 8** we measured adipokine production in an *in vitro* co-culture model of adipocytes and macrophages. Co-incubation of adipocytes and macrophages may synergistically amplify adipokine expression, presumably leading to an aggravated inflammatory response. Overproduction of adipokines may evoke potent proatherogenic effects in the vasculature. This may provide a pathological link between obesity and its associated cardiovascular complications. Finally, in **chapter 9** the results are summarized and discussed.

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Procoagulant Activity
of Endothelial Cells after
Infection with Respiratory Viruses

## **Summary**

Influenza virus epidemics are associated with excess mortality due to cardiovascular diseases. There are several case reports of excessive coagulation during generalized influenza virus infection. In this study, we demonstrate the ability of respiratory viruses (influenza A, influenza B, parainfluenza-1, respiratory syncytial virus, adenovirus, cytomegalovirus) to infect lung fibroblasts and human umbilical vein endothelial cells in culture. All viral pathogens induced procoagulant activity in infected endothelial cells, as determined in a one-stage clotting assay, by causing an average 55% reduction in the clotting time. When factor VII deficient plasma was used clotting time was not reduced. The induction of procoagulant activity was associated with a 4- to 5-fold increase in the expression of tissue factor, as measured by the generation of factor Xa. Both experiments indicate that the procoagulant activity of endothelial cells in response to infection with respiratory viruses is caused by up regulation of the extrinsic pathway. Although both enveloped viruses and a non-enveloped virus (adenovirus) induced procoagulant activity in endothelial cells by stimulating tissue factor expression, the role of the viral envelope in the assembly of the prothrombinase complex remains uncertain.

We conclude that both enveloped and non-enveloped respiratory viruses are capable of infecting cultured human endothelial cells and causing a shift from anticoagulant to procoagulant activity associated with the induction of tissue factor expression.

## Introduction

Endothelial cells form a functional and dynamic non-thrombogenic surface in the vascular lumen, and damage to these cells may be a starting point for inflammation, thrombosis and eventually atherosclerosis. There is increasing evidence that viral pathogens, in particular members of the herpes virus family (cytomegalovirus (CMV) and herpes simplex virus (HSV)) can damage the endothelium<sup>1-4</sup>. Herpes viruses<sup>5-7</sup> and measles virus<sup>8</sup> induce procoagulant activity in endothelial cells and cause increased adherence of polymorph nuclear leukocytes<sup>9</sup> and monocytes<sup>10</sup> to endothelial cell monolayers. In an overview Hajjar describes the potential mechanisms of viral activation of coagulation by herpes viruses<sup>11</sup>.

Influenza epidemics are associated with excess mortality and morbidity due to cardiovascular diseases<sup>12-15</sup>. Although age and background diseases appeared to be associated risk factors for influenza-related morbidity and mortality, the exact

mechanism by which influenza has this effect is not known. An influenza-like syndrome resembling acute respiratory infection often precedes myocardial infarction 16, 17. Also, symptoms of chronic upper airway infection predicted the risk of coronary disease independent of the known major cardiovascular risk factors<sup>18</sup>. Influenza virus is able to invade myocardial and pericardial tissue<sup>19-23</sup> and enhances platelet aggregation *in vitro* <sup>24-26</sup> and *in vivo*<sup>27</sup>. In a clinical survey, Bogomolov et al. demonstrated a pronounced acceleration of coagulation in elderly patients with an influenza virus infection<sup>27</sup>. During convalescence, these patients had low concentrations of antithrombin III. A pregnant female is reported to have generalized influenza infection resulting in disseminated intravascular coagulation (DIC)<sup>28</sup>. Multiple thrombosis was observed in the microcirculation. In an animal model, the most pronounced changes observed after infection with influenza were found in the vessels of the lung and brain microcirculation. These changes included alteration of the luminal surface profile and damage to the super membrane layer<sup>26</sup>. In some studies human endothelial cells were infected with influenza virus in vitro<sup>29-32</sup>. Infected cells produced interferon- $\alpha$  and were thus able to up regulate the immune system<sup>29</sup>. Moreover, leukocytes adhered to human endothelial cells infected with influenza virus<sup>31</sup>. In order to investigate the thrombogenic effects of viral infection of endothelial cells and the potential role of virus infection in initiating atherothrombosis, we investigated the ability of respiratory viruses, including influenza viruses, to infect human endothelial cells and cells of other cell lines in vitro and the response of endothelial cells, in terms of procoagulant activity, to infection.

# **Methods**

# **Cell culture**

Endothelial cells. Human umbilical vein endothelial cells were harvested and cultured as described by Jaffe et al<sup>33</sup> with minor modifications. Briefly, the umbilical veins were canulated at each end and washed through with phosphate buffered saline (PBS). A solution of 0.05% trypsin-0.02% EDTA (Trypsin-EDTA, Life Technologies Inc., Paisley, Scotland) was introduced into the vessel lumen and left at 37 °C for 20 minutes. The cells were collected by flushing the vein with sterile PBS and centrifuging the suspension at 1000 rpm for 10 minutes. The pellet was resuspended in endothelial cell growth medium (EGM, Clonetics, San Diego, USA), supplemented with 2% fetal bovine serum, 10 mg/L epidermal growth factor, 12 mg/L bovine brain extract, 1 mg/L hydrocortisone, 50 mg/L gentamycin sulphate and 50 mg/L amphotericin-B. Cells were

cultured in tissue culture flasks (Nunc, Roskilde, Denmark) in a humidified incubator at 37 °C with 5% CO<sub>2</sub>. The growing surfaces of the tissue culture flasks were coated with fibronectin (a generous gift from dr. J.A. van Mourik, CLB, Amsterdam, the Netherlands). For all experiments cultures of passage 3 were used. For procoagulant studies cells were cultured in coagulometer cups (Amelung, Lemgo, Germany) pretreated with fibronectin or for immunocytochemical studies in 24-well tissue culture plates (Nunc, Roskilde, Denmark). The purity of the cultures was verified by morphological analysis as well as by immunostaining (Dako, Glostrup, Denmark) for human Von Willebrand factor antigen (> 97% staining).

Fibroblasts. Human embryonic lung fibroblasts were cultured in Eagle Minimal Essential Medium (EMEM) (Life Technologies Ltd., Paisley, Scotland) supplemented with 10% fetal calf serum (Life Technologies Ltd., Paisley, Scotland), 4 mg/L amphotericin-B (Bristol-Myers-Squibb, Epernon, France), 10 mg/L gentamycin (Schering-Plough, Heist op den Berg, Belgium), 10 mg/L vancomycin (Eli Lilly, Indianapolis, USA), 2 mM L-glutamine (Life Technologies Ltd., Paisley, Scotland) and 1% non-essential amino acids (Life Technologies Ltd., Paisley, Scotland) at 37 °C up to passage 4 in 80-cm² tissue culture flasks (Nunc, Roskilde, Denmark) and stored in liquid nitrogen until further use. Cell passages 5 to 20 were used for experiments.

*Kidney cells*. Rhesus monkey kidney cells (LLC-MK<sub>2</sub>: ICN Biomedical Inc., Costa Mesa, USA) were cultured in tissue culture flasks and maintained in EMEM (described above). Cells were stored in liquid nitrogen.

#### Virus preparations

The influenza A clinical isolates H1N1 (A/Singapore/6/86-like), H3N2 (A/Beijing/32/92-like), Influenza B (B/Panama/45/90) and Parainfluenza type 1 were used. In the experiments with factor deficient plasma, the influenza strains H1N1 (A/Taiwan/1/81), H3N2 (A/Beijing/32/92-like) and B/Netherlands/580/89 were used. Virus strains were identified by immunofluorescence staining with monoclonal antibodies against influenza A and B (Imagen K6105), parainfluenza (Imagen K6103), adenovirus type 2 (Imagen K6100), respiratory syncytial virus (RSV; Imagen K6102), and cytomegalovirus (CMV; 12-003 CMV IEA clone E13). The antibody against CMV was from Argene, Varilhes, France and the other antibodies were from Dako, Glostrup, Denmark. Parainfluenza

and influenza strains were propagated in a monkey kidney cell line (LLC-MK,) with M199 medium (Life Technologies Ltd., Paisley, Scotland) free of serum and supplemented with 0.25% trypsin, 0.2% glucose, 0.1% bovine serum albumin (Behring, Marburg, Germany) and 25 mM Hepes. Clinical isolates of adenovirus type 2 and respiratory syncytial virus (RSV) were propagated in a HEP-2 cell line with EMEM (Life Technologies Ltd., Paisley, Scotland) supplemented with 4 mg/L amphotericin-B, 10 mg/L gentamycin, 10 mg/L vancomycin, 2 mM L-glutamine, 1% non-essential amino acids, 2% fetal calf serum and 25 mM Hepes. Cytomegalovirus was isolated from a newborn infant with a congenital infection and was propagated in human embryonic lung cells in order to get a pool of CMV. After more than 80% of all cells showed changes in morphology due to a virus infection e.g. cytopathologic effect (CPE), as determined by light microscopy evaluation, all virus suspensions were frozen at -90 °C after addition of GLY-medium (1:1 dilution). To determine the 50% tissue culture infecting dose (TCID<sub>50</sub>) virus stocks were titrated against human embryonic lung cells (CMV) and HEP-2 cells (adenovirus and RSV) according to the method of Reed and Muench <sup>34</sup>. For the influenza strains H1N1, H3N2, influenza B and parainfluenza, hemagglutinin (HA) titers were determined by titration. To do this, turkey erythrocytes (0.5%) obtained from the National Institute for Public Health (RIVM, Bilthoven, the Netherlands) were suspended in McIlvaine buffer with 0.2% bovine albumin and 0.01% sodiumazide. Twenty-five microlitres of erythrocyte suspension and 25 µL virus suspension in two-fold dilutions were added to microtiter plates with V-shaped wells (Nunc, Denmark), shaken for 1 minute and incubated for 1 hour at 4 °C. The highest dilution which induced agglutination was considered the HA titer. Influenza growth medium was used as negative control and did not have a detectable HA titer. As a positive control, egg suspensions infected with a known HA titer were used. These suspensions were obtained from the National Institute for Public Health (RIVM, Bilthoven, the Netherlands).

# Viral inoculation of cell monolayers

Endothelial cell monolayers grown to confluence were used for infection experiments. The growth medium was removed from the monolayers and maintenance medium (EMEM, 2% fetal calf serum, 25mM Hepes) was added as described previously. For the procoagulant assay endothelial cell monolayers were incubated with 200  $\mu$ L virus suspension for 3 and 24 hours. For comparison of infectivity in different cell lines, 200  $\mu$ L virus suspension was added to cell monolayers grown in 24 well tissue culture plates, centrifuged at 800g for 60 minutes and incubated for 24 hours. For quantifi-

cation of infection, cells were fixed for 30 minutes in absolute methanol/acetone, air dried, and stained with fluorescein isothiocyanate (FITC)-labeled monoclonal antibodies for 30 minutes at 37 °C. Infected monolayers from different cell lines were examined by fluorescence microscopy. Each virus-infected cell showing apple green fluorescence was scored as one fluorescent unit. Fluorescent units were counted in 10 random fields at 200x magnification. The number of infected cells was expressed as percentage of the infection of the control cell line (the standard cell line was assigned to an infection-% of 100%). The actual infection percentage of the standard cell line was approximately 20%. Infection with all viral strains did not influence cell viability or integrity, as judged by trypan blue exclusion after the 24-hours incubation.

# **Cell procoagulant assay**

Procoagulant activity was measured on the surface of intact endothelial cell monolayers in coagulometer cups (Amelung, Germany), using a one-stage clotting assay. After different incubation times (3 and 24 hours) culture medium was removed and cells were washed with PBS. Monolayers were incubated with 100 µL Michaelis buffer (sodium-acetate 28.5 mM, sodium-barbital 28.5 mM, NaCl 50 mM, CaCl 33 mM, pH 7.35) at 37 °C for 60 seconds after which 100 μL human pooled citrated plasma or factor VII or XI deficient plasma (Biopool, Burlington, Canada) was added. The time necessary for fibrin to form was measured on the coagulometer. Endothelial cells pre-treated with 200 pM/well tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ; gift from Department of Hematology, University Hospital Utrecht, the Netherlands) were used as positive control; uninfected endothelial cells served as negative control. The human plasma pool was obtained from six healthy male volunteers. Clotting times in this assay were converted to percentage thromboplastin by relating clotting times to a standard thromboplastin dilution curve as described by Pronk et al<sup>35</sup>. Briefly, various dilutions of a rabbit brain thromboplastin suspension (Neoplastin, Boehringer Mannheim, Mannheim, Germany) in Ca<sup>2+</sup>-Michaelis buffer were pre-warmed for 1 minute at 37 <sup>o</sup>C in the cups of a coagulometer. Coagulation was started by addition of an equal volume of human pooled citrated plasma. The relation between clotting times and thromboplastin dilutions was used as a calibration curve. The procoagulant activities of the endothelial cell monolayers were read from the curve and expressed as percentages of thromboplastin time.

# Factor Xa generation assay

All studies were performed in 48-well tissue culture plates with confluent endothelial cell monolayers at 37 °C. Confluent endothelial cell monolayers were inoculated as previously described and incubated for 3 and 24 hours. The cells pre-treated with 200 pM/well TNF- $\alpha$  for 6 hours served as positive control. The cells were washed with Hepes buffer (Hepes 25 mM, NaCl 135 mM, KCl 4.5 mM, glucose 4.5 mM, pH 7.4, 0.3% BSA), and then washed twice with the same buffer containing CaCl<sub>3</sub> (5 mM). Factor VII and X were purified from plasma essentially as described by Bajaj<sup>36</sup> and Miletich <sup>37</sup>. They had a specific activity of 2000 U/mg (factor VII) and 100 U/mg (factor X) and were diluted in the Hepes-CaCl, buffer. Endothelial cells were pre-incubated with 100 µL factor VII (2.0 nM) for 5 minutes, after which 50 µL factor X (168 nM) was added and the cells were incubated for 15 minutes under gentle shaking. The reaction was terminated by addition of 25  $\mu$ L of the sample to 50  $\mu$ L EDTA (25 mM in Hepes buffer) in a 96-well plastic assay plate. To determine the total amount of activated factor X (factor Xa) formed, 100 µL 0.2 mM chromogenic substrate S2765 (Chromogenics, Mondal, Sweden) was added and the absorbance at 405 nm was measured for 1 minute at 5 second intervals in a V-max ELISA reader (Molecular Devices, Menlo Park, CA, USA). The concentration of factor Xa was calculated by using a standard curve for purified factor Xa<sup>38</sup>. Tissue factor activity was expressed as pmol factor Xa formed per minute by 1\*10<sup>5</sup> endothelial cells. The quantitative determination of tissue factor was performed using the Imubind Tissue Factor ELISA Kit (American diagnostics Inc, Greenwich, USA) recognizing TF-apo, TF and TF-VII complexes. Infected endothelial cell monolayers were lysed with 1 % triton-X-100 before measuring TF antigen.

# **Data analysis**

Results are expressed as means  $\pm$  standard error of mean (SEM). All data were analyzed by using the Statistical Package of Social Sciences (SPSS). Differences in quantitative measures were tested for significance by using the unpaired two-tailed Student's t-test, unless otherwise stated.

# **Results**

# Hemagglutinin, virus titers

Virus titrations were performed with fibroblasts and larynx carcinoma cells (HEP-2). The highest dilution of virus at which  $\geq$ 50% of 10 separate cell monolayers showed morphological changes due to infection (tissue culture infecting dose; TCID<sub>50</sub>) was determined for each pathogen. The TCID<sub>50</sub> values were 10<sup>-1</sup> for RSV; 10<sup>-2.7</sup> for CMV and 10<sup>-5.4</sup> for adenovirus. For the influenza strains the hemagglutinin titer was measured: 1:32 for influenza A (H1N1), 1:64 for influenza A (H3N2), 1:32 for influenza B and 1:16 for parainfluenza 1.

# Susceptibility of cell lines to virus infection

We assessed the *in vitro* susceptibility of different cell monolayers to infection with respiratory viruses. Endothelial cells, fibroblasts, monkey kidney cells and larynx carcinoma cells were all susceptible to infection with influenza A (H1N1, H3N2), influenza B, parainfluenza type 1, CMV, RSV and adenovirus (**Table 1**). Inoculated cell monolayers were centrifuged at 200g for 60 minutes, to increase virus infectivity. After infection all monolayers were intact as determined by phase contrast microscopy. Cell viability was ≥98 % as judged by trypan blue exclusion and the release of lactate dehydrogenase (LDH) after 24 hours' incubation with virus. This was of special concern because influenza viruses A and B are lytic viruses and destroy their host cell after infection.

 Table 1
 Susceptibility of different cell lines to infection by respiratory viruses

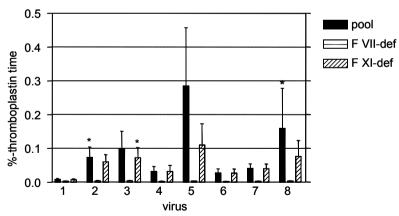
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	LLC-MK2	HEL	HEP-2	HUVEC
Inf A (H1N1)	100ª	108 + 21.0		67 + 19.2
Inf A (H3N2)	100ª	64 + 3.2		36 + 7.3
Inf B	100 <sup>a</sup>	70 + 15.2		78 + 11.7
Para	100 <sup>a</sup>	91 + 5.5		83 + 10.1
$CMV^b$		100 <sup>a</sup>		72 + 2.8
RSV		61 + 5.8	100ª	69 + 7.1
Adeno		68 + 10.3	100ª	31 + 0.6

Susceptibility  $\% \pm SD$  after a 24-hour incubation. a Cells were assigned a susceptibility of 100%. Actual infection percentage of the standard cells is approximately 20%. b 48-hour incubation. In each experiment 10 fields of the monolayer were examined. The values are the mean of 3 experiments in duplicate. Abbreviations: LLC-MK2, Rhesus monkey kidney cells; HEL, Human embryonic lung fibroblasts; HEP-2, Human epidermoid larynx carcinoma cells; HUVEC, Human umbilical vein endothelial cells.

In a pilot study, influenza virus infection caused major damage to all cell monolayers after 36 to 48 hours (data not shown). Endothelial cells were highly susceptible to infection with most strains of virus. Influenza A and B and parainfluenza also infected fibroblasts *in vitro*.

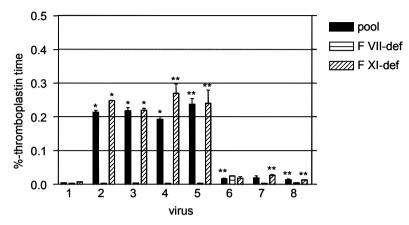
# **Procoagulant activity**

We determined the effect of infection with various respiratory viruses on the procoagulant activity of confluent, intact endothelial cell monolayers. Procoagulant activity was measured 3 and 24 hours after infection. The conversion of fibrinogen to fibrin was measured in a one-stage clotting assay with human pooled plasma or plasma deficient in factor VII or XI. After 3 hours' incubation all strains reduced the clotting time compared to that of uninfected cells (**Figure 1**), with the influenza strains reducing the clotting time by 55%. After 24 hours' incubation (**Figure 2**) the reduction in clotting time was even more pronounced (66% reduction) except for adenovirus. When factor VII deficient plasma was used, the clotting time was not reduced. Results showed that the reduction in clotting time was factor VII dependent and therefore mediated by the extrinsic coagulation pathway.



**Figure 1.** Procoagulant activity of human umbilical vein endothelial cells after incubation with respiratory viruses for 3 hours. Procoagulant activity was measured in a coagulometer with pooled human plasma (pool), factor VII-deficient plasma (F VII-def) or factor XI-deficient plasma (F XI-def). The procoagulant activity of the endothelial cell monolayers is expressed as a percentage of the thromboplastin time. X-axis: 1: uninfected confluent endothelial cell monolayer, 2: influenza A/H1N1/Taiwan/1/81, 3: influenza A/H3N2/Beijing/32/92-like, 4: influenza B/Netherlands/580/89, 5: parainfluenza-1, 6: cytomegalovirus, 7: respiratory syncytial virus, 8: adenovirus type 2. Levels are expressed as means  $\pm$  S.E., based on 4 separate experiments in duplicate. \* P < 0.05 compared with uninfected cells; paired t-test.

This indicates an increased expression of tissue factor (TF) on the endothelial cell membrane. There was no relationship between the percentage of endothelial cells infected by the different strains of virus and the reduction in clotting time.

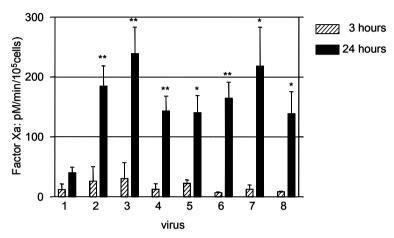


**Figure 2.** Procoagulant activity of human umbilical vein endothelial cells after incubation with respiratory viruses for 24 hours. Procoagulant activity was measured in a coagulometer with pooled human plasma (*pool*), factor VII-deficient plasma (*F VII-def*) or factor XI-deficient plasma (*F XI-def*). The procoagulant activity of the endothelial cell monolayers is expressed as a percentage of the thromboplastin time. X-axis: 1: uninfected confluent endothelial cell monolayer, 2: influenza A/H1N1/Taiwan/1/81, 3: influenza A/H3N2/Beijing/32/92-like, 4: influenza B/Netherlands/580/89, 5: parainfluenza-1, 6: cytomegalovirus, 7: respiratory syncytial virus, 8: adenovirus type 2. Levels are expressed as means  $\pm$  S.E., based on 4 separate experiments in duplicate. \*P < 0.001 compared with uninfected cells, \*\*P < 0.05 compared with uninfected cells; paired t-test.

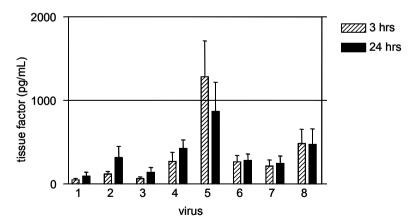
#### **Factor Xa generation**

To establish whether the reduction in clotting time was mediated by the production of tissue factor by endothelial cells, we decided to measure TF production as the generation of factor Xa, using a chromogenic substrate technique. Experiments were performed after incubation with virus for 3 and 24 hours. These time points were chosen to evaluate the effect of a 'full blown' infection without morphological damage to the monolayer seen with longer (36 hours) incubation. After 3 hours' incubation low levels of TF expression was observed on infected as well as on uninfected monolayers (**Figure 3**). After 24 hours' incubation all virus strains induced TF expression at different levels. Compared with the negative (uninfected) control there was a 3 to 4-fold increase in TF production. The confluent endothelial cell monolayers were morphologically intact after the 3- and 24-hour incubations with viruses.

Tissue factor antigen was measured in order to determine whether increased TF activity was the result of increased TF antigen expression on the endothelial cell surface. After 3 and 24 hours' infection increased TF antigen concentrations were measured in the endothelial cell homogenate (**Figure 4**).



**Figure 3.** Generation of factor Xa by human umbilical vein endothelial cells after incubation with respiratory viruses and incubation for 3 and 24 hours. X-axis: 1: uninfected confluent endothelial cell monolayer, 2: influenza A/H1N1/Taiwan/1/81, 3: influenza A/H3N2/Beijing/32/92-like, 4: influenza B/Netherlands/580/89, 5: parainfluenza-1, 6: cytomegalovirus, 7: respiratory syncytial virus, 8: adenovirus type 2. Levels are expressed as means + S.E., based on 5 separate experiments in duplicate. \* P < 0.001 compared with uninfected cells, \*\* P < 0.05 compared with uninfected cells; paired t-test.



**Figure 4.** Tissue factor antigen expression by endothelial cells after infection with viruses and incubation for 3 and 24 hours. X-axis: 1: uninfected confluent endothelial cell monolayer, 2: influenza A/H1N1/Tai-wan/1/81, 3: influenza A/H3N2/Beijing/32/92-like, 4: influenza B/Netherlands/580/89, 5: parainfluenza-1, 6: cytomegalovirus, 7: respiratory syncytial virus, 8: adenovirus type 2. Levels are expressed as means + S.E., based on 5 separate experiments in duplicate.

# Discussion

The mechanism by which infection with viruses contributes to increased cardiovascular morbidity and mortality is not known but may include alterations in the haemostatic balance. Endothelial cells may be a target for viruses in vivo, and infection may result in changes in endothelial cell functions. The association between infection with respiratory viruses and cardiovascular disease has not been confirmed in serological studies, but there is circumstantial evidence that these viruses may be a trigger for cardiovascular disease<sup>14,16,17</sup>. In this study, we found that viruses (influenza A, influenza B, parainfluenza, RSV, adenovirus and CMV) were able to infect different cell monolayers including human umbilical vein endothelial cells and human embryonic lung fibroblasts in vitro. Infection rates for CMV were similar to those previously published<sup>5</sup>. The susceptibility of endothelial cells to infection with respiratory viruses is less well documented. The observation that most respiratory viruses surveyed in this study were able to infect lung fibroblasts as well as endothelial cells may have implications for the *in vivo* situation where infection of the respiratory tract by respiratory viruses subsequently leads to infection of lung fibroblasts. Respiratory viruses then enter the vascular system and cause a more general infection, or viremia. In this way respiratory viruses can spread throughout the body, infecting other organs and other sites of the vascular tree. Endothelial cells, which line the lumen of blood vessels, can be targets for infection with these viruses.

Tissue factor is the physiological activator of the extrinsic coagulation pathway. Several stimuli (e.g. thrombin, IL-1ß, TNF-α) are known to induce the production of TF in endothelial cells, with detectable levels being reached after 2-4 hours and peak expression after 4-6 hours<sup>39</sup>. High concentrations of TF antigen and activity are found in coronary arteries from patients with unstable angina or myocardial infarction<sup>40</sup>. In this study viruses induced endothelial cell procoagulant activity *in vitro*. All viruses tested reduced the clotting time in a time-dependent manner. By different experimental set-ups, we could show that this reduction in clotting time was a result of TF expression on endothelial cells. Low levels of TF activity were detectable on infected as well as on uninfected cell monolayers after 3 hours' incubation (**Figure 3**), and there was a factor VII-dependent reduction in the clotting time at this time point (**Figure 1**). Tissue factor antigen levels were already increased after 3 hours' incubation (**Figure 4**). At 24 hours, TF antigen levels were similarly increased as at the 3 hours time point, but this was accompanied with a marked increase in TF activity. Apparently, the

increased activity of TF was not the result of increased antigen expression only. Most likely changed phospholipid expression on the cell membrane, as a result of the virus infection, leads to increased TF activity. Increased TF activity due to elevated levels of phosphatidylserine<sup>41</sup> on the cell membrane was not likely, because apoptosis was not observed after infection as determined by the lack of annexin V binding to infected endothelial cells. Another explanation for the discrepancy between levels of TF activity and TF antigen may be the fact that TF antigen was measured in the cell homogenate. After 3 hours' incubation TF antigen was probably still intracellular and at the 24 hours measurement TF was expressed on the cell membrane which subsequently resulted in the measured increase in TF activity. Thus, the clotting time reduction on endothelial cells after virus infection is due to increased TF activity as a result of up regulation of TF antigen expression in combination with increased activity of the expressed TF possibly by changed co-expression of phospholipids on the cell outer membrane. Our results are in line with the studies by Key et al<sup>42,43</sup>. They showed TF expression on the cell surface of herpes simplex infected endothelial cells. In contrast to our study, they observed that TF activity returned to baseline after 20 hours' infection. However, an other study has shown that TF does not play a major role at all in endothelial cell procoagulant activity induction by CMV<sup>5</sup>. In that study, the facilitated formation of the prothrombinase complex on endothelial cells was considered important. Key et al<sup>43</sup> demonstrated TF dependent procoagulant activity in endothelial cells infected with herpes simplex virus-1 (HSV-1) by using a neutralizing antibody to human TF. In both studies<sup>5, 43</sup> and in a study by Pryzdal and coworkers<sup>44</sup> the facilitated assembly of the prothrombinase complex could be related to the viral envelope of HSV-1 and CMV. Presentation of antigens of the viral envelope on the surface of endothelial cells promotes procoagulant activity. However, we found that adenovirus, a non-enveloped virus, could also induce procoagulant activity in endothelial cells in conjunction with TF expression. Viral antigens may enhance prothrombinase complex assembly on the surface of endothelial cells which in turn facilitates thrombin generation<sup>45</sup>. This in turn stimulates endothelial cells to express TF. Moreover, the endogenous production of IL-1ß by endothelial cells, in response to viral infection, might induce TF expression of the same or neighboring cells.

The observation that influenza viruses, due to their lytic properties, caused major damage to the endothelium after a 36-hour incubation may be of importance. A dying cell expresses phosphatidylserine on its surface, and after lysis the prothrombotic extracellular matrix is exposed to the vascular lumen. We could not demonstrate

annexin V binding to infected endothelial cells but cell lysis may be a mechanism by which respiratory infections could contribute to alterations in the procoagulantanticoagulant balance. We did not find a relationship between the proportion of endothelial cells infected with virus and the induction of procoagulant activity. A linear relationship has been described between the number of endothelial cells infected with HSV-1 and TF production<sup>42</sup>. It is not possible to compare the infectivity of different strains of virus as long as the mechanism of TF activity induction is not known. It is possible that the viruses induce IL-1 $\Omega$  or TNF- $\alpha$  production in the endothelial cells. In turn, is IL-1ß a mediator of TF expression on the endothelium. However, we could not detect TNF- $\alpha$  or IL-1 $\beta$  in the supernatant after infection of endothelial cells by the different viruses (data not shown) but it is possible that very low (undetectable) TNF- $\alpha$ and IL-1ß concentrations are responsible for TF expression on endothelial cells. In conclusion, we have shown that respiratory viruses can infect human endothelial cells and fibroblasts in vitro. Both enveloped and non-enveloped respiratory viruses induce procoagulant activity on cultured endothelial cells by stimulating the expression of TF antigen and enhancing the activity of expressed TF antigen. Also, facilitated prothrombinase complex assembly may contribute to procoagulant activity of infected endothelial cells. If similar processes occur in vivo, then infections with respiratory viruses give rise to endothelial procoagulant activity and hence contribute to an increased risk of atherothrombosis

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3

Procoagulant and inflammatory response of virus-infected monocytes

### **Abstract**

# **Background**

Monocytes play a prominent role in inflammation, coagulation and atherosclerosis by their ability to produce tissue factor (TF) and cytokines. The aim of the present study was to establish whether virus-infected monocytes initiate coagulation. In addition, the production of cytokines by monocytes may accelerate the chronic process of atherosclerosis and may contribute to coronary syndromes by eliciting plaque instability.

#### Materials and methods

Monocytes were isolated by Vacutainer\*, BD Biosciences, Alphen aan den Rijn, Netherlands and subsequent magnetic cell sorting (MACS\*, Miltenyi Biotec, Bergish Gladbach, Germany). Coagulation times in normal pooled plasma and Factor VII-deficient plasma were measured after infection with cytomegalovirus (CMV), Chlamydia pneumoniae (Cp) and influenza A\H1N1. Anti-TF antibodies were added to neutralize TF expressed on monocytes. Interleukins (IL) 6, 8 and 10 were measured in the supernatants.

### Results

Chlamydia pneumoniae- and CMV-infected monocytes decreased the clotting time by 60% and influenza-infected monocytes by 19%, as compared to uninfected monocytes. Procoagulant activity was absent when Factor VII-deficient plasma or anti-TF antibodies were used. Monocytes produced both IL-6 and IL-8 after infection with CMV (317 pg/mL and 250 pg/mL) or *Cp* (733 pg/mL and 268 pg/mL). Similar results were obtained for influenza virus-infected monocytes, but the levels of both cytokines were 3-5 fold higher (1797 pg/mL and 725 pg/mL). Interleukin-10 was not produced by infected monocytes.

#### Conclusion

The procoagulant activity of virus-infected monocytes is TF-dependent. Although influenza infection did not generate a significant reduction in clotting time, the pronounced expression of IL-6 and IL-8 may induce local and/or systemic inflammatory reactions, which may be associated with plaque rupture and atherosclerosis. The lack of production of the anti-inflammatory cytokine IL-10 may even accelerate these processes.

Keywords: Chlamydia pneumoniae, cytomegalovirus, influenza A virus, interleukins, monocytes, tissue factor.

# Introduction

Atherosclerosis is thought to be an inflammatory disease<sup>1-3</sup>. The cause of this chronic low-grade inflammation is not unambiguous, but may well be a combination of classical risk factors or novel risk factors such as micro-organisms. There is increasing evidence that viruses have a role in the etiology of atherosclerosis<sup>4,5</sup> and several studies have identified viruses and bacteria as potential agents contributing to atherogenesis<sup>6-10</sup>. Cytomegalovirus (CMV) and Chlamydia pneumoniae (Cp) have been found in atherosclerotic plagues and smooth muscle cells<sup>11-13</sup>. Also, several seroepidemiological studies have shown association between seropositivity for Cp and increased risk of cardiovascular diseases<sup>14,15</sup>. At present, controversy exists about the role of chronic CMV infection in relation to the risk of cardiovascular disease. In general predisposition of CMV for inflammation and subsequent susceptibility for chronic coronary artery disease (CAD) is well accepted, but seropositivity for CMV alone is unlikely to be a strong risk factor for CAD<sup>16-19</sup>. Cytomegalovirus-lgG titers rather than seropositivity, seropositivity for more than one agent, and pathogen burden, are associated with enhanced inflammatory responses, potentially accelerating the atherothrombotic process 20-22. However, most predictive indicators for the risk of CAD are seropositivity for CMV with increased IgG titers in correlation with elevated IL-6 or CRP levels, reflecting a (subclinical) inflammatory response<sup>23–26</sup>. Infection of the endothelium shifts endothelial properties from anticoagulant to procoagulant as a result of the production and expression of tissue factor (TF)<sup>27,28</sup>. Although normally absent from all intravascular cell types, TF can be synthesized by endothelial cells, smooth muscle cells and macrophages in response to inflammatory stimuli, endotoxins or immunological compounds such as lipopolysaccharides (LPS), interleukin-1 (IL-1) and tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) <sup>29,30</sup>. Elevated expression of TF on monocytes/ macrophages in atherosclerotic plaques suggests a role for TF in fatal thrombosis associated with plaque rupture<sup>31,32</sup>. Virus infection of the endothelium results in the promotion of monocyte adhesion<sup>33-35</sup> as well as cytokine production<sup>36-38</sup>. The cytokines IL-6 and IL-8 may induce monocyte procoagulant activity by stimulating TF expression. This may link inflammation to thrombotic events<sup>39</sup>. Undoubtedly cross-talk between cells of the immune system and the vessel wall mediated by cytokines and other pro-inflammatory substances is responsible for initiating cascade effects leading to vascular disorders<sup>40</sup>. Studies conducted on the effects of infection on monocyte activation reveal inflammatory responses by production of tissue factor<sup>41,42</sup> and various

proinflammatory cytokines like IL-1 and IL-6, but also potent chemo-attractants like monocyte chemo attractant protein1 (MCP-1) are readily produced<sup>43-45</sup>. Next to this, foam cell formation and lipid oxidation are induced and loss of anti-inflammatory properties of high-density lipoproteins has been recorded<sup>46,47</sup>. In combination with up regulation of adhesion molecules by endothelial cells<sup>48</sup>, monocytes harboring virus and oxidized low-density lipoproteins eventually infiltrate the vessel wall<sup>46,49</sup>. In the present study, we investigated the response of monocytes to infection with *Cp*, CMV and influenza A *in vitro*. *Chlamydia pneumoniae* and CMV cause more chronic events while influenza represents an agent responsible for acute cardiovascular complications. Considering the distinct properties of these viruses, distinct modes of action by monocytes infected with these viruses may be observed.

# **Methods**

# **Isolation of monocytes**

For isolation of the mononuclear fraction, fresh blood from healthy volunteers was drawn by venapuncture either into citrated tubes for Ficoll/Hypaque isolation or into mononuclear cell preparation tubes (Vacutainer CPT°, BD Biosciences, Alphen aan den Rijn, Netherlands). Ficoll/ Hypaque isolation was performed according to the well defined protocol of Boyum 50. Vacutainer isolation tubes were inverted a few times and centrifuged at 1400g for 20 min. At this speed a gel barrier separates the mononuclear fraction from plasma and the fraction containing erythrocytes and polymorph nuclear neutrophils. Plasma from each donor was stored at - 70 °C. Subsequently, monocytes were purified from the mononuclear fraction by means of a magnetic-activated cell sorting protocol (MACS°, Miltenyi Biotec, Germany). Monocytes can be isolated by two protocols [1]: depletion of non-monocytes with a cocktail of CD3, CD7, CD19, CD45RA, CD56 and anti-IgE antibodies, leaving an untouched monocyte fraction or [2] positive selection of CD14+ cells (monocytes). For reasons discussed later in this paper, positive selection was our method of choice. For the isolation of monocytes by the MACS positive-selection protocol, 10<sup>7</sup> cells were suspended in 80 μL cold phosphatebuffered saline (PBS) pH 7.2 supplemented with 0.5% fetal calf serum (FBS) and 2 mM EDTA. Cells were magnetically labeled with 20 μL CD14-microbeads<sup>®</sup> (Miltenyi Biotec) per 10<sup>7</sup> cells, mixed well and incubated at 4 °C for 15 min. Then cells were washed with 10 volumes of cold buffer and centrifuged at 300g and 4 °C for 10 min. The cell pellet was resuspended in buffer (500 μL buffer/108 cells) and applied to a prewashed LS+

positive selection column (Miltenyi Biotec), which was placed in the magnetic field of a MACS separator\* (Miltenyi Biotec). Unlabelled (CD14-) cells passed through the column and were discarded with the effluent. The column was rinsed three times with PBS. After the column was removed from the magnetic field, the retained CD14+ cells were eluted in cold PBS. The yield and purity of the isolated cells were determined by flow cytometry. Cells were kept on ice until experiments were performed.

# **Preparation of virus stocks**

Human embryonic lung (HEL), human epidermoid larynx carcinoma (HEP-2) and LLC-MK2 cells were cultured at 37 °C and 5% CO<sub>3</sub> in Eagle's minimum essential medium with Earle's salts (EMEM, Life Technologies Ltd, Paisley, Scotland) with 10% FBS (Life Technologies Ltd). This culture medium was supplemented with 2 mmol/L L-glutamine (Life Technologies Ltd), 5 mL nonessential amino acids (Life Technologies Ltd), 10 mg/L vancomycin (Faulding Pharmaceuticals, Brussels, Belgium), 4 mg/L fungizone (Bristol-Meyers Squibb, Woerden, the Netherlands) and 10 mg/L gentamycin (Schering Plough, Maarssen, the Netherlands). The same supplements were also added to the media used for the propagation of the virus strains. A clinical isolate of CMV was propagated in HEL cells in EMEM containing 2% FBS, 20 mM Hepes and supplements. At >80% cytopathologic effect (CPE), CMV-infected HEL cells were detached with trypsin/EDTA solution (Life Technologies Ltd) and centrifuged. The cell pellet was resuspended in EMEM containing 2% FBS, 10% DMSO (Sigma-Aldrich, Zwiindrecht, the Netherlands) and supplements. Cytomegalovirus stock was aliquoted and stored at -70 °C. Chlamydia pneumoniae-strain AR 39 was propagated in HEP-2 cells in EMEM 10% FBS with 0.1% cycloheximide (Sigma-Aldrich) and supplements at 37 °C / 5% CO<sub>2</sub>. After 72 h cells were frozen and thawed to release the elementary bodies. After a short centrifugation step, cell debris was discarded and 0.2 M SPG medium (2.088 g/L K<sub>2</sub>HPO<sub>4</sub>, 1.088 g/L KH<sub>2</sub>PO<sub>4</sub>, 68.46 g/L saccharose, 7.16 g/L L-glutamine, 10% FBS, 2.5 mg/L fungizone, 23 mg/L vancomycin and 18 mg/L gentamycin) was added (1:1 v/v). Chlamydia pneumoniae stock was aliquoted and stored at - 70 °C. Influenza A/ H1N1/1/86/Singapore was propagated in LLC-MK2 cells in serum-free medium M199 with 20 mM Hepes and supplements. At > 80% CPE, the virus was harvested by freezing and thawing the cells. After centrifugation, the pellet was discarded and 0.2 M sucrose phosphate medium was added to the supernatant (1:1 v/v). Influenza A stock was aliquoted and stored at -70 °C. The 50% tissue culture infective dose (TCID<sub>50</sub>) of each virus stock was calculated using the method of Reed & Muench<sup>51</sup>.

For CMV and influenza A reading CPE was used to calculate  $TCID_{50}$  titers. In the case of Cp, the  $TCID_{50}$  titer was calculated based on the number of immunofluorescent units per field in the infected HEP-2 cells after staining with Chlamydia culture confirmation monoclonal antibodies (de Beer, Diessen, the Netherlands).

# Infection of monocytes

Isolated monocytes were counted and brought to a concentration of  $10^5$  cells/mL in RPMI medium with 10% autologous plasma. Coagulation tubes (P1000 sample cups, Amelung; Boom BV, Meppel, the Netherlands) were precoated, for at least 1 h, with fibronectin (CLB, Amsterdam, the Netherlands) diluted 1:100 in PBS. Aliquots of 1 mL ( $10^5$  monocytes) were distributed over the tubes. Subsequently, 100 mL of each virus stock was added to one tube. The multiplicity of infection (MOI) was 0.4 for CMV, 0.1 for *Cp* and 0.04 for Influenza A. As negative control,  $100 \, \mu L$  of RPMI instead of the virus was added. For the positive controls,  $10^5$  monocytes were incubated with  $1 \, \mu g/mL$  LPS (prepared in endotoxin-free water; Sigma-Aldrich). Tubes were incubated at 37 °C and 5%  $CO_2$  for 4 h and then centrifuged at 300g and 4 °C for 5 min. Supernatants were collected and frozen at -70 °C for cytokine assays. Monocytes were washed immediately in cold PBS, refreshed with  $100 \, \mu L$  PBS, and kept on ice until clotting assays were performed.

# **Coagulation assay**

Normal pooled plasma was prepared from the blood of nine healthy donors who did not use aspirin or oral anticonception. Freeze-dried Factor VII-deficient plasma (Kordia Biopool, Leiden the Netherlands) was reconstituted for use with 1 mL of distilled water. Clotting assays were performed at 37 °C in a coagulometer (KC4A Amelung/Boom BV), which records the time taken for fibrin formation. To 100  $\mu$ L (10<sup>5</sup> cells) of infected monocytes, incubated in a coagulation tube, were added: 100  $\mu$ L of Michaelis buffer (28.5 mM sodium barbital, 28.5 mM sodium acetate, 50 mM sodium chloride and 33 mM calcium dichloride), one metal bead, and 100  $\mu$ L of normal pooled plasma or Factor VII-deficient plasma. All components were kept cold until use. Clotting times at 37 °C were measured as described. To investigate whether TF played a central role in the reduction of clotting time, samples in normal pooled plasma were also incubated with 1:100 diluted neutralizing antibodies against TF (American Diagnostica Greenwich, CT).

# **Cytokine assays**

Interleukins IL-6, IL-8 and IL-10 were measured in the supernatants of virus-infected monocytes. Sandwich-Elisa's were used (IL-6, IL-8 and IL-10 Pelikine compact human ELISA kits; CLB the Netherlands) and performed as prescribed by the manufacturer. Before the assay, the frozen supernatants were thawed quickly in a 37 °C water bath and brought to room temperature. Samples were prediluted in buffer, at least 1:2 or more, depending on the expected IL titer. A standard curve (serial dilution of the human recombinant cytokine) was included in each experiment. All experiments were performed at least three times in duplicate. The absorbance of standards and samples was read at 450 nm using an ELISA reader (ELx800, Biotek Instruments Inc). The cytokine concentration of the samples was calculated by interpolation from the standard curve and corrected for the dilution factor.

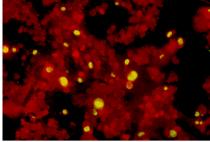
#### **Statistics**

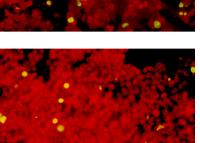
Results are expressed as means  $\pm$  standard errors (SE) of three experiments. ANOVA analysis of variances was performed to check for variances among means in the test populations. P-values of means  $\pm$  SE were calculated by comparing all samples to the negative control using Dunnett's multiple comparisons test, which corrects for unequal variances.

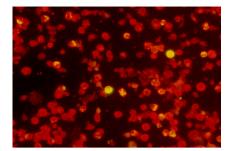
# Results

#### Infection of monocytes

After infection,  $TCID_{50}$ -titres of the virus stocks in standard cell lines and MOI values in monocytes were calculated for each virus strain.  $TCID_{50}$  values were:  $10^{-4.25}$  for CMV,  $10^{-3.75}$  for Cp and  $10^{-3.25}$  for influenza A. The *in vitro* susceptibility of monocytes to the virus strains was determined with immunofluorescent staining for the presence of virus in monocytes. Cytomegalovirus-infected monocytes demonstrated the typical nuclear staining of infected cells (**Figure 1a**), whereas influenza- or Cp-infected monocytes showed cytoplasmatic fluorescence (**Figure 1b, c**). It appeared that all strains could infect monocytes, but in all cases the infection percentage was below 5% when undiluted virus stocks were used (MOI: 0.4, 0.1 and 0.04 for CMV, Cp and influenza virus, respectively).







**Figure 1** Fluorescence microscopy images of virus-infected monocytes. Magnification 200x after overnight incubation; Red background: uninfected monocytes. Green areas: virus-infected monocytes stained with FITC-labeled anti-virus antibodies.

(a) CMV, specific staining of CMV in the nuclei of the infected cells. (b) Influenza A, smooth staining pattern of influenza A in monocytes. (c) Cp, denser staining pattern of Cp in the cytoplasm of monocytes. (*For color figure see page 218*)

# **Isolation of monocytes**

In both (Ficoll and Vacutainer) fractions, obtained by MACS-depletion, large amounts of debris were present and some granulocytes were left. In both Ficoll and Vacutainer depletion prepared suspensions, 80% of the monocytes were covered with activated (CD41+/CD42+) platelets. Practically no free platelets were present in the suspensions. In suspensions prepared by the positive selection method little debris was present and 78% of all isolated cells were monocytes with only a few platelets (3%) attached (**Table 1**).

**Table 1** Immunophenotyping of isolated leucocytes

	Ficoll	VPT	VPT
	+ depletion *	+ depletion †	+ positive selection *
CD14+/CD45+	78%	95%	78%
CD42+/CD14+	77%	83%	2%
CD42-/CD14+	22%	16%	97%
CD41+	47%	78%	3%
CD41+/CD42+	46%	78%	3%

FACS-analysis of isolated monocyte fractions; determination of yield and purity of monocyte fractions obtained from three different isolation protocols: \* Ficoll isolation followed by MACS depletion of non-monocytes; † Vacutainer CPT isolation followed by MACS depletion of non-monocytes; ‡ Vacutainer CPT isolation followed by MACS CD14-isolation protocol (positive selection). CD45+: leucocytes; CD14+: monocytes; CD41+: platelets; CD42+: activated platelets.

# **Procoagulant activity**

Infection of monocytes with *Cp* or CMV for 4 h resulted in a significant reduction of the clotting time by approximately 60% (140  $\pm$  3.8 and 134  $\pm$  8.9 s, respectively; p<0.05) compared with that of uninfected monocytes (333  $\pm$  22.1 s) [**Figure 2**]. The clotting time of monocytes infected with influenza A/H1N1 virus (273  $\pm$  81 s) was reduced by 19% (p>0.05). The same experiments were performed with Factor VII-deficient plasma. In these experiments no differences between clotting times were recorded (p>0.05). Moreover, addition of anti-TF antibodies during incubation of the monocyte suspensions with virus completely blocked the reduction in clotting time to the level of the negative control. ANOVA analysis on variances showed that the variances between test samples in the normal pooled plasma were significantly different (F=6.4/ P=0.0079). In contrast, the analysis of variances among Factor VII-deficient- and anti-TF neutralized sample populations no longer indicated any differences (F=0.75/ p=0.58 and F=0.96/p=0.47, respectively).

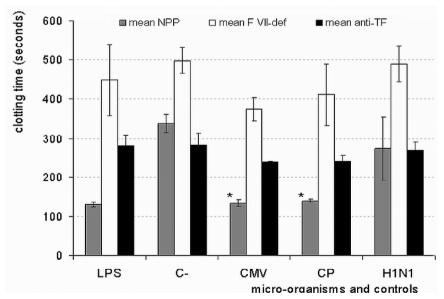


Figure 2 Procoagulant activity of infected monocytes.

Mean clotting times of virus-infected monocytes in seconds  $\pm$  SE based on three experiments in duplicate. LPS: positive control (1 µg/mL), C-: negative control (culture medium/no virus); H1N1: Influenza A/ H1N1/1/86/Singapore; *Cp: Chlamydia pneumoniae*, CMV: Cytomegalovirus; NPP: normal pooled plasma; F VII-def: Factor VII-deficient plasma; anti-TF: with anti-tissue factor antibodies pre-incubated samples in NPP. \*P < 0.05 indicating significant difference from the negative control (Dunnett's multiple comparisons test).

# **Cytokine assays**

Results from IL-6 and IL-8 experiments in the supernatants of the infected monocytes were more or less identical. *Chlamydia pneumoniae*-infected monocytes produced significantly more IL-6 (733  $\pm$  84 pg/mL; p<0.01) than did the uninfected monocytes (90  $\pm$  61 pg/mL), whereas the difference was not significant for CMV-infected monocytes (317  $\pm$  94 pg/mL IL-6; p>0.05). IL-8 production by *Cp*- and CMV-infected monocytes was marginally elevated (268  $\pm$  58 pg/mL and 250  $\pm$  48 pg/mL, respectively; p>0.05). Influenza A/H1N1-infected monocytes produced high levels of IL-6 (1797  $\pm$  244 pg/mL; p<0.001) and IL-8 (725  $\pm$  92 pg/ mL; P<0.001) [**Figure 3**]. Interleukin 10 was not detected in the supernatants or in the LPS control.

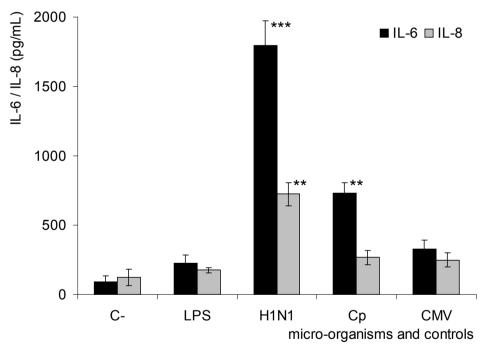


Figure 3 Interleukine production of infected monocytes.

Mean  $\pm$  SE concentrations of interleukin-6 (IL-6) and interleukin 8 (IL-8) in the supernatants of virus-infected monocytes, based on three experiments in duplicate. LPS: positive control (1 µg/mL), C-: , negative control (culture medium/no virus); CMV: cytomegalovirus; H1N1: Influenza A/H1N1/1/86/Singapore; *Cp: Chlamydia pneumoniae.* \*\*\*P < 0.001, indicating extremely significant difference from the negative control (Dunnett's multiple comparisons test). \*\*P < 0.01, indicating very significant difference from the negative control (Dunnett's multiple comparisons test).

### Discussion

# **Isolation of monocytes**

Ficoll isolation is the most commonly used method for isolation of peripheral blood mononuclear cells<sup>50,52,53</sup>. Care must be taken to prevent monocytes becoming activated during isolation procedures and to obtain platelet-free monocytes. Platelets adherent to monocytes substantially contribute to procoagulant activity by expressing glycoprotein IIb/IIIa complexes; the receptor complexes for fibrinogen. Moreover, P-selectin on activated platelets is able to up-regulate TF expression. Several authors have emphasized the importance of obtaining platelet-free monocyte preparations<sup>54-58</sup>. After the MACS depletion procedure, a large number of monocytes had platelets attached to them. This is probably because the cocktail of haptene conjugated antibodies, which was used in the depletion protocol, contains antibodies against almost all cell types (CD3, CD7, CD19, CD45RA, CD56 and IgE) but not against platelets (CD41 or CD61). Fluorescence-activated cell sorter analysis showed that platelets were activated and attached to 80% of the monocytes. Compared with the depletion protocol, the positive selection method yielded monocyte suspensions of greater purity and with only marginal interference from platelets; after all, only 3% of monocytes had platelets on their membrane. In our hands, Vacutainer CPT<sup>\*</sup> isolation combined with MACS° CD14-positive selection provided a reliable technique for the isolation of unstimulated monocytes from peripheral blood.

# Procoagulant activity and acute-phase response

In previous research we described the procoagulant activity of endothelial cells after infection with respiratory viruses. Viral pathogens induced a TF-dependent reduction of clotting time by approximately 55% <sup>27</sup>. Thiruvikraman et al. found TF to be localized extracellular in high levels in atherosclerotic plaques<sup>31</sup>. Thrombosis can be initiated rapidly when arterial plaques rupture and expose their contents to the bloodstream. Co-culturing of endothelial cells with monocytes revealed that TF induction occurred in monocytes rather than in endothelial cells<sup>59</sup>. Levels of TF are already detectable 2-4 h after infection/stimulation and TF expression reaches a maximum at 4-6 h after infection. In our coagulation experiments, CMV and *Cp* as well as influenza reduced the clotting time by 19-60%. The initiation of coagulation by virus-infected monocytes was a result of the expression of TF. By using Factor VII-deficient plasma and anti-TF

antibodies, we showed that TF plays an important role in virus-induced monocytic procoagulant activity. Infection with CMV and Cp induced the production of modest levels of IL-6 and IL-8, whereas infection with influenza A strongly stimulated the production of IL-6 and IL-8. Our results indicate that only small doses of an infectious virus are needed to stimulate monocytes to exert considerable immunological effects. There have been conflicting reports about whether influenza virus is able to infect monocytes. It is imaginable that viral proteins are synthesized de novo, but that there is no or little productive infection. Using immunofluorescence, we showed that influenza virus did infect monocytes (Fig. 1b), although the infection percentage was very low (< 5%). Secondary bacterial infections are not uncommon after influenza infections. It was assumed that increased sensitivity to bacterial products is the result of 'priming' of cells<sup>60</sup> because low amounts of LPS can accelerate cytokine production substantially<sup>61,62</sup>. As described earlier, TF-expression is elevated after infection, which may be a direct virus effect but which may also be triggered by IL-6. By producing IL-6, infected monocytes stimulate uninfected cells to express TF. Continuing viral infection may sustain excess TF production and exhaust the inhibitory effects of tissue factor pathway inhibitor in vivo<sup>63</sup>. We did not detect IL-10 production by virus-infected monocytes, not even in the LPS control. However, because cytokine production is maximal at 24 h after infection, it is possible that the infection times we used were too short to stimulate IL-10 production. After 4 h IL-10 production may still be minimal or only detectable at the mRNA level. During viral infections monocytes predominantly produce the inflammatory cytokines IL-6 and IL-8. Interleukin-10 is detected predominantly in atherosclerotic lesions, but monocytes may produce this cytokine transiently. Nevertheless detection of IL-10 is of utmost importance because it is a potent down regulator of pro-inflammatory cytokine production, thus being a potent immunosuppressor with potential for therapeutic use. The balance between pro-inflammatory and anti-inflammatory cytokines in the atherosclerotic plague may be decisive for the progression of the lesion. This concept is confirmed by a recent study showing that patients with unstable angina had significantly lower serum IL-10 concentrations compared with patients with chronic stable angina<sup>64</sup>. Moreover experimental work in animals confirms the protective role of IL-10 in atherosclerosis<sup>65,</sup> 66. We conclude that virus infection of monocytes triggers pro-atherosclerotic and prothrombotic processes in vitro.

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Azithromycin inhibits interleukin-6 but not fibrinogen production in hepatocytes infected with Cytomegalovirus and Chlamydia pneumoniae

# **Summary**

Chlamydia pneumoniae and cytomegalovirus (CMV) have been associated with the development of atherosclerosis. Inflammatory stimuli initiate the biosynthesis of fibrinogen, interleukin (IL)-6 and plasminogen activator inhibitor 1 (PAI-1) in the liver. Chronic infection may perpetuate the inflammatory status. We hypothesized that infection of human hepatocytes with the intracellular pathogens *C. pneumoniae a*nd CMV accelerates biosynthesis of fibrinogen, IL-6, and PAI-1 but that this biosynthesis can be reduced with the use of azithromycin.

HepG2 human hepatocytes were infected with *C. pneumoniae* and CMV *in vitro* in the presence of 0, 0.016, 0.125, or 1  $\mu$ g/mL azithromycin. We measured IL-6, PAI-1, and fibrinogen after 24, 48, 72, and 96 hours. *C. pneumoniae*–infected hepatocytes produce IL-6 (2667  $\pm$  309 pg/mL vs. 137  $\pm$  120 pg/mL in uninfected cells after 96 hours. Incubation with 0.016  $\mu$ g/mL azithromycin decreased IL-6 levels to a mean of 1516  $\pm$  402 pg/mL, and incubation with 0.125 and 1  $\mu$ g/mL azithromycin decreased IL-6 to 871  $\pm$  364pg/mL and 752  $\pm$  403 pg/mL, respectively. *C pneumoniae*–induced IL-6 production was time- and dose dependent.

The interaction of *C. pneumoniae* with azithromycin treatment was significant, indicating an inhibitory effect of azithromycin on *C. pneumoniae*–induced IL-6 production. CMV infection did not lead to IL-6 production by hepatocytes.

*C. pneumoniae* and CMV infection did not induce any changes in PAI-1 production. Fibrinogen production was increased by CMV infection after 72 hours (838  $\pm$  88 ng/mL; p<0.01) and after 96 hours by infection with both *C. pneumoniae* and CMV (765  $\pm$  100 ng/mL and 846  $\pm$  123 ng/mL, respectively; p<0.05). Azithromycin did not suppress CMV- or *C. pneumoniae*—induced fibrinogen production. Moreover, we could not confirm an anti-inflammatory effect of azithromycin in experiments with crosstitrations of azithromycin against either IL-1 or IL-6 (p>0.05). Azithromycin reduces *C. pneumoniae*—induced IL-6 production, but not fibrinogen production, by human hepatocytes. This is a result of the antimicrobial properties of azithromycin and not a direct anti-inflammatory effect.

#### **Abbreviations**

ANOVA: analysis of variance; APP: acute-phase protein; APR: acute phase reaction; BGM: Buffalo green monkey kidney cells; CMV: cytomegalovirus; *Cp: Chlamydia pneumoniae*; CRP: C-reactive protein; EMEM Earle's minimal essential medium; FBS: fetal bovine serum; HEL: human embryonic lung cells: Hep: human epithelioma; IL-6: interleukin 6; Lp(a): lipoprotein A; MBC: minimal bactericidal concentration; MIC: minimal inhibitory concentration; MOI: multiplification of infection; PAI: plasminogen-activator inhibitor; TCID<sub>50</sub>: 50% tissue culture infective dose; TNF: tumor necrosis factor

#### Introduction

Chronic low-grade inflammation plays an important role in the development of atherosclerosis<sup>1</sup>. Whether infectious agents play a causative role in atherogenesis is still a matter of debate. Several studies have indicated that Chlamydia pneumoniae and CMV are associated with the development of atherosclerosis<sup>2-4</sup>. Direct vesselwall infection may trigger atherogenic processes such as procoagulant activity, local inflammation, and endothelial dysfunction, leading to the development of atherosclerotic plague and eventually to plague rupture, exposing the thrombogenic plaque content to the bloodstream<sup>5,6</sup>. The liver is able to produce a variety of APP's such as IL-6, CRP, TNF-alpha, fibrinogen, haptoglobin, PAI-1, and complement factors C3 and C4, leading to plasma concentrations as high as 2 to 100 times normal<sup>8-11</sup>. In pathophysiological conditions, plasma levels can remain increased during chronic inflammatory processes such as atherosclerosis<sup>12–15</sup>. Fibrinogen is synthesized by hepatocytes under control of inflammatory cytokines. Oncostatin M and the pleiotropic cytokine IL-6 are considered the most potent stimulators of fibrinogen expression. PAI-1 is a major physiologic inhibitor of plasminogen activation. High plasma levels of PAI-1 have been associated with thrombotic and arterial disease. Increased synthesis of PAI-1 in the liver and cellular components of the atherosclerotic plaque may contribute to the thrombotic complications associated with plaque rupture and the accumulation of extracellular matrix deposits <sup>26,27</sup>. Macrolide antibiotics are effective in the treatment of intracellular infections because of their high cellular accumulation and tissue retention<sup>28-30</sup>. The effects of neomacrolides such as roxithromycin and azithromycin on inflammation have been delineated in animal studies identifying anti-inflammatory activities on top of the expected antibiotic properties of these antibiotics<sup>31,32</sup>. Treatment of *C. pneumoniae* infection with macrolide antibiotics has been shown to lead to decreased plasma concentrations of CRP, TNF-α, and IL-6 but did not reduce the incidence of cardiovascular events<sup>19,33-35</sup>. This may be due to insufficient dose regimens, short duration of treatment and follow-up, underpowered studies, or ineffective treatment of C. pneumoniae in monocytes.

In this *in vitro* study we investigated whether azithromycin was able to inhibit *C. pneumoniae* infection of human hepatocytes and reduce the inflammatory response of hepatocytes to infection.

## **Methods**

# C pneumoniae and CMV

We propagated *C. pneumoniae* strain AR 39 and a clinical isolate of CMV in Buffalo green monkey kidney cells (BGM) and human embryonic lung cells (HEL), respectively. Both strains were harvested at greater than 80% CPE(cytopathologic effect) and virus stocks were aliquoted and frozen at -90 °C until the time of infection. In an attempt to determine the infectious doses for the experiments, we calculated the  $TCID_{50}$  of each strain in accordance with the method of Reed and Muench.

# Hep-G2 cell culture

We grew 2 cell lines of Hep-G2 human hepatoma cells (1 donated by the oncology department of the University Hospital of Utrecht and the other obtained from the European Collection of Cell Cultures no. 85011430) in 80 cm² tissue-culture flasks (Nunc, Roskilde, Denmark) using EMEM (Life Technologies Ltd, Paisley, Scotland) supplemented with 10 % FBS (Life Technologies) and 2 mM L-glutamine (Life Technologies), 5 mL of nonessential amino acids (Life Technologies) and 4 mg/L fungizone (Bristol-Meyers Squibb, Woerden, the Netherlands). Before the experiments, we propagated HepG2 cells and maintained them for at least 3 passages in EMEM without any antibiotics. In the infection experiments, cells were seeded in 48-well plates (Corning Costar, Corning Inc, Corning, NY) and grown to confluence at 37 °C under 5% CO<sub>2</sub>, also in an antibiotic-free medium. To determine the number of cells per well, we detached cells using trypsin solution and counted cells in suspensions using a Bürker-Türk chamber.

## Azithromycin

We pre-dissolved 262.5 mg of purified azithromycin (Zithromax; 94.6% azithromycin dihydrate, a kind gift from Pfizer BV) in 5 mL of ultra pure water in the presence of 192.3 mg of sodium citrate (approximate pH 3.0). We also prepared a stock solution of 25 mg/mL azithromycin by adding 99.15 mg NaOH and ultrapure water for a final volume of 10 mL (approximate final pH 6.3). We stored this solution at -20 °C until it was needed

#### MIC/MBC

We determined the MIC and MBC of azithromycin for *C. pneumoniae* in HepG2 cells against a  $100*TCID_{50}$  dose of *C pneumoniae*. MIC and MBC were calculated on the basis of immunofluorescence for *C. pneumoniae* on the HepG2 cells.

# **Infection experiments**

Twenty-four hours before inoculation, we refreshed the cells and pre-treated them with 10% EMEM supplemented with 1, 0.125, 0.016 or 0 µg/mL azithromycin. Immediately before inoculation, we performed this procedure again and added 0.4 mL of fresh medium to each well. Cells were infected with 0.1 mL of 2 or 3 doses from both the *C. pneumoniae* AR39 strain and the CMV strain (MOI 0.6, 0.06, and 0.006, respectively). To the uninfected cells we added 0.1 mL of EMEM with the corresponding azithromycin concentration. IL-1 (100 pg/mL for IL-6 experiments) or IL-6 (1 ng/mL for fibrinogen and PAI-1 experiments) was added to the positive control wells. We conducted all incubations 3 times in duplicate. After inoculation, the cells were centrifuged for 1 hour at 800g and incubated for 24, 48, 72, or 96 hours. At each time point, we harvested supernatants and froze them at -35 °C until measurements could be taken. We verified infection rates by immunostaining the infected cells with antibodies to *C. pneumoniae* (30701 Pathfinder *Chlamydia* Culture Confirmation System, BioRad, Hercules, Calif) and CMV (anti-CMV immediate early antigen clone E13, no.12-003; Argene, Varilhes, France).

# **Cytokine incubations**

To investigate the anti-inflammatory effects of azithromycin, we incubated HepG2 cells with panels of IL-1 and IL-6 (200-01B and 200-06; Peprotech Inc, Rocky Hill, NJ) in low to (sub)maximal stimulating concentrations of these cytokines (0, 10 or 100 pg/mL; 1, 10, or 50 ng/mL). Cells were pre-treated in the same fashion as those used in the infection experiments.

# **APR** assays

We conducted sandwich Elisa's for Fibrinogen (FG-EIA Kordia; Affinity Biologicals, Inc, Leiden, the Netherlands), PAI-1 (Imulyse PAI-1 no. 211000; Kordia/ Biopool), and IL-6 (Pelikine IL-6 assay M1916; CLB, Amsterdam, the Netherlands) in accordance with the manufacturer's instructions. For fibrinogen, we constructed a standard curve from

normal haemostasis reference plasma (no. 50720; Kordia/Biopool). Standard curves for PAI-1 and IL-6 were supplied with the kits. For PAI-1, we used a standard fibrinolysis reference plasma (no. 210540; Kordia/Biopool) as a positive control.

# Statistical analysis

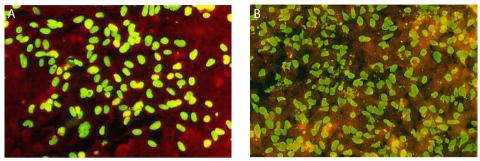
In all infection experiments we compared the production levels of infected hepatocytes with those of the negative controls (uninfected HepG2) at each time point. We analyzed data using Dunnett's multiple-comparisons g-test for differences between pairs of columns, which corrects for unequal variances. P values of less than 0.05 were taken to indicate a significant difference between production of IL-6, PAI-1, or fibrinogen by infected hepatocytes and that by uninfected hepatocytes. We also conducted analysis of linear trends in the IL-6 data to establish whether the effects of treatment were time- or dose-dependent. Linear regression was used to analyze the effects of C. pneumoniae-infection, time, and azithromycin on IL-6 production after transformation of the response variable on the log scale. On the other hand, our objective in the anti-inflammatory experiments was to assess whether azithromycin was capable of inhibiting production of IL-6, PAI-1 or fibring en by interleukin-stimulated hepatocytes, regardless of infection. For this reason it was necessary to compare the levels of production in all incubations with different concentrations of azithromycin and IL's among themselves. For this purpose we used Bonferroni's multiple-comparisons t-test. Beforehand, we performed ANOVA to rule out the possibility that any effect might have been due to a difference in variations instead of azithromycin alone. p-values greater than 0.05 were considered not significant.

#### Results

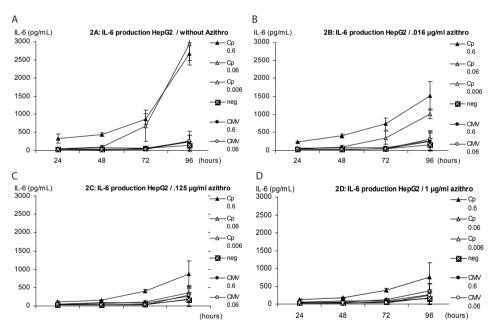
# C. pneumoniae and CMV infection in human hepatocytes

Infection rates of HepG2 cells were assessed by determining the TCID $_{50}$ -titres for CMV and *Cp* and the number of cells counted on the growth areas in each well of a 48-well TC-plate. TCID $_{50}$ -values of the virus stocks were  $10^{5.5}/50~\mu L$  and  $10^{4.5}/50~\mu L$  for *C. pneumoniae* and CMV, respectively, and  $10^5$  cells were counted on the growth surfaces. For *C. pneumoniae*,  $100~\mu L$  of a 10x, 100x and a 1000x dilution was inoculated on the HepG2 cells and for CMV  $100~\mu L$  of undiluted and 10x-diluted virus stock. Therefore actual MOI's were 0.6, 0.06, and 0.006 for *C. pneumoniae* and 0.6 and 0.06 for CMV (**Figure 1 and 2**). The MIC and MBC values for azithromycin on *C. pneumoniae*—infected

HepG2 cells were assessed at 0.063 and 0.125  $\mu$ g/mL, respectively. Hence, on the basis of these MIC and MBC values, we selected azithromycin concentrations of 0.016, 0.125, and 1  $\mu$ g /mL to be added to the culture medium in all experiments.



**Figure 1** Infection of HepG2 cells. **A**: Fluorescence staining of CMV-infected HepG2 cells. Note apple-green fluorescence of CMV immediate early antigen in the nuclei of infected cells. **B**: Fluorescence staining of C. pneumoniae–infected HepG2 cells. Apple-green fluorescence (genus-specific) denotes Chlamydia antigens in the cytoplasm of infected cells. Background of each photo contains uninfected cells counterstained with Evans blue (red). Original magnification 100x. (*For color figure see page 218*)

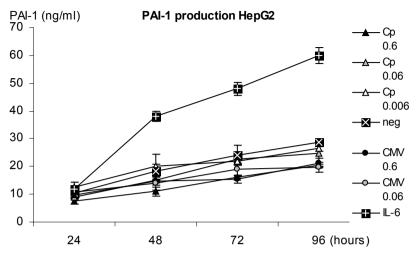


**Figure 2** IL-6 production by infected HepG2 cells. Cp = C pneumoniae; neg = uninfected HepG2 cells. Values of 0.6, 0.06, and 0.006 for C pneumoniae and CMV refer to the MOI. **A**, Incubations without azithromycin. **B**, HepG2 cells incubated with 0.016 µg/mL azithromycin. **C**, HepG2 cells incubated with 0.125 µg/mL azithromycin. **D**, HepG2 cells incubated with 1 µg/mL azithromycin. Y-axis denotes IL-6 levels in picograms per milliliter (mean  $\pm$  SEM); x-axis denotes incubation time in hours. All data are from 3 experiments performed in duplicate.

# IL-6, PAI-1, and fibrinogen production by hepatocytes after infection and effects of azithromycin.

The basal secretion of IL-6 by uninfected hepatocytes ranged from  $25 \pm 5$  pg/mL after 24 hours to 137 ± 120 pg/mL after 96 hours. Immediately after inoculation, no IL-6 (<20 pg/mL) was detected in the supernatants (data not shown). CMV infection did not induce IL-6 production by HepG2 cells at any point. C. pneumoniae infection with the highest dose of *C. pneumoniae* resulted in IL-6 production at all time points up to 2667 ± 309 pg/mL IL-6 after 96 hours, whereas infection with a lower concentration of C. pneumoniae did not induce IL-6 production until 96 hours had elapsed (2930 ± 447 pg/mL IL-6). After 96 hours of high-dose C. pneumoniae infection, 0.125 and 1 μg/mL, azithromycin suppressed IL-6 production (871±364 and 752±403 pg/mL, respectively) although azithromycin did not entirely reduce IL-6 production to the level of the uninfected HepG2 cells. C. pneumoniae-infected HepG2 cells incubated without azithromycin and with 0.016  $\mu$ g/mL azithromycin still produced IL-6 (2667  $\pm$ 309 and 1516  $\pm$  402 pg/mL, respectively (**Figure 2, A–D**). To substantiate the effects of C. pneumoniae on IL-6 production and the inhibitory effects of azithromycin, we subjected IL-6 responses to analysis for linear trends. After log transformation of the data, analysis for linear trends in a regression model showed dose-dependent effects on IL-6 production by C. pneumoniae doses with MOI's of 0.6 and 0.06 (p<0.001) but not for the lowest dose (MOI 0.006; p:0.39). The effects of time and azithromycin concentration on IL-6 production were also significant (p<0.001). After azithromycin treatment, C. pneumoniae-induced IL-6 production was no longer significant (p-values 0.23, 0.21, and 0.32, respectively). ANOVA revealed significant interaction of C. pneumoniae with time (p<0.001), indicating a time dependent effect of *C. pneumoniae* infection on IL-6 production. Interaction of *C. pneumoniae* with azithromycin was also significant (p:0.02) establishing the inhibitory effect of azithromycin on *C. pneumoniae*–induced IL-6 production. The interaction of azithromycin with time was not significant (p:0.20), indicating that azithromycin did not have additional effects on log IL-6 responses over time.

Although PAI-1 levels gradually increased over time, PAI-1 production by hepatocytes was not influenced by infection (**Figure 3**). Average PAI-1 production ranged from 10.3  $\pm$  1.7, 7.7  $\pm$  1.2 and 9.7  $\pm$  1.2 ng/mL after 24 hours to 28.7  $\pm$  0.5, 20.3  $\pm$  0.5 ng/mL and 21.3  $\pm$  1.7 ng/mL after 96 hours for uninfected cells, *C pneumoniae*–infected cells and CMV-infected cells, respectively (all p>0.05). Production of PAI by IL-6 stimulated hepatocytes was quite significant (p<0.01), ranging from 12 ng/mL at 24 hours to 60 ng/

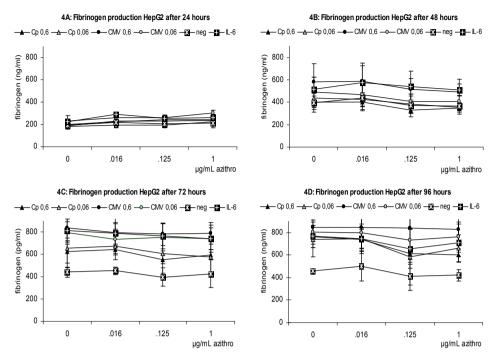


**Figure 3** PAI-1 production by infected HepG2 cells. *Cp: C. pneumoniae; neg:* uninfected HepG2 cells; *IL-6:* HepG2 cells incubated with 1 ng/mL IL-6 (positive control). Values of 0.6. 0.06, and 0.006 for *C. pneumoniae* and CMV refer to the MOI. *Y-axis* denotes PAI levels in nanograms per milliliter (mean±SEM); *x-axis:* incubation time in hours. All data are from 3 experiments performed in duplicate.

mL at 96 hours. Incubation with azithromycin did not alter these findings (p>0.05).

# HepG2 cells constitutionally produce fibrinogen

After 24 and 48 hours, we found no significant differences among fibrinogen levels in infected cells, uninfected cells, and IL-6–stimulated cells. Seventy-two hours stimulation with IL-6 and both CMV doses induced significant increases in fibrinogen production by hepatocytes compared with that in the negative control (838  $\pm$  88 and 794  $\pm$  98 ng/mL for both CMV concentrations (p<0.01) and 818  $\pm$  103 ng/mL (p:0.02) for IL-6, compared with 442  $\pm$  44 ng/mL for the negative control. After 96 hours, both *C. pneumoniae* and CMV infections induced significant fibrinogen production: 765  $\pm$  100 and 741  $\pm$  155 ng/mL (both p<0.05) for *C. pneumoniae* and 846  $\pm$  123 ng/mL (p<0.01) and 804  $\pm$  106 ng/mL (p<0.05) for CMV, compared with the negative control (460  $\pm$  16 ng/mL). Ninety-six hours incubation with IL-6 induced 766  $\pm$  99 ng/mL fibrinogen (p:0.03). Fibrinogen production after 72 (CMV) and 96 hours (CMV and *C. pneumoniae*) was not changed by the addition of 0.016, 0.125, or 1 µg/mL azithromycin, indicating that azithromycin did not suppress fibrinogen production (Bonferroni tests, all p>0.05). ANOVA showed no significant differences in variance among these groups (all p>0.05;



**Figure 4** Fibrinogen production by infected HepG2 cells. *Cp:* C pneumoniae; *neg:* uninfected HepG2 cells; *IL-6:* HepG2 cells incubated with 1 ng/mL IL-6 (positive control). Values of 0.6 and 0.06 for *C pneumoniae* and CMV refer to the MOI. *Y-axis* denotes fibrinogen levels in nanograms per milliliter (mean  $\pm$  SEM); *X-axis* denotes azithromycin concentrations (micrograms per milliliter) added to the incubations. All data are from 3 experiments performed in duplicate. **A:** Fibrinogen production by HepG2 cells after 24 hours. **B:** Fibrinogen production by HepG2 cells after 72 hours. **D:** Fibrinogen production by HepG2 cells after 96 hours.

# Figure 4, A-D).

# Anti-inflammatory effects of azithromycin

We also investigated the possibility of anti-inflammatory action on the part of azithromycin by performing cross-titrations of azithromycin against dilution panels (0, 10, and 100 pg/mL; 1, 10, and 50 ng/mL) of both IL-1  $\beta$  and IL-6. Hepatocytes did not produce significant concentrations of IL-6 at any point during incubation with a low concentration of IL-1 $\beta$  (10 pg/mL). Incubation with 100 pg/mL IL-1 $\beta$ , however, had induced IL-6 concentrations of as much as 1360 pg/mL after 96 hours, and the higher IL-1 $\beta$  concentrations of 1, 10, and 50 ng/mL had induced maximal IL-6 levels of 2299,

Table 1 Anti-inflammatory effects of azithromyci

Azithromycine						
dosage						
(hg/mL)	Unstimulated	10 pg/mL IL	100 pg/mLIL	1 ng/mL IL	10 ng/mL IL	50 ng/mL IL
Fibrinogen (ng/mL)	(7)					
0	616 (100%)	780 (100%)	758 (100%)	781 (100%)	761 (100%)	775 (100%)
.016	615 (100%)	777 (100%)	739 ( 97%)	740 (95%)	734 ( 96%)	(%66 ) 992
.125	577(94%)	744 (95%)	787 (104%)	757 ( 97%)	720 (95%)	728 (94%)
_	621 (101%)	100 ( 95%)	753 ( 99%)	748 (96%)	752 ( 99%)	745 (96%)
IL-6 (pg/mL)						
0	138 (100%)	268 (100%)	1360 (100%)	2299 (100%)	2617 (100%)	3403 (100%)
.016	129 ( 94%)	257 ( 96%)	1261 (93%)	2301 (100%)	2370 (91%)	3416 (100%)
.125	118 (86%)	257 ( 96%)	1278 ( 94%)	2312 (101%)	2656 (101%)	3609 (106%)
_	126 (91%)	264 (98%)	1226 (90%)	2411 (105%)	2581 (99%)	3618 (106%)
PAI-1 (ng/mL)						
0	32.7 (100%)	37.7 (100%)	54.3 (100%)	55.3 (100%)	25.6 (100%)	39.2 (100%)
.016	33.3 (102%)	35.0 (93%)	50.4 (93%)	51.1 ( 92%)	18.3 (71%)	28.7 (73%)
.125	28.1 (86%)	27.5 (73%)	43.7 (81%)	51.4 ( 93%)	22.3 (87%)	34.3 (88%)
_	35.0 (107%)	31.9 (85%)	52.5 ( 97%)	55.4 (100%)	25.2 ( 98%)	31.4 (80%)
Overview of fibring	en and PAI-1 production	n hv II -6 stimulated hena	Overview of fibrinonen and DAL-1 production by II-6 stimulated benatocutes incubated with 1-1 and incubates	zithromycin For II -6 hens	atocytes were stimulated	with II -1 and incubate

Overview of fibrinogen and PAI-1 production by IL-6 stimulated hepatocytes incubated with azithromycin. For IL-6 hepatocytes were stimulated with IL-1 and incubated with azithromycin. IL: dilution series of IL-1 or IL-6 from 0 (unstimulated) – 50 ng/ml. Unstimulated: HepG2 cells in medium without azithromycin and IL-1 or IL-6 (negative control). First row: azithromycin concentrations in µg/ml. Results are means of 3 experiments in duplicate and expressed in ng/ml (fibrinogen and PAI) or pg/ml (IL-6). Between brackets: each incubation without azithromycin (0 µg/ml Azithro) was set at 100%. For each azithromycin concentration expressions were related to expressions without Azithromycin incubation and expressed as a percentage of these values. 2617, and 3403 pg/mL, respectively, after 96 hours. Co-incubation with azithromycin did not affect IL-6 production by hepatocytes (Bonferroni tests, all p>0.05; **Table 1**).

# IL-6 is a stimulus for hepatocytes to produce PAI-1

Maximal PAI production in hepatocytes was induced by 100 and 1 ng/mL IL-6, resulting in PAI-1 levels of  $18.0 \pm 7.2$  ng/mL (p<0.05) and  $20.6 \pm 6.2$  ng/ml (p<0.05), respectively, after 24 hours, compared with  $8.6 \pm 3.5$  ng/mL for unstimulated hepatocytes. After 96 hours, PAI-1 levels had increased to  $54.3 \pm 7.5$  ng/mL (p<0.05) and  $55.3 \pm 6.1$  ng/mL (p<0.05) after incubation with 100 pg/mL and 1 ng/mL IL-6, respectively and to  $32.7 \pm 11.6$  ng/mL in unstimulated cells. Nevertheless, co-incubation with azithromycin had no effect on PAI levels (Bonferroni test, p >0.05; **Table 1**).

Basal production of fibrinogen by hepatocytes ranged from 466  $\pm$  31 ng/mL after 24 hours to 616  $\pm$  20 ng/mL after 96 hours. Stimulation of hepatocytes with IL-6 concentrations of up to 1 ng/ml slightly increased fibrinogen production from 584  $\pm$  19 ng/mL after 24 hours (p:0.03) to 664  $\pm$  42 ng/mL (p:0.04), 766  $\pm$  12 ng/mL (p<0.01), and 781  $\pm$  18 ng/mL (p<0.01) after 48, 72, and 96 hours, respectively. Co-incubation with azithromycin did not affect fibrinogen synthesis by HepG2 (Bonferroni multiple-comparisons tests, all p<0.05; **Table I**).

# **Discussion**

In this study we designed an *in vitro* model for infection of hepatoma cells and the effect of azithromycin in the outcome of these infections. We have shown that *in vitro* infection of hepatocytes with *C. pneumoniae* leads to time and dose-dependent production of IL-6 but only marginal production of fibrinogen. Co-incubation with azithromycin strongly inhibited this inflammatory response.

Now that inflammation is considered the underlying pathophysiological cause of atherosclerosis, the driving force behind chronic low-grade inflammation becomes a matter of question. The liver is the most important production site for inflammatory mediators such as IL-6 and CRP. IL-6 is the major regulator of hepatic CRP production and is an inducer of proatherogenic processes. We have previously shown that infection of liver cells contributes to the production of inflammatory proteins *in vitro*<sup>7</sup> possibly leading to increased plasma concentrations.

In vivo studies have revealed an anti-inflammatory effect of azithromycin, as measured on the basis of reduced plasma concentrations of IL-6 and CRP after treatment<sup>36,37</sup>.

In this study we showed that azithromycin reduces the inflammatory response of infected hepatocytes *in vitro*. We conclude that this reduction is a manifestation of an antimicrobial effect because a cytokine-induced inflammatory response could not be inhibited by azithromycin. Similarly, azithromycin did not affect IL-6 production by CMV-infected hepatocytes. The findings of several studies however, suggest anti-inflammatory potential for macrolides. Although the exact mechanisms seem quite heterogeneous and still must be elucidated, it is recognized that macrolides can prevent or inhibit the production of proinflammatory mediators independent of their antibacterial activities<sup>31,32,35</sup>. Whether these effects have therapeutic consequences in vivo is still under investigation.

The addition of 0.016 mg/L azithromycin to C pneumoniae-infected hepatocytes inhibited IL-6 production, but not entirely (p<0.05). Concentrations of 0.125 and 1 mg/L azithromycin reduced IL-6 production to almost the same level seen in the negative control (p>0.05). Nevertheless, even 1 µg/mL azithromycin was not sufficient to inhibit IL-6 production in the Cpneumoniae-infection experiment entirely, possibly indicating an alternative IL-6-stimulating mechanism insensitive to the action of azithromycin. In contrast to IL-6 production, fibrinogen production by hepatocytes could not be suppressed with azithromycin treatment. Taking into account that the inoculation materials did not contain IL-6 (<20 pg/mL; data not shown) and that we used azithromycin concentrations well above the MIC and MBC values for *C. pneumoniae*, this suggests that fibrinogen production by azithromycin-treated infections was induced by factors other than C. pneumoniae infection alone. It is possible that IL-6 produced by neighboring cells led to paracrine effects stimulating fibrinogen expression. These findings are supported by the results of other studies describing alternative pathways in fibrinogen synthesis. Northemann et al<sup>38</sup> suggested that different rat hepatoma lines each secrete different characteristic sets of hepatocyte stimulating factor activities and establish unambiquously that IL-6 is secreted by at least some of these lines. Another consideration is the fact that hepatoma cell lines may be quite heterogeneous and may contain distinct subsets of cells<sup>39</sup>. Zuraw and Lotz also described a striking phenomenon in studies of cultured HepG2 cells incubated with supernatants from previous HepG2 cultures. The stimulatory effect of these supernatants was neutralized by IL-6 antibodies, indicating that IL-6 may be active in the same culture in which it was produced, confirming its role as the major hepatocyte stimulating factor<sup>40</sup>. Paysant et al<sup>41</sup> used fibrinogen production in HepG2 cells as an index of IL-6 responsiveness; a concentration range of 1 to 5 ng/mL IL-6 was considered to yield the maximal response

of fibrinogen biosynthesis. In our experiments, hepatocytes still produced IL-6 levels approaching this range (~800 pg/mL after 72 and 96 hours), even in the presence of azithromycin. This IL-6 activity in our cultures may represent an alternative pathway for fibrinogen production, accounting for the failure of azithromycin to inhibit fibrinogen production. Apparently this is due to underlying processes other than infection alone.

On the basis of the concept of atherosclerosis being an inflammatory disease, a role of increasing importance has been assigned to *C. pneumoniae*<sup>3,42</sup>. Increasing evidence for a possible causal role of *C. pneumoniae* in atherogenesis prompted studies with macrolide antibiotics in patients at high risk for cardiovascular complications<sup>43</sup>. Several recent clinical studies, including WIZARD<sup>44</sup>, ACES<sup>45</sup>, AZACS<sup>46</sup>, and ACADEMIC<sup>47</sup>, in addition to animal studies<sup>33,48</sup> have addressed the properties and possible benefits of azithromycin in particular, as well as other macrolides<sup>49,50</sup>. Although some investigators foresee improvement in clinical outcome after macrolide treatment 33,43,49,51 most of these trials demonstrate no significant or only marginally beneficial effects of azithromycin in the treatment of coronary-artery disease. Recently the limitations of these studies were reviewed extensively by Grayston<sup>52</sup>. From the findings of 12 published and 2 ongoing clinical studies of antibiotic treatment for the secondary prevention of atherosclerotic cardiovascular disease, it was concluded that a "substantial number of these studies were performed on an inadequate number of subjects and length of observation to provide convincing results". Treatment of C. pneumoniae for just 7 days was considered entirely inadequate for a chronic infection. Unsuccessful treatment can also be explained from the life cycle of C. pneumoniae. Elementary bodies of C. pneumoniae may be unsusceptible to antibiotics, and the initially susceptible reticulate bodies may be able to enter a persistent phase<sup>52</sup>. This concept was also supported by Gieffers et al<sup>53</sup>. In this study we confirm the antibacterial properties of azithromycin on C. pneumoniae. We could not show direct anti-inflammatory properties of this drug in vitro.

#### **Acknowledgements**

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Azithromycin reduces

Chlamydia pneumoniaeinduced attenuation of
eNOS and cGMP production
by endothelial cells

# **Summary**

#### **Background**

Intracellular infections with cytomegalovirus (CMV) or *Chlamydia pneumoniae* (*Cp*) may play a role in the etiology of atherosclerosis. Nitric oxide (NO) is a key regulator of endothelial function. Under pathological conditions uncoupling of endothelial nitric oxide synthase (eNOS) leads to vessel damage as a result of production of oxygen radicals instead of NO. We hypothesized that infection-induced atherosclerosis is initiated by changes in NO metabolism and may be reversed by azithromycin treatment.

#### Methods

Confluent human umbilical vein endothelial cells (HUVECs) were infected with *Cp* or CMV. After 48 h of infection, production of eNOS, cyclic guanosine monophosphate (cGMP) and reactive oxygen species (ROS) was measured. Detection of cGMP was used as a reporter assay for the bioavailability of NO. Subsequently, *Cp*- and CMV-infected HUVECs were co-incubated with 0.016 mg/Land 1 mg/L azithromycin.

#### Results

Infection with Cp (MOI 1 and MOI 0.1) and CMV (MOI 1) caused a dose- and time-dependent reduction of eNOS production in the HUVECs: Cp MOI 1: 1141  $\pm$  74 pg/mL (p< 0.01); Cp MOI 0.1: 3189  $\pm$  30 pg/mL (p< 0.01); CMV: 3213  $\pm$  11 pg/mL (p< 0.01) vs. 3868  $\pm$  83 pg/mL for uninfected HUVECs. *Chlamydia pneumoniae*- but not CMV-infection also reduced cGMP-production (Cp: 0.195  $\pm$  0.030 pmol/mL (p< 0.01); CMV: 0.371  $\pm$  0.027 pmol/L (p>0.05) vs. 0.378  $\pm$  0.019 pmol/mL for uninfected HUVECs. CMV-infection did not affect ROS production either, but Cp-infection reduced ROS-production by 21% (p>0.05; Cp MOI 0.1) to 68% (p 0.01; Cp MOI 1). Azithromycin treatment restored Cp-induced eNOS, cGMP and ROS production in a dose-dependent manner.

#### Conclusions

Infection with *Cp* in endothelial cells *in vitro* attenuates eNOS, cGMP and ROS production in HUVECs and azithromycin reverses *Cp*-induced effects on eNOS, cGMP and ROS-production. The results from our *in vitro* research support the role of antibiotic therapy for infection-induced atherosclerosis by indicating that azithromycin does actually improve endothelial function.

## Introduction

Chronic infection may contribute to the pathogenesis of atherosclerosis. The major organisms that have been studied are *Chlamydia pneumoniae* (*Cp*) and cytomegalovirus (CMV)<sup>1-3</sup>. Chronic infection could act by a number of mechanisms including direct vascular injury and induction of a systemic inflammatory state; both leading to endothelial dysfunction<sup>4,5</sup>. Early evidence for involvement of *Cp* in atherogenesis came from sero-epidemiological studies<sup>6</sup> but also from animal models and histopathological studies<sup>7,8</sup>. Incidentally, even viable Cp was isolated from atherosclerotic vascular tissue<sup>9,10</sup>. Nevertheless, the relationship between *Cp* and atherosclerosis is still under debate. It now appears that some of the inconsistency of results from study to study may be owing, in part, to a lack of standardized methods. In addition to the lack of consistent serologic criteria, recent evaluations have demonstrated inherent problems with the performance of the most widely used serologic methods. Most importantly, there is no gold standard nor a reliable serologic marker for chronic or persistent *Cp* infection<sup>11-13</sup>. Plasma concentrations of CMV antibodies in patients with coronary disease are higher compared with control subjects<sup>14,15</sup> and those with carotid intimal thickening (a measure of subclinical atherosclerosis)<sup>16</sup>. Patients with antibodies directed against CMV more frequently have endothelial dysfunction and impaired responses to NO<sup>17</sup>. In addition, CMV antigens, nucleic acid sequences and DNA have been detected in smooth muscle cells of carotid artery plaques obtained from patients undergoing endarterectomy<sup>18</sup>. There is also an association between antibodies to CMV and elevated markers of inflammation, including C-reactive protein and interleukin-6<sup>19,20</sup>. Impaired endothelial function, induced by vascular risk factors such as dyslipidemia, hypertension and diabetes and by factors such as oxidative stress and inflammation, is a key process in the initiation and progression of atherosclerosis. Impaired vasoreactivity has already been documented in asymptomatic young adults with a family history of premature coronary disease but no other risk factors<sup>21</sup>. Nitric oxide (NO) is a multifunctional molecule with effects on vessel dilatation, inhibition of platelet adhesion/aggregation and monocyte adhesion to the vessel wall. In this way NO has anti-atherosclerotic properties<sup>22</sup>. Nitric oxide activates guanylate cyclase, resulting in an increase in the concentration of intracellular cGMP. Cyclic GMP is a second messenger, mediating the vasodilatating effects of NO. Under normal conditions, NO is synthesized in endothelial cells from L-arginin in a reaction catalyzed by endothelial nitric oxide synthase (eNOS). Under the influence of cardiovascular risk factors this process can be disturbed, for instance by suppression or uncoupling of eNOS leading to vessel damage as a result of production of oxygen radicals instead of NO. Excessive formation of reactive oxygen species (ROS) leads to inflammation, endothelial barrier dysfunction and increased adhesion for leucocytes<sup>23</sup>. In the present study we investigated the effects of intracellular infections with CMV or *Cp* on the production of cGMP, eNOS and ROS by human endothelial cells. In addition, the effects of azithromycin on infection induced endothelial dysfunction were also studied.

# **Methods**

#### **Endothelial cell culture**

Human umbilical vein endothelial cells (HUVEC) were harvested from umbilical veins by canulating the veins at both ends. Subsequently, veins were rinsed with warm phosphate-buffered saline solution. Then warm trypsin/EDTA solution (trypsin 0.05%/ EDTA 0.02%, Life Technologies Inc, Paisley, UK) was injected in the vein and incubated for 30 min at 37 °C. After gently massaging the vein, the cell suspension was collected in a 50 mL centrifugation tube and 10% fetal calf serum was added for neutralization. Cells were spun for 10 min at 500g. The pellet was resuspended in endothelial growth medium, consisting of endothelial basal medium (CC-3121, Clonetics, Cambrex Corporation, East Rutherford, NJ) and an EGM Bullet kit (CC-4133, Clonetics). The obtained cell suspension was brought into a 75 cm<sup>2</sup> tissue culture flask (3376, Costar<sup>®</sup>, Corning Inc., Corning, NY), which was precoated with fibronectin-solution (1:200 in PBS, a generous gift from M. Mul, CLB, Amsterdam, the Netherlands). Cells were cultured in a humidified CO<sub>2</sub>-incubator at 37 °C and 5% CO<sub>2</sub>. At confluence the cells were detached with trypsin-EDTA solution and seeded into new tissue culture flasks (split ratio 1:2-1:3, passages 1 and 2) or 48-well tissue culture plates (passage 3) precoated with fibronectin. Cell viability was regularly checked by trypan blue staining. For all experiments cultures of passage 3 were used unless otherwise stated. Human umbilical vein endothelial cells were identified by immunostaining with antibodies against factor VIII-related antigen (anti-von Willebrand, M0616 DAKO, Glostrup, Denmark).

# **Preparation of Cp- and CMV-stock**

Human embryonic lung (HEL) and Buffalo Green Monkey cells (BGM) were cultured at  $37 \, ^{\circ}$ C and  $5\% \, CO_2$  in Minimal Essential Medium Eagle with Earle's Salts (EMEM, Gibco) with 10% fetal calf serum (Gibco). This culture medium was supplemented with  $2 \, \text{mM}$ 

L-glutamine (Gibco), 5 mL non-essential amino acids (Gibco), 10 mg/Lvancomycin (Faulding Pharmaceuticals, Brussels, Belgium), 4 mg/L amphothericin B (Fungizone; Bristol-Meyers Squibb, Woerden, the Netherlands) and 10 mg/L gentamicin (Schering Plough, Maarssen, the Netherlands). The same supplements were also added to the media used for the propagation of the virus strains. A clinical isolate of CMV was propagated in HEL cells with minimal essential medium containing 2% FBS, 20 mM Hepes and supplements. At > 80% cytopathologic effect, CMV-infected HEL cells were detached with trypsin/EDTA solution (Gibco) and centrifuged. The cell pellet was resuspended in the same medium containing 2% fetal calf serum and 10% DMSO (Sigma-Aldrich, Zwijndrecht the Netherlands) and supplements. Cp-strain AR 139 was propagated in BGM cells at 37 °C/5% CO<sub>3</sub> in EMEM with 10% fetal calf serum and 0.1% cycloheximide (Sigma-Aldrich Zwijndrecht, the Netherlands) and supplements. After 72 h of growth, infected cells were frozen and thawed to release the elementary bodies. After a short centrifugation step, cell debris was discarded and 0.2M SPG-medium (2.088 g/L K<sub>2</sub>HPO<sub>4</sub>, 1.088 g/L KH<sub>2</sub>PO<sub>4</sub>, 68.46 g/L saccharose, 7.16 g/L L-glutamine, 10% FBS, 2.5 mg/L amphothericin B, 23 mg/L vancomycin and 18 mg/L gentamicin) was added (1:1 v/v). Chlamydia pneumoniae and CMV stock suspensions were aliquoted and stored at -80 °C until further use. Also, aliquots of inactivated CMV and Cp were made by overnight UV-inactivation. The 50% tissue culture infective doses (TCID<sub>so</sub>) of live virus stocks were calculated using the method of Reed & Muench<sup>24</sup>. For calculation of TCID<sub>50</sub>-titres for CMV the HEL cells were examined daily for cytopathologic effects during 1 week. The TCID<sub>50</sub>-titre of the Cp stock was calculated otherwise, based on the number of immunofluorescent units per field in the infected BGM cells after staining with Chlamydia culture confirmation monoclonal antibodies (de Beer, Diessen, the Netherlands). Infection concentrations were chosen to mimic chronic low-grade infection (MOI 1 and 0.1; Figure 1).

# MIC and MBC concentrations of azithromycin

Minimal inhibitory (MIC) and bactericidal (MBC) concentrations of azithromycin concentrations in Cp-infected HUVECs cells were calculated with minor changes according to the protocol of Suchland et al<sup>25</sup>. Shortly, a twofold dilution series with final concentrations of azithromycin from 4.096 mg/L to 0.008 mg/L of azithromycin in antibiotic-free medium was made and brought onto HUVEC monolayers. The cells were infected with a  $100x \, \text{TCID}_{50}$  dose of Cp and incubated for 48 h. The azithromycin concentration in which no or morphologically changed inclusions were formed was recorded as the MIC-value. For MBC assessment, all cells were freeze-thawed and passed

onto new HUVEC monolayers. After 48 h the MBC-value was read as the concentration of azithromycin in which no inclusions were formed. Accumulation of azithromycin, associated with high tissue retention and a half-life time of 50-90 h, results in tissue concentrations up to 100-fold higher than plasma levels. Mean peak plasma levels after a 500mg dose of azithromycin approximate 0.40–0.45 mg/L after 2.5 h <sup>26</sup>. Azithromycin concentrations in tissue usually peak between 48 and 96 h after administration. Baldwin et al. determined mean peak tissue and serum concentrations after 48 h in bronchial epithelial lining fluid (1.56 mg/L vs. 0.03 mg/L; ratio 52:1) and bronchoalveolar macrophages (23 mg/L vs. 0.02 mg/L; ratio 1150:1) <sup>27</sup>. In 2000, Schneider et al. described the in vivo uptake of azithromycin in human coronary plaques. The median concentration in all plaques was 0.284 mg/L (Cl: 0.163-517 mg/L; n=10)<sup>28</sup>. Recently, Reveneau et al. described the superior bactericidal effect of azithromycin on persistent chlamydial infection in interferon gamma-treated epithelial cells, which was correlated with an enhanced uptake of the drug<sup>29</sup>. Azithromycin concentrations found in these studies are within the in vitro susceptibility range. In general, MICs and MBCs of azithromycin for *Cp* infections range from  $\leq$  0.03 to 1 mg/L<sup>30-32</sup>. Hence, in this study we chose to perform neutralization experiments with 0.016 mg/L (MIC) and 1 mg/L azithromycin doses.

#### **Set-up of experiments**

Based on the outcome of the first eNOS and cGMP experiments (Figs 2 and 3) incubation times of 48 h were chosen for all (neutralization) experiments. In previous optimization experiments the influence of the HUVEC passage number on eNOS production was only marginal, in contrast to cGMP production for which cells in passage 3 produced 50%-90% greater cGMP than in previous or higher passages, indicating an optimum production in passage 3 cells (data not shown).

# Inoculation of endothelial cells

For all experiments HUVEC cells were subcultured at passage 3 in 48-well plates (Costar\*Corning Inc.) until confluence was reached. All experiments were performed at least three times in duplicate on cells obtained from different donors. One day before inoculation, cells were refreshed with endothelial growth medium.

Purified azithromycin 262.5 mg (94.6% azithromycin dihydrate; a kind gift from Pfizer BV.) was predissolved in 5 mL ultra pure water in the presence of 192.3 mg sodium

citrate (pH approx. 3.0). A stock solution of 25 mg/mL azithromycin was prepared further by adding 99.15 mg NaOH and ultra pure water to a final volume of 10 mL (final pH approximately 6.3). This stock solution was stored at -20 °C until further use. For neutralization experiments, concentrations of 1 mg/L and 0.016 mg/L azithromycin were added to the endothelial growth medium. Chlamydia pneumoniae and CMV were prediluted in endothelial growth medium and added at a multiplicity of infectious units per cell (MOI) of 1 and 0.1 for both Cp and CMV. Uninfected cells and inactivated virus were used as negative controls. After inoculation, cell cultures were centrifuged for 1 h at 800 g and re-fed with the corresponding culture medium. 48 hours after infection, production of eNOS, cGMP and ROS was measured. All infections were monitored by immunostaining for either *Cp* or CMV (Fig 1b-d). Cell-free supernatants were collected and frozen at -80 °C for cytotoxicity testing. For eNOS and cGMP testing, cells were washed with PBS. For cell lysis 200 µL of lysis buffer (eNOS-assay) or 200 µL of 0.1 M HCl (cGMP-assay) was added to each well. After 30 min on an orbital shaker the cells were inspected microscopically to verify complete cell lysis. Lysates were stored at -80 °C until the assay was scheduled. ROS-detection was performed immediately.

# **Immunoassays**

In order to obtain valid results for eNOS, cGMP and ROS production it was mandatory to assess the extent of cytotoxicity beforehand. The viability of the HUVECs cells was measured using a LDH activity assay (cytotoxicity detection kit 1644793, Roche Applied Science, Germany). For eNOS detection a quantitative solid-phase sandwich enzyme immunoassay (Quantikine Human eNOS assay, DENO0, R & D Systems Europe Ltd., Abingdon, UK) was used, and cGMP was measured in a competitive immunoassay (Correlate-EIA #900-014, Assay Design Inc., Ann Arbor, MI).

For ROS measurement HUVEC were grown in 96-microwell plates (Nunclon 167008, NUNC A/S, Roskilde, Denmark). Cells were washed once with 100  $\mu$ L PBS+++ (1 mM CaCl2, 0.5 mM MgCl<sub>2</sub>, 0.1% (w/v) glucose in phosphate-buffered saline). Cells were loaded with 75  $\mu$ L of 10  $\mu$ M dichlorodihydrofluorescein diacetate probe (CM-H2DCFDA; cat nr C6827, Molecular Probes Europe, Leiden, the Netherlands) for 30 min at 37 °C in the dark. The DCF-probe is a cell-permeant indicator for reactive oxygen species that is nonfluorescent until the acetate groups are removed by intracellular esterases and oxidation occurs within the cell. Cells were washed twice with 100  $\mu$ L PBS+++ and 190  $\mu$ L PBS+++ was added to each well. Subsequently, 10  $\mu$ L 20 mM H<sub>2</sub>O<sub>2</sub> (final concentration 1 mM) or 10  $\mu$ L 2 mM Angiotensin II (final concentration 100  $\mu$ M) was

added. Fluorescence (DCF-signal) was read every 5 min during 70 min at 485 nm excitation and 538 nm emission. ROS-values were expressed as fluorescent signals from the DCF-probe we used without an indication of units. In general, the signals would be low to such a degree that relating them to the signals of controls (Angiotensin and  $H_2O_2$ ) would have led to marginal values and that any effects, if present, might remain unnoted. Alternatively, incubation with inactivated Cp and CMV was considered to be the most accurate (negative) infection control and fluorescence values of these incubations were fixed at 100%. Subsequently, all other fluorescence values were related to this and expressed as a percentage of these controls to give an impression of the relative ROS-production after infection with or without azithromycin treatment.

# **Statistical analysis**

The GraphPad Instat™ software package version 2.05a (Graphpad Software Inc., San Diego, CA) was used for data analysis. After anova-analysis on variation between means, the production levels of infected HUVECs were compared with those of uninfected HUVEC cells (or cells incubated with inactivated virus) at each separate time point. Data were analyzed using Dunnett's multiple comparisons q-test (which corrects for possibly unequal variances) for differences between controls and incubations; p-values less than 0.05 indicate a significant difference between results. Analyses on linear trends were performed to warrant that the decrease of eNOS-production in time was owing to infection rather than merely a time effect. The null hypothesis assumes that there is no correlation between means and time points. Therefore, low p-values suggest a linear trend.

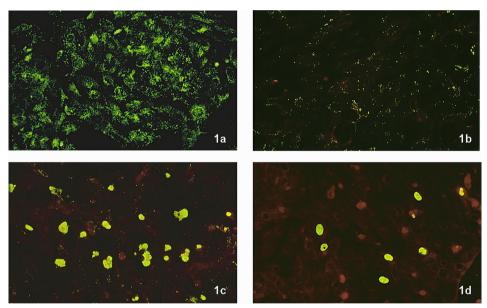
#### Results

# Chlamydia pneumoniae and Cytomegalovirus infection of endothelial cells

Log TCID<sub>50</sub>/mL-values of the virus- and Chlamydia stocks were calculated on 5.3 and 4.3 for Cp and CMV, respectively. At confluence the mean density of cells grown in 48-well tissue culture plates was  $4 \times 10^4$  per well. One hundred  $\mu$ L of each dilution (Cp:  $\times 10$  and  $\times 100$ ; CMV: undiluted and  $\times 10$ ) was inoculated on HUVECs, thus obtaining MOI values of 1 and 0.1 for both Cp and CMV. In addition, HUVEC cells of each donor were identified as such, by immunostaining on von Willebrand factor (**Figure 1a**). The minimal inhibitory (MIC) and bactericidal (MBC) concentrations of azithromycin in Cp-infected HUVEC cells were calculated to be 0.016 mg/L and 0.125 mg/L, respectively.

# **Cytotoxicity testing**

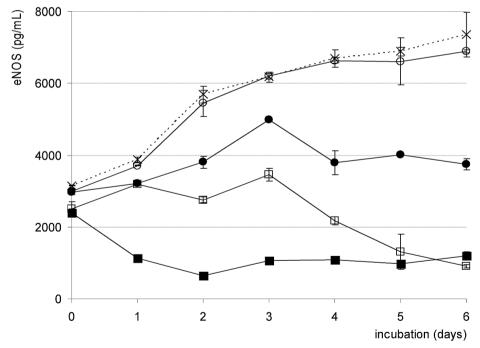
Human umbilical vein endothelial cell suspensions used for seeding the plates did not show any loss of viability (≥ 99% trypan blue exclusion); neither did HUVEC monolayers after visual examination by phase-contrast microscopy before infection (data not shown). In addition, infections were monitored by immunostaining. Care was taken to obtain low-grade infections with minimal cell damage (**Figure 1b-d**). Cell injury was determined by measuring LDH leakage in the supernatants. Relative cytotoxicity in uninfected cells ranged from 1.3 to 6.4%. Cytotoxicity in HUVEC cells incubated with inactive or infectious Cp and CMV was approximately 2-3-fold higher than in uninfected cells (data not shown).



**Figure 1**. Fluorescence microscopy images ( $\times$ 200) of human umbilical vein endothelial cells (HUVECs) 48 h after inoculation. Indirect immunostaining with antihuman von Willebrand factor (a), anti-Cp (b,c) and anti-CMV (d) antibodies. Secondary antibody: FITC-labelled rabbit antimouse conjugate; counterstaining: Evans Blue. Red background: uninfected cells. Apple green fluorescence: positive identification of von Willebrand factor (a), inactivated *Cp* (b), *Cp* (MOI = 1) (c) and CMV (MOI = 1) (d). (*For color figure see page 219*)

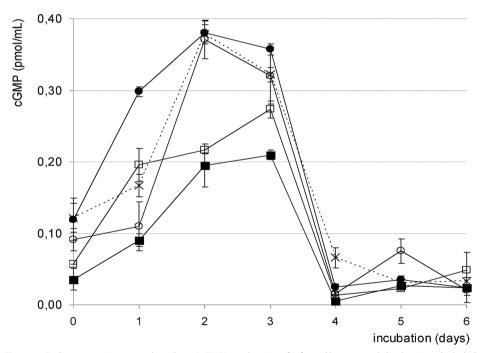
# eNOS and cGMP production after infection with Cp or CMV

In uninfected HUVECs, eNOS production gradually increased from 3156  $\pm$  19 pg/mL to 7350  $\pm$  621 pg/mL after 6 days, indicating a linear trend in eNOS production in time (p<0.0001). After 1 day eNOS production did not differ from time point zero, but as from 48 h eNOS production was significantly different from time point zero (Dunnet: p<0.01). Infection with the lowest CMV concentration (MOI 0.1) did not alter eNOS expression at any time point (Dunnett: all p>0.05) and increase of eNOS in time was still linear, also indicating that there was no effect of this infection on eNOS-production (p<0.001). On the other hand, incubation of 24 h or longer with *Cp* MOI 1, 0.1 and CMV MOI 1 suppressed eNOS expression significantly (1141  $\pm$  74 pg/mL, 3189  $\pm$  30 pg/mL and 3213  $\pm$  11 pg/mL after 24 h, respectively; Dunnett: all p<0.01) when compared with uninfected HUVECs (3868  $\pm$  83 pg/mL after 24 h). There was no longer a linear relation between eNOS and time (p>0.05), also indicating the suppressive effect of infection on eNOS production (**Figure 2**).



**Figure 2** eNOS expression in infected human umbilical vein endothelial cells (HUVECs). Means  $\pm$  standard error (SE) of eNOS expression in three experiments in duplicate (n = 3) and incubation times from 0 to 6 days. X-axis: incubation time in days; Y-axis: eNOS expression in pg/mL; – x -- uninfected HUVECs;  $\blacksquare$  infection with *Cp* MOI 1;  $\Box$  infection with *Cp* MOI 0.1; • infection with CMV MOI 0.1.

Cyclic guanosine monophosphate production in *Cp*-infected HUVECs did not differ from cGMP production in uninfected cells during the first 48 h of incubation (p>0.05). After 48 h, infection with both *Cp*-doses suppressed cGMP-production significantly (MOI 1: 0.195  $\pm$  0.030 pmol/mL, p< 0.01; MOI 0.1: 0.216  $\pm$  0.064 pmol mL, p<0.05) compared with uninfected HUVECs (0.378  $\pm$  0.019 pmol/mL). Cytomegalovirus infection did not alter cGMP-production at all (0.371  $\pm$  0.027 and 0.380  $\pm$  0.023 pmol mL, p>0.05). After 72 h, infection with *Cp* MOI 1 still gave a significant reduction of cGMP-production (0.210  $\pm$  0.099 vs. 0.322  $\pm$  0014 pmol/mL for uninfected cells; p<0.05) but after incubation times of 96 h and longer overall cGMP-production was decreased and was only marginal, indicating abolished cGMP-synthesis (**Figure 3**). Cyclic guanosine monophosphate production-synthesis in this culture model seems self-aborting, probably owing to exhaustion of one more components in the eNOS-NO-cGMP pathway. Consequently, an incubation time of 48 h was kept for all upcoming azithromycin experiments.



**Figure 3** Cyclic guanosine monophosphate (cGMP) production of infected human umbilical vein endothelial cells (HUVECs). Means  $\pm$  standard error (SE) of cGMP production in three experiments in duplicate (n = 3) and incubation times from 0 to 6 days. X-axis: incubation time in hours; Y-axis: cGMP production in pmol/mL; − x − uninfected HUVECs; ■ infection with *Cp* MOI 1; □ infection with *Cp* MOI 0.1; • infection with CMV MOI 1; ○ infection with CMV MOI 0.1.

# Effect of azithromycin on ROS production by infected HUVECs

After 48 h of incubation with infectious and inactivated CMV (MOI 1), ROS-production in HUVEC was unchanged compared with uninfected HUVEC (DCF signals for infectious CMV:  $1.128 \pm 0.117$  (p>0·05); inactivated CMV:  $0.863 \pm 0.266$  (p>0.05) vs.  $0.883 \pm 0.129$  for uninfected HUVEC.

In contrast, infectious Cp (MOI 1), but not inactivated Cp, suppressed ROS-production significantly (infectious Cp: 0.342  $\pm$  0.029; p<0.05, inactivated Cp: 1.080  $\pm$  0.19; p>0.05) compared with uninfected HUVECs (0.883  $\pm$  0.129), indicating a reduction of 61% in ROS-production.

Co-incubation with azithromycin restored relative ROSproduction of Cp-infected HUVECs (MOI 1) compared with HUVECs with inactivated Cp from 31.7% without azithromycin (342  $\pm$  29; p<0.01) to 46.2% with 0.016 mg/L azithromycin (480  $\pm$  24; p<0.05) and 83.4% with 1 mg/L azithromycin (723  $\pm$  54; p>0.05), indicating a dosedependent effect of azithromycin on ROS-production. Infection with Cp MOI 0.1 still reduced ROS-production by 21.2% compared with inactivated Cp-incubation (0.852 $\pm$ 0.176 vs. 1.080 $\pm$ 0.197 for inactivated Cp) but this reduction was not statistically significant (p>0.05). Although not significant, azithromycin was still able to enhance ROS-production of MOI 0.1 Cp-infected cells from 78.8% (DCF: 0.852 $\pm$ 0.176) to 91.5% for 0.016 mg/L azithromycin (DCF: 0.951 $\pm$ 0.060) and 96.4% for 1 mg/L azithromycin (DCF: 0.836 $\pm$ 0.060), respectively (**Figure 4**).

# Effect of azithromycin on eNOS-expression by infected HUVECs

After 48 h, eNOS-expression in HUVECs was 4970  $\pm$  128 pg/mL. Incubation with either infectious or inactivated CMV did not alter eNOS-expression (4940  $\pm$  222 pg/mL and 5010  $\pm$  262 pg/mL resp. both p>0.05; (**Figure 5**). *Chlamydia pneumoniae*-infection, however, did suppress eNOS-expression significantly. Infection with *Cp* MOI 1 and MOI 0.1 *Cp* reduced eNOS-expression to 2649  $\pm$  381 pg/mL (p< 0.001) and 4214  $\pm$  87 pg/mL (p<0.05), respectively. eNOS-expression after incubation with both inactivated *Cp* stocks was not different (p>0.05) from eNOS-expression in uninfected HUVECs (4425  $\pm$  298 pg/mL, p>0.05 for inactivated *Cp* MOI 1 and 4640  $\pm$  489 pg/mL (p>0.05) for inactivated Cp MOI 0.1 vs. 4970  $\pm$  128 pg/mL for uninfected HUVECs). Addition of azithromycin to the *Cp*-infected cells restored eNOS-expression. Co-incubation with 0.016 mg/L azithromycin increased eNOS-expression only moderately from 2649  $\pm$  381 pg/mL (p<0.001) to 3018  $\pm$  pg/mL for *Cp* MOI 1 (p<0.001) and from

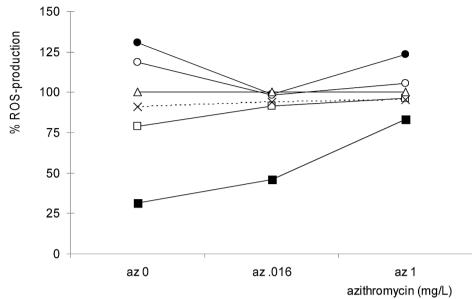


Figure 4: ROS production of infected human umbilical vein endothelial cells (HUVECs) and effect of azithromycin. Means  $\pm$  standard error (SE) of ROS production in three experiments in duplicate (n = 3); incubation time 48 h. ROS = reactive oxygen species; Cp = Chlamydia pneumoniae; CMV = cytomegalovirus; MOI = multiplification of infection. X-axis: azithromycin concentration (mg/L); Y-axis: relative ROS-production in percentage of inactivated virus control; incubations with inactivated Cp and inactivated CMV were both adjusted to 100% (control). ♦ inactivated Cp/CMV; -- x -- uninfected HUVECs; ■ infection with Cp MOI 1; □ infection with Cp MOI 0.1; • infection with CMV MOI 0.1.

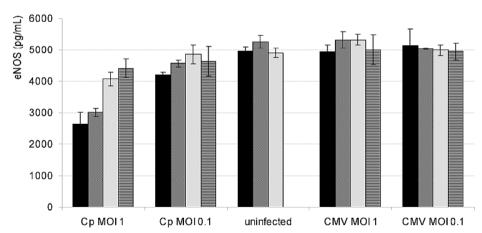
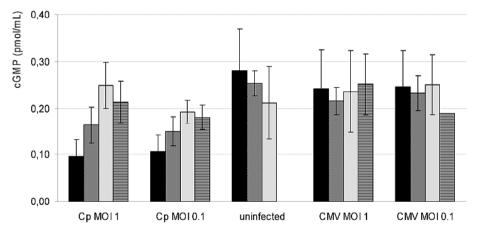


Figure 5. Effect of azithromycin on eNOS-expression in human umbilical vein endothelial cells (HUVECs). Means ± standard error (SE) of eNOS expression in three experiments in duplicate (n = 3); Incubation time 48 h; *Cp* = *Chlamydia pneumoniae*; CMV = cytomegalovirus; uninfected: uninfected HUVECs cells; MOI = multiplification of infection; Y-axis: eNOS expression in pg/mL; X-axis: incubation type; infection without azithromycin; 0·016 mg/L azithromycin; 1 mg/L azithromycin; incubation with inactivated Cp or CMV.

4214  $\pm$  87 pg/mL (p<0.05) to 4575  $\pm$  108 mg/L (p<0.05) for *Cp* MOI 0.1 infected HUVECs compared with azithromycin-treated uninfected HUVECs (5262  $\pm$  197 pg/mL). On the other hand, co-incubation with 1 mg/L azithromycin almost completely restored eNOS-expression to 4084  $\pm$  219 pg/mL (p>0.05) for *Cp* MOI and 4857  $\pm$  294 pg/mL (p>0.05) for 0.1 MOI when compared with azithromycin-treated uninfected HUVECs (4909  $\pm$  151 pg/mL) (**Figure 5**).

# Effect of azithromycin on cGMP-production by infected HUVECs

After 48 h, cGMP-expression in uninfected HUVECs cells was  $0.280 \pm 0.077$  pmol/mL. Addition of azithromycin did not alter cGMP-production of uninfected HUVECs. Initially, CMV-infection did not affect cGMP-production in HUVECs ( $0.241 \pm 0.089$  pmol/mL; p>0.05). Therefore, addition of azithromycin to CMV-infected HUVECs did not induce any changes in cGMP-production either. *Chlamydia pneumoniae*-infection however, did suppress cGMP-production ( $0.096 \pm 0.037$  pmol mL). Co-incubation of azithromycin (0.016 and 0.016 mg/L) with the *Cp*-infected HUVEC cells blunted cGMP-suppression ( $0.0164 \pm 0.038$  pmol/mL and 0.0016 mg/L and 0.0016 mg/L azithromycin, respectively). Although these results were not statistically significant, there was an upward trend in cGMP-production after co-incubation with azithromycin (**Figure 6**).



**Figure 6.** Effect of azithromycin on Cyclic guanosine monophosphate (cGMP)-production in human umbilical vein endothelial cells (HUVECs). Means  $\pm$  standard error (SE) of cGMP production in three experiments in duplicate (n = 3); incubation time 48 h; Cp = Chlamydia pneumoniae; CMV = cytomegalovirus; uninfected: uninfected HUVECs cells; MOI = multiplification of infection;

Y-axis: cGMP production in pmol/mL; X-axis: incubation type; ■ infection without azithromycin; ■ 0·016 mg/L azithromycin; ■ 1 mg/L azithromycin; ■ incubation with inactivated Cp or CMV.

## Discussion

Endothelial dysfunction plays a pivotal role in atherogenesis and development of vascular complications. Classical risk factors contribute to endothelial function, but also non classical risk factors like oxidation products and inflammatory mediators may lead to endothelial dysfunction. Also, several studies indicate a role for infections with *Cp* and CMV in the pathogenesis of atherosclerosis<sup>33-35</sup>. Infection of the endothelium causes continuous activation and loss of anti-atherogenic cellular mechanisms like a decrease in NO-bioavailability<sup>36,37</sup>. In the present study we investigated the effects of intracellular endothelial infection with CMV and *Cp* on NO-metabolism. Endothelial nitric oxide synthase, cGMP and ROS production were attenuated in *Cp*-infected HUVECs. CMV-infection did not induce any changes in cGMP- or ROS-production and only marginally decreased eNOS-production. Subsequent co-incubations with azithromycin restored eNOS, cGMP and ROS-production of *Cp*-infected HUVECs to levels comparable with those in uninfected HUVECs.

Inflammation is considered to be a major underlying mechanism contributing to endothelial dysfunction. Uncoupling of eNOS and subsequent vascular production of ROS may be the result of inflammatory mediators<sup>38</sup>. Reports about the relationship between CMV and Cp with atherosclerosis and endothelial dysfunction are guite inconsistent. Several studies show an association between Cp and CMV and endothelial dysfunction<sup>39,40</sup> and other studies do not<sup>41-43</sup>. In our study, CMV-infection of endothelial cells did not influence eNOS, cGMP or ROS-production, suggesting that CMV does not induce direct effects on vascular function by influencing eNOSdependent NO-production as determined by cGMP levels. We did not demonstrate involvement of CMV in decreased NO-production; this may indicate that involvement in atherogenesis is mediated by other mechanisms. Infection with CMV may lead to atherosclerosis by other mechanisms, including production of inflammatory cytokines in the liver, expression of endothelial adhesion molecules and production of tissue factor on endothelial cells<sup>44-46</sup>. Other investigators suggested that CMV elicits a general subclinical inflammatory response in certain individuals who are therefore susceptible to the development of atherosclerosis; those without an inflammatory response are expected to be resistant<sup>47</sup>.

In this study *Cp*-infection attenuated both eNOS- and cGMP-production in HUVECs significantly, indicating decreased NO-bioavailability. Inactivated *Cp* did not alter eNOS or cGMP-expression, suggesting that active infection is involved in reduced eNOS

production. Uncoupling of eNOS should lead to elevated production of ROS instead of NO<sup>48</sup>. Surprisingly enough, ROS-production by Cp-infected HUVECs was also reduced but this was not caused by cytotoxicity owing to infection. These findings are in contradiction with most studies in which Cp-infection generates ROS and subsequent activation of the NF-kappa B pathway, resulting in synthesis of pro-inflammatory cytokines and procoagulant activity. Our group previously demonstrated that coincubation of Cp-infected HUVECs with radical scavengers elevated IL-6 production, indicating NF-kappa B activation<sup>49</sup>. In the present infection-experiments eNOS may have become degraded, inactivated or uncoupled. Subsequently, in the tissue culture environment the dual action of eNOS may have resulted in production of NO as well as superoxide. The superoxide radical is frequently involved in oxidative inactivation of NO. Likely, enhanced degradation of NO occurs by reaction of NO with superoxide. This reaction kinetically exceeds other NO-reactions and leads to formation of peroxynitrite<sup>50</sup>. Nevertheless, the used DCF-probe should be sensitive to peroxynitrite as well as other radical oxygen species (e.g. superoxide, peroxynitrite, hydroxyl radical and hydrogen peroxide). However, in the process of degrading NO, ROS themselves might have been consumed. Addition of radical scavengers might reveal which radicals are involved or not, but seems unrealistic if ROS levels appear to be reduced. In addition, development of *Cp*-infection inside the cytoplasm of cultured HUVEC cells may have down-regulated cell metabolism, possibly leading to unstable eNOS and diminished NO and ROS-synthesis.

The suppressive effect of *Cp*-infection on eNOS-expression in human endothelial cells and loss of functionality leads to decreased NO-bioavailability and may trigger atherogenic processes. The results of the present *in vitro* study support the feasibility of an infectious etiology in atherogenesis and confirm a link between *Cp*-infection and endothelial dysfunction. In vivo this hypothesis is supported by clinical studies on the relation between inflammation and vascular function in patients with documented (coronary) artery disease. Impaired vasodilator function as measured by plethysmography demonstrated CRP to be an independent predictor for endothelial function <sup>51,52</sup>. Similar studies have shown that macrolide therapy effectively improved flow-mediated vasodilatation in *Cp*-seropositive patients<sup>53,54</sup>.

Endothelial nitric oxide synthase expression was measured in a protein assay, which does not provide any information about the functionality of the eNOS enzyme. Actual detection of NO provides information about eNOS activity. However, detection of NO *in vitro* is cumbersome. The NO molecule itself is a very reactive gas with short half-

life time and is present in endothelial cells only at a nanomolar level lying far beyond detection levels of current (commercial) detection methods. We acknowledge this as one of the potential pitfalls in our study. Alternative methods for detection of NO may be detection of degradation products like nitrate or nitrite; by-products like citruline or second messenger products like cGMP. Furthermore, the relationship between local NO concentration and the quanylate cyclase activity might be more complex than a straightforward 1:1 relation<sup>55</sup>. Moreover, cGMP formation might be influenced by other stimuli than NO. In the experimental setup it appeared that cell culture itself was of great influence, particularly on cGMP expression. Apart from individual variation in cell donors we found that cGMP expression was optimal in HUVEC cells of passage 3; in lower and higher passages cGMP levels were at least 50% lower (data not shown). Pooling of results obtained from cells in different passages would subdue cGMP measurement. As Figure 3 shows, cGMP production was completely abolished after 3 days. A plausible explanation for this phenomenon may be exhaustion of cofactors or substrate (L-arginin), required for NO-synthesis in the tissue culture environment, leading to uncoupling of eNOS. Preventing this from happening, all incubation times for the azithromycin experiments were set at 48 h. Finally, the low dynamic range of the cGMP-assay should be mentioned. With maximal absorbance values of 0.500 for this assay, small variations in culture and detection procedures may yield undesirable uncertainty in calculated cGMP levels.

At present, there are several therapeutic options for increasing the bioavailability of NO, aiming to improve endothelial function. Drugs like statins, ACE inhibitors and antioxidants seem to have beneficial effects on endothelial dysfunction. Sildenafil (Viagra\*) therapy has been shown to be a clinically effective treatment for erectile dysfunction. With respect to the possible role of infectious pathogens in atherosclerosis it has been suggested that antimicrobial therapy might be beneficial for vascular function and therefore protective for the development of atherosclerotic complications. Application and effects of quinolone- and neomacrolide antibiotics like roxithromycin and azithromycin on inflammation have been evaluated in prospective clinical trials<sup>56-59</sup>. Until today results from studies upon antibiotics, that might be beneficial in preventing CHD, remain inconclusive. As for azithromycin, some even tend to indicate that this drug is ineffective for secondary coronary heart disease prevention. Results from the WIZARD and ACES & PROVE IT trials showed no benefit of long-term antibiotic use after myocardial infarction<sup>60,61</sup>. In previous research other reservations were made about the effectiveness of antibiotic therapy; Gieffers et al.

postulated that intracellular Cp in circulating monocytes may be refractory to antibiotic therapy because of the inability of antibiotics to eliminate Cp from the circulation  $^{62}$ . In addition, nonviable Cp present in atherosclerotic lesions may still be potentially immunogenic, thus maintaining an inflammatory status  $^{63}$ . Still, antimicrobial drugs may contribute to improvement of endothelial dysfunction  $^{64,65}$  although it remains debatable whether any observed benefits of antibiotic drugs should be contributed to antimicrobial or anti-inflammatory effects. In a previous study we already demonstrated that azithromycin suppressed inflammatory responses to infection of hepatocytes although the observed benefits should be contributed to its antimicrobial effects rather than anti-inflammatory properties  $^{66}$ . In addition, the results from our present research support the role of antibiotic therapy for infection-induced atherosclerosis by indicating that azithromycin actually does improve endothelial function.

In conclusion we demonstrated that infection with *Cp* in endothelial cells *in vitro* attenuates eNOS, cGMP and ROS production in HUVECs and that azithromycin reversed *Cp*-induced effects on eNOS, cGMP and ROS-production.

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6

Iron enhances endothelial cell activation in response to Cytomegalovirus or *Chlamydia pneumoniae*-infection

# **Summary**

# **Background**

Chronic inflammation has been implemented in the pathogenesis of inflammatory diseases like atherosclerosis. Several pathogens like *Chlamydia pneumoniae* (*Cp*) and cytomegalovirus (CMV) result in inflammation and thereby are potentially atherogenic. Those infections could trigger endothelial activation, the starting point of the atherogenic inflammatory cascade. Considering the role of iron in a wide range of infection processes, the presence of iron may complicate infection-mediated endothelial activation.

# **Materials and methods**

Endothelial intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (VCAM-1) and endothelial selectin (E-selectin) expression were measured using flow cytometry, as an indication of endothelial activation. Cytotoxicity was monitored using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. Immunostaining was applied to measure *Cp* and CMV infectivity to endothelial cells.

#### Results

An increased number of infected endothelial cells in a monolayer population leads to a raised expression of adhesion molecules of the whole cell population, suggesting paracrine interactions. Iron additively up-regulated *Cp*-induced VCAM-1 expression, whereas synergistically potentiated *Cp*-induced ICAM-1 expression. Together with CMV, iron also enhanced ICAM-1 and VCAM-1 expression. These iron effects were observed without modulation of the initial infectivity of both microorganisms. Moreover, the effects of iron could be reversed by intracellular iron chelation or radical scavenging, conforming modulating effects of iron on endothelial activation after infections.

#### Conclusions

Endothelial response towards chronic infections depends on intracellular iron levels. Iron status in populations positive for *Cp* or CMV infections should be considered as a potential determinant for the development of atherosclerosis.

**Keywords:** Adhesion molecules, atherosclerosis, infection.

#### Introduction

Chronic inflammation plays a crucial role in coronary artery disease (CAD) and other manifestations of atherosclerosis<sup>1</sup>. The pathogenic inflammatory event is characterized by over-recruitment of leukocytes to the sites of inflammation. This event is mediated by activation of vascular endothelial cells. Endothelial cell activation, the key mechanism of atherosclerotic inflammation, is characterized by up-regulation of adhesion molecule expression, including vascular cell adhesion molecule-1 (VCAM-1), intercellular adhesion molecule-1 (ICAM-1) and endothelial selectin (E-selectin)<sup>2</sup>. These adhesion molecules have been found in human atherosclerotic lesions<sup>3-6</sup>. Chlamydia pneumoniae (Cp), the gram-negative obligate intracellular bacterium, is capable of infecting endothelial cells<sup>7-10</sup> as well as inducing the expression of adhesion molecules on these cells<sup>11,12</sup>. Cp-infection also leads to increased soluble adhesion molecules in human<sup>13</sup>. Differential induction of adhesion molecule expression by cytomegalovirus (CMV) has also been demonstrated 14-18. The up-regulation of adhesion molecules by Cp and CMV infections suggests a mechanism whereby infections could induce arterial disease. In addition, CMV and at a lesser extent Cp show high infection prevalence in the community 19,20. Cp has been established as a respiratory pathogen and contributed 10–20% of community-acquired pneumonia<sup>19</sup>, whereas the herpes virus CMV is associated with persistent, latent and recurrent infections due to reactivation of latent virus, with a prevalence of 50–90% in adults<sup>20</sup>.

Coronary artery disease (CAD) risk has been linked to certain persistent microorganism infections, like *Chlamydia pneumoniae* (*Cp*) and cytomegalovirus (CMV). Lack of correlations between *Cp*-serology and atherosclerotic lesion, however, has been observed in several studies<sup>21</sup>. Moreover, the evidence for the role of CMV in atherogenesis is conflicting<sup>22</sup>. Further studies are therefore warranted especially to unravel the pathological mechanisms of infections in arterial disease. Recently, we and other researchers have shown that iron status influences the endothelial activation state<sup>23–35</sup>.

There are many abnormal conditions that may cause increased body iron stores, and the formation of low molecular weight labile forms of iron that are capable of freely entering cells with no feedback-regulated process<sup>36</sup>. These conditions include hereditary haemochromatosis and secondary iron overload like in thalassaemia with frequent blood transfusions<sup>37</sup>. The labile forms of iron may play a role in the development of atherosclerotic vascular disease<sup>38,39</sup>. Coincidently, iron has been found accumulating

in human atheroma<sup>40</sup>. Several other studies show reduced formation of early atherosclerotic lesions by means of iron chelation or iron-deficient diets in experimental animals<sup>41-44</sup>. In this study, we investigated the effects of iron-rich and iron-withholding conditions during *Cp*- or CMV-infection on endothelial cells, in particular on the expression levels of VCAM-1, ICAM-1 and E-selectin. The possible involvement of iron-catalyzed oxygen-derived radical formation was also investigated. In light of the high frequencies of *Cp*- and CMV- infections in the population, together with the tendency of having increased body iron stores in conditions like hereditary haemochromatosis and secondary iron overload, this study provides important new insights and advances to the knowledge of the pathological mechanism of infections in atherosclerotic artery disease

#### **Methods**

#### **Baseline iron level**

The iron content of the endothelial growth medium-2 (EGM-2, Clonetics, Wallkersville, MD, USA) was measured by Vitros® 950 Chemistry System (Ortho-Clinical Diagnostics, Tilburg, the Netherlands) to monitor the baseline iron level in all of the experiments involving human umbilical cord endothelial cells (HUVECs). To avoid any external iron contamination, plastic materials having an affinity for iron lower than glass were used in all experiments.

#### **HUVEC** isolation and culture

HUVECs were isolated and cultured as described by Jaffe *et al.*<sup>45</sup>. To minimize donor-to-donor variability, HUVECs were pooled from three to four donors for each experiment. Experiments were performed at least three times on cells from sets of different donors. Moreover, the cells were always used and maintained at a cobblestone confluent density for all conducted experiments.

#### **Propagation of microorganisms**

Human embryonic lung (HEL) and buffalo green monkey (BGM) cells were cultured at 37 °C and 5% CO<sub>2</sub> in minimal essential medium Eagle with Earle's salts (EMEM, Gibco, Breda, the Netherlands) containing 10% Fetal Bovine Serum (FBS, Gibco). This culture medium was supplemented with 2 mmol/L L-glutamine (Gibco), 5 mL non-essential amino acids (Gibco), 10 mg/L vancomycin (Faulding Pharmaceuticals, Brussels,

Belgium), 4 mg/L amphotericin B (Fungizone; Bristol-Meyers Squibb, Woerden, the Netherlands) and 10 mg/L gentamycin (Schering Plough, Maarssen, the Netherlands). The same supplements were also added to the media used for the propagation of the virus strains. A clinical isolate of CMV was propagated in HEL cells with EMEM containing 2% FBS, 20 mmol/L Hepes and supplements. At > 80% cytopathologic effect, CMV-infected HEL cells were detached with trypsin/EDTA solution (Gibco) and centrifuged. The cell pellet was re-suspended in the same medium containing 2% FBS, 10% dimethyl sulphoxide (DMSO, Sigma-Aldrich, Zwijndrecht, the Netherlands) and supplements.

*Cp*-strain AR39 was propagated in BGM cells at 37 °C, 5% CO $_2$  in EMEM containing 10% FBS and 0.1% cycloheximide (Sigma-Aldrich) and supplements. After 72 h of growth, infected cells were frozen and thawed to release the elementary bodies. After a short centrifugation step, cell debris was discarded and 0.2 mol/L sucrose phosphate-glutamic acid (SPG) medium (2.088 g/L K2HPO4, 1.088 g/L KH2PO4, 68.46 g/L saccharose, 7.16 g/L L-glutamine, 10% FBS, 2.5 mg/L amphotericin B, 23 mg/L vancomycin and 18 mg/L gentamycin) was added (1:1 v/v). Both *Cp* and CMV stock suspensions were aliquoted and stored at -80 °C until further use. The tissue culture infective dose (TCID $_{50}$ ) of CMV was determined by daily examination of the infected HEL cells for cytopathologic effects during 1 week, whereas the TCID $_{50}$  of the *Cp* stock was calculated based on the number of immunofluorescent units per field in the infected BGM cells after staining with chlamydia culture confirmation monoclonal antibodies (de Beer, Diessen, the Netherlands)<sup>10</sup>.

#### Inoculation and immunostaining of endothelial cells

*Cp* and CMV were pre-diluted in endothelial EGM-2 medium and added at a multiplicity of infection (MOI) of 0.1 for both *Cp* and CMV. Uninfected cells and filtrate of microorganisms through a 100 kDa Microcon filter (Millipore, Bedford, MA, USA) were used as negative controls. 2 days after *Cp*-inoculation or 4 days after CMV inoculation, cells were harvested for flow cytometry. The infectivity of *Cp* and CMV to HUVECs was verified by immunostaining the infected cells with antibodies to *Cp* (30701 pathfinder chlamydia culture confirmation system, Bio-Rad, Redmond, WA, USA) and CMV (anti-CMV immediate early antigen clone E13, no. 12-003, Argene, Varilhes, France).

# Confocal scanning laser microscopy

For visualization purpose, infected immunostained cells were fixed in phosphate buffered saline (PBS) containing 3% paraformaldehyde (Polysciences, Warrington, PA, USA) and 0.02% glutaraldehyde (Merck, Darmstadt, Germany) and visualized using a Leica TCS-SP2 confocal scanning laser microscope and Leica confocal software (Leica Microsystems, GmbH, Heidelberg, Germany).

#### Preparation of iron, iron chelators and radical scavengers

A 10 mmol/L Fe(III) citrate (1:6 iron-citrate molar ratio; Sigma-Aldrich) solution was made by dissolving the iron crystals in distilled water at 56 °C for 30 min. Iron solutions were always freshly prepared and filter-sterilized prior to use. The iron chelators were prepared as stocks in PBS; 10 mmol/L deferoxamine (Novartis, Arnhem, the Netherlands) and 30 mmol/L deferiprone (Duchefa Biochemie, Haarlem, the Netherlands), and stored at –20 °C prior to use. The final pH in incubation medium was maintained at 7.8. Several radical scavengers were used in some experiments, including tempol (Sigma-Aldrich), thiourea (OPG, Utrecht, the Netherlands) and 1,3-dimethyl-2-thiourea (DMTU, Aldrich Chemical, Milwaukee, WI, USA).

# Viability assay

Cellular viability of HUVECs was monitored by the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazoliumbromide (MTT, Sigma-Aldrich) method<sup>46</sup>. Compound cytotoxicity was expressed as a  $TC_{50}$  denoting the concentration resulting in 50% loss of cell viability, as calculated by Calcusyn<sup>47</sup>.

# Fluorescence-activated cell sorting (FACS)

Human umbilical cord endothelial cells (HUVECs) were harvested by incubating with 0.2% trypsin-EDTA at 37 °C for 3 min. The cells were then incubated with fluorescence labelled monoclonal antibodies against the surface proteins, fluorescein isothiocyanate (FITC)-conjugated ICAM-1 antibody (R&D systems, Minneapolis, MN, USA), phycoerythrin (PE)-conjugated VCAM-1 antibody (BDBiosciences, San Diego, CA, USA), or Cychrome-conjugated E-selectin antibody (BDBiosciences), for 30 min at 4 °C. Each flow cytometric measurement was performed using a Becton Dickinson (San Jose, CA, USA) FACScan and 10.000 events were analyzed.

#### Measurement of oxidative stress

2,7-Dichlorofluorescein diacetate (DCFH-DA, Molecular Probes) is a non-polar compound that is converted into a membrane-impermeable non-fluorescent polar derivative, 2,7-dichlorofluorescein (DCFH) by cellular esterase after incorporation into cells. The trapped DCFH is rapidly oxidized to fluorescent 2,7-dichlorofluorescein (DCF) by intracellular hydrogen peroxide and hydroxyl radicals  $^{48}$ . Human umbilical cord endothelial cells (HUVECs) were harvested by incubating with 0.2% trypsin-EDTA at 37 °C for 3 min. Cells were then re-suspended in DCFHDA at a final concentration of 5  $\mu$ M, incubated for 30 min at room temperature and washed. The emission of the trapped, oxidized DCF in 10.000 cells was analysed on a FACScan.

# **Calcein assay**

In this assay<sup>49</sup>, cells were incubated with 0.125  $\mu$ M calcein-AM (30 min at 37 °C). The cells were washed twice to remove the remaining extracellular calcein-AM before fluorescence signal of calcein (excitation = 485 nm; emission = 530 nm) was followed using the Flexstation (Molecular Devices, Workingham, UK) at 37 °C. After a stable basal fluorescence signal was observed, iron was added to the incubation medium. Addition of iron quenches the fluorescence intensity of calcein signal. The accumulation of labile iron within cells due to addition of iron was expressed as the level of quenched calcein fluorescence adjusted to untreated controls at indicated time periods.

# **Data analysis**

Results are expressed as means  $\pm$  standard error of the mean (SEM). Differences in quantitative measures were tested for significance by using the unpaired two-tailed Student's *t*-test, unless otherwise stated. Significance was established when p < 0.05.

#### Results

#### Cp and CMV infections of endothelial cells

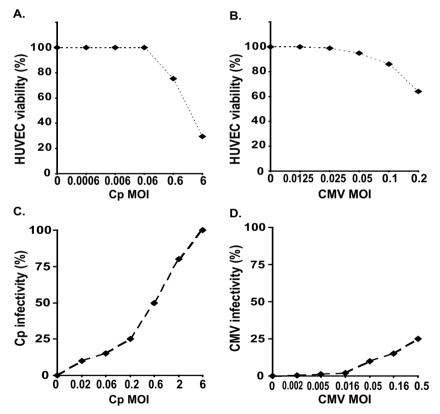
The logTCID $_{50}$  values of the stock Cp and CMV were calculated as 5.5 and 4, respectively, indicating titers of 6x10 $^6$  Cp/mL and 2x10 $^5$  CMV/mL. As HUVEC density was 10 $^5$  cells/cm $^2$  at a cobblestone confluence, in order to obtain an MOI of 0.1 for Cp and CMV, Cp was diluted 300-fold, whereas CMV was diluted 10-fold before being used for HUVEC inoculation.

# **Cytotoxicity testing**

The viability of HUVECs, after *Cp* (**Figure 1a**) or CMV infection (**Figure 1b**) at an MOI of 0.1, was > 85%.

# Cp and CMV infectivity

Infectivity of Cp and CMV to HUVECs were counted using fluorescence microscopy on a random and blind basis after immunostaining procedure, using monoclonal antibodies specific towards Cp or CMV. The complete developmental cycle of Cp in cell culture models is between 48 and 72 h<sup>50</sup>, whereas the slow-replicating CMV enters the early stage of infection at 72–96 h post-infection, and reaching the late stage of



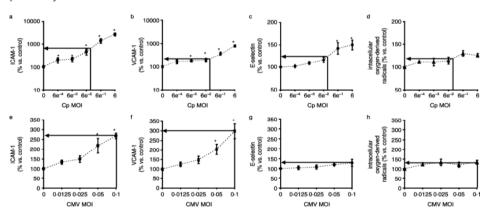
**Figure 1** Endothelial viability and microorganism infectivity. (a) Shows the percentage of human umbilical vein endothelial cell (HUVEC) viability determined by the 3-(4,5-dimethylthiazol- 2-yl)-2,5-diphenyltetrazoliumbromide (MTT) cytotoxicity assay, 2 days after *Chlamydia pneumoniae* (Cp) infection at an indicated multiplicity of infection (MOI), whereas (b) shows HUVEC viability 4 days after cytomegalovirus (CMV) infection. (c) shows the percentage of Cp infectivity towards HUVECs determined by immunostaining, and (d) shows CMV infectivity. The interpolated values at MOI of 0.1 are indicated by arrows (all data, n = 3).

infection at the 7th day post-infection<sup>51</sup>. To mimic the conditions of chronic steady-grade infections without having secondary infections, HUVECs were inoculated for 2 days with Cp or 4 days with CMV. Cp at an MOI of 0.1 resulted in < 20% infection (**Figure 1c**). Additionally, CMV at an MOI of 0.1, gave rise to < 10% infection (**Figure 1d**).

# Low-grade *Cp*- and CMV-infections induced endothelial adhesion molecule expression without an increased oxygen-derived radical production

Chlamydiapneumoniae (Cp) infection markedly induced ICAM-1 (**Figure 2a**) and VCAM-1 expression (**Figure 2b**). A two-fold increase in ICAM-1 expression was observed with Cp-infection at an MOI of as low as 0.0006. E-selectin was significantly up-regulated by Cp starting at an MOI of 0.6 (**Figure 2c**).

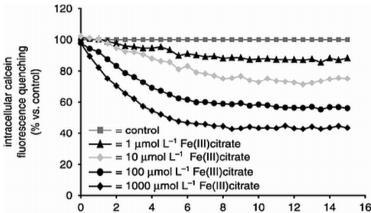
Cytomegalovirus (CMV) at an MOI of as low as 0.05 induced a more than two-fold increase in both ICAM-1 and VCAM-1 expression, whereas E-selectin up-regulation was negligibly noticed (**Figure 2e-g**). The filtrate of microorganisms through a 100 kD Microcon filter did not result in up regulation of adhesion molecule expression (not shown), confirming the specific effects of *Cp* and CMV infections on the induction of endothelial adhesion molecule expression. No increase in intracellular oxygen-derived radicals was observed in both *Cp*- (**Figure 2d**) and CMV-infected HUVECs (**Figure 2h**), indicating that the induction of ICAM-1 and VCAM-1 in infected HUVECs could not primarily be attributed to radical formation.



**Figure 2** Induction of adhesion molecule expression by Cp or CMV infection. The expression of (a) intercellular adhesion molecule-1 (ICAM-1); (b) vascular cell adhesion molecule-1 (VCAM-1); (c) endothelial selectin (E-selectin); and (d) the levels of intracellular oxygen derived radicals in human umbilical vein endothelial cells (HUVECs) after *Chlamydia pneumoniae* (Cp) infection at an indicated multiplicity of infection (MOI) (mean  $\pm$  SEM, n = 4, \*P < 0.05). The expression of (e) ICAM-1; (f) VCAM-1; (g) E-selectin; and (h) the levels of intracellular oxygen-derived radicals in HUVECs after cytomegalovirus (CMV) infection at an indicated MOI. The interpolated values at MOI of 0.1 are indicated by arrows (mean  $\pm$  SEM, n = 4, \*P < 0.05).

# Increased level of endothelial intracellular iron due to the addition of low molecular weight iron

With no external iron addition, the baseline iron level in the basal cell culture growth medium, EGM-2, was 0.36  $\mu$ mol/L. Accumulation of intracellular labile iron within HUVECs was monitored by following the fluorescence of calcein for up to 15 h (**Figure 3**). An iron concentration of as low as 1  $\mu$ mol/L was able to quench 5% of calcein signal after 2 h and 10% after 5 h, indicating an increase in the intracellular labile iron level. This result indicates that the addition of low molecular weight iron augmented the level of endothelial cytoplasmatic labile iron.



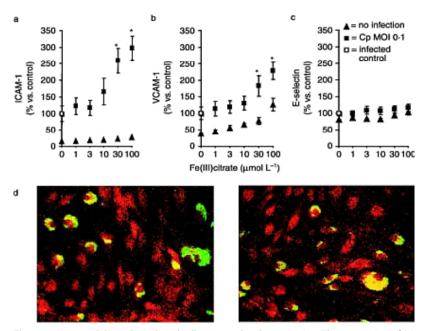
**Figure 3** Endothelial intracellular iron level is modulated by addition of low molecular weight iron. Data represent the normalized mean fluorescence of intracellular calcein in human umbilical vein endothelial cells (HUVECs) in the presence of specified concentrations of Fe(III) citrate, monitored up to 15 h (n = 3). Quenching indicates the presence of intracellular labile iron.

# *Cp*-induced endothelial adhesion molecule expression was markedly potentiated by iron

At an MOI of 0.1, *Cp*-induced VCAM-1 expression was additively up-regulated by iron, whereas ICAM-1 expression was synergistically up-regulated by iron (**Figure 4a-c**). Iron of 30 µmol/L significantly enhanced *Cp*-induced VCAM-1 expression two-fold and ICAM-1 expression almost threefold.

Infectivity and the size of inclusions of Cp in HUVECs were not affected by various concentrations of iron, ranging between 0 and 1000  $\mu$ mol/L. This result was obtained by examination using fluorescence microscopy after immunostaining with monoclonal

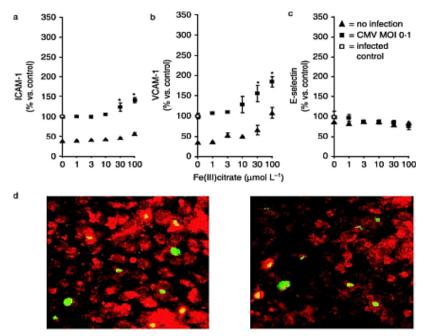
antibody against *Cp* (**Figure 4d**) on a blind and random basis. This finding indicates that iron could modulate endothelial response towards chronic *Cp*-infection without affecting initial infectivity and the growth of *Cp*.



**Figure 4** Iron modulates Cp-induced adhesion molecule expression. The expression of (a) intercellular adhesion molecule-1 (ICAM-1); (b) vascular cell adhesion molecule-1 (VCAM-1); and (c) endothelial selectin (E-selectin) on human umbilical vein endothelial cells (HUVECs) 2 days after *Chlamydia pneumoniae* (Cp) infection in the presence of increasing iron concentrations (mean  $\pm$  SEM, n=4, \*P<0.05). (d) Confocal laser micrographs, representing four different slides, visualize the infectivity of Cp (green) on HUVECs (red) in the absence and presence of iron. (*For color figure see page 220*)

# CMV-induced endothelial adhesion molecule expression was enhanced by iron

Both CMV-induced VCAM-1 and ICAM-1 expression were additively up-regulated by iron (**Figure 5a-b**), whereas E-selectin was not affected (**Figure 5c**). Iron of 30  $\mu$ mol/L significantly enhanced CMV-induced VCAM-1expression 1.5-fold, and CMV-induced ICAM-1 expression 1.3-fold. As noted for *Cp*, the infectivity of CMV towards HUVECs and the size of CMV inclusions were not affected by various concentration of iron, ranging between 0 and 1000  $\mu$ mol/L (**Figure 5d**). This result indicates that iron could also modulate endothelial response towards chronic CMV-infection, without affecting initial infectivity and the growth of CMV.

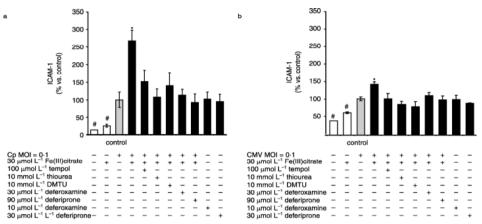


**Figure 5** Iron modulates CMV-induced adhesion molecule expression. The expression of (a) intercellular adhesion molecule-1 (ICAM-1); (b) vascular cell adhesion molecule-1 (VCAM-1); and (c) endothelial selectin (E-selectin) on human umbilical vein endothelial cells (HUVECs) 4 days after cytomegalovirus (CMV) infection in the presence of increasing iron concentrations. (mean  $\pm$  SEM, n=4, \*p < 0.05). (d) Confocal laser micrographs, representing four different slides, visualize the infectivity of CMV (green) on HUVECs (red) in the absence and presence of iron. (*For color figure see page 220*)

# Iron chelation and radical scavenging could counteract potentiating effects of iron on infections

Addition of chelator-bound iron no longer modulated *Cp*- or CMV-induced endothelial adhesion molecule expression (**Figure 6a-b** for ICAM-1 expression), confirming the specific modulating effects of iron on endothelial response towards infections. The scavengers, including tempol, thiourea and DMTU, were also able to counteract the modulating effects of iron on *Cp*- or CMV-infection, indicating the involvement of oxygen-derived radicals in this process (**Figure 6a-b**). Furthermore, addition of iron chelator alone did not significantly down-regulate *Cp*- or CMV-induced adhesion molecule expression (**Figure 6a-b**), suggesting that infections may induce endothelial activation through a distinct pathway than iron. Concentrations of 10 µmol/L for

deferoxamine and 30  $\mu$ mol/L for the bidentate deferiprone were chosen to cover the possible highest level of iron in HUVECs. These concentrations are below the TC<sub>50</sub> values (deferoxamine = 15 ± 2.3  $\mu$ mol/L, deferiprone = 100 ± 11.3  $\mu$ mol/L) <sup>24</sup>.



**Figure 6** Effects of iron chelation and radical scavenging on infections. The expression of intercellular adhesion molecule-1 (ICAM-1) on (a) *Chlamydia pneumoniae (Cp)* or (b) cytomegalovirus (CMV) infected human umbilical vein endothelial cells (HUVECs) in the presence of indicated compounds. (mean  $\pm$  SEM, n=3, \*higher than control p<0.05; #lower than control p<0.05).

#### Discussion

Endothelial dysfunction plays an important role, not only at the initial step in the development of atherosclerosis, but also at a critical late step of thrombosis that leads to vessel occlusion and acute cardiovascular events<sup>1</sup>. As serological associations were found between *Cp* or CMV infections and acute myocardial infarctions as well as chronic coronary heart disease, there has been much effort on determining how endothelial infection by *Cp* or CMV causes endothelial dysfunction. One well-characterized phenotype of endothelial dysfunction is increased expression of the endothelial adhesion molecules, E-selectin, ICAM-1 and VCAM-1<sup>2</sup>. In this study, we analysed endothelial activation after low-grade *Cp*- or CMV-infections that resulted in less than 20% infection of an endothelial cell population. Both low-grade chronic *Cp*- and CMV-infections had readily up-regulated the expression of adhesion molecules. The whole endothelial cell population showed relatively homogenous induction of adhesion molecule expression despite a low number of infections. This finding indicates that paracrine interactions such as through secretion of pro-inflammatory

cytokine interleukin-6 by the infected cells are crucial to generate response towards infections<sup>52</sup>. Our results therefore support the hypothesis that *Cp*- or CMV- infections are likely to contribute to the chronic inflammatory events in the vasculature associated with atherosclerosis.

In this study, we show that the expression of adhesion molecules in HUVECs infected with *Cp* was further enhanced when iron-rich medium was used during incubation. This iron-rich medium modulated intracellular iron level. It has been previously described that the infectivity and the growth of a relatively high-grade *Cp*-infection with a long incubation time could be restricted by iron chelation in epithelial cell line<sup>53,54</sup>. Using the current experimental settings mimicking a chronic low-grade vascular Cp infection, we observed modulating effects of iron on endothelial response towards Cp, before the infectivity or the growth of the microorganism were noticed. We also observed counteracting effects of iron chelation as well as radical scavenging on the effects of iron. These findings suggest that the modulating role of iron in endothelial response towards chronic infection is not by way of increasing the infectivity or the growth of the microorganism. Instead, our findings suggest that iron primarily exerted its effects through priming of the endothelial cells by generating oxidative stress. These ironprimed endothelial cells may consequently be more responsive towards the paracrine effects of infection. Additionally, it is known that for Cp, attachment is sufficient to initiate an endothelial response, while uptake may not be required<sup>11</sup>. Furthermore, we observed additive effects of iron on Cp-induced VCAM-1 expression and synergistic effects on Cp-induced ICAM-1 expression. This could be due to the differential signal transduction activated by Cp on endothelial cells11,12 that in turn was potentiated by iron <sup>30</sup>. Iron chelators alone, however, did not down-regulate *Cp* or CMV-induced adhesion molecule expression, because there were no changes in the initial Cp or CMV infectivity to HUVECs due to addition of iron chelators. This could be because the resting iron levels in HUVECs passages 2-3 used in this study were already low, that is, in the order of 10 folds less than in freshly isolated cells <sup>26,28</sup> suggesting that further chelation probably would give little or no effects. Additionally, the level of intracellular oxygen-derived radicals was not changed due to infections. This may explain the absence of iron chelator effects as antioxidants in reducing infection enhanced endothelial adhesion molecule expression. This finding also suggests that infections may exert their effects on endothelial activation through a pathway that is different from the formation of oxygen-derived radicals.

Cytomegalovirus (CMV) infection has a relatively slow development cycle<sup>51</sup>. The early

stage of infection is started at 72-96 h post-infection<sup>51</sup>. During this time CMV has produced the immediate early gene products that are capable of inducing endothelial activation<sup>15,17</sup>. In this study we harvested HUVECs at the 4th day post infection, to allow CMV-induced endothelial activation to take place without the propagation of the microorganism. Up-regulation of both ICAM-1 and VCAM-1 was observed, whereas E-selectin was not affected significantly. This result confirms previous studies<sup>14-16</sup> showing differential induction of endothelial adhesion molecule expression by CMV. Furthermore, compared with controls, the iron primed endothelial cells generated higher levels of CMV-induced ICAM-1 and VCAM-1 expression without affecting the initial infectivity of CMV. These results again demonstrate the potentiating effects of iron on endothelial response towards infections without modulation of the initial infectivity of the virus.

The role of infections in promoting or accelerating atherosclerosis has been extensively demonstrated, although some studies showed lack of evidence<sup>21,22</sup>. Our *in vitro* study demonstrated modulating effects of iron on the endothelial response towards chronic infections of *Cp* and CMV. These findings imply that iron status in populations positive for *Cp*- or CMV- infections could be a potential determinant for the development of atherosclerosis. Further studies showing functional consequences of these recent findings such as the extent of leukocyte infiltration through endothelial cells are indeed warranted, in order to clarify the true role of iron in atherosclerosis. Furthermore, this study also implies that populations with increased body iron levels such as hereditary haemochromatosis and thalassaemia with frequent blood transfusions <sup>37</sup>, and populations with disturbed iron homeostasis such as chronic renal insufficiency with iron substitution <sup>55</sup> may have an aggravated susceptibility towards atherosclerotic vascular disease when they are positive for *Cp*- or CMV-infections.

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Infection-induced inflammatory response of adipocytes in vitro

### **Abstract**

# **Background**

Abdominal obesity plays an important role in the development of insulin resistance, diabetes mellitus and atherosclerosis. The exact pathophysiological mechanisms are unclear but adipocyte dysfunction is thought to be crucial. Infections are associated with the development of atherosclerosis as well as diabetes. In this study we investigated whether adipocytes can be infected and whether this results in production of inflammatory cytokines relevant for the development of atherosclerosis and diabetes.

#### **Methods**

Pre-adipocytes were cultured and differentiated into mature adipocytes *in vitro*. Adipocytes and pre-adipocytes were incubated with infective and heat inactivated *Chlamydia pneumoniae (Cp)*, Cytomegalovirus (CMV), Adenovirus (Ad) subtypes 2 & 36, Influenza A (Inf A) and Respiratory Syncitial Virus (RSV). After 48 hours, adiponectin, Interleukin-6 (IL-6), Tumor necrosis factor-alpha (TNF-α) and Plasminogen Activator Inhibitor-1 (PAI-1) were measured in supernatants.

#### Results

Infection of adipocytes with Ad 36, CMV and RSV resulted in increased IL-6 production from  $192\pm22$  pg/mL (uninfected) to  $1030\pm86$  pg/mL,  $838\pm59$  pg/mL and  $1241\pm191$  pg/mL respectively (all p<0.01 vs. control). In addition, Ad 36 infection slightly reduced PAI-production in adipocytes ( $285\pm26.8$  ng/mL vs. uninfected:  $477\pm71.2$  ng/mL; p=0.05) and pre-adipocytes ( $709\pm43.3$  ng/mL vs. uninfected:  $1071\pm71.8$  ng/mL; p<0.01). In contrast, human adenovirus type 2 did not exert any effect on IL-6 or PAI-production. None of the micro organisms induced significant changes in adiponectin and/or TNF- $\alpha$  production.

#### Conclusions

Adipocytes can be infected with several micro organisms *in vitro*. Infection of adipocytes with Ad 36, but not Ad 2 leads to increased production of IL-6 which might contribute to chronic low-grade inflammation, a process known to be involved in the development of cardiovascular diseases and type 2 diabetes.

### Introduction

Obese patients have an increased risk for the development of atherosclerosis and diabetes mellitus type 2 and myocardial infarction<sup>1</sup>. Although the exact pathophysiological mechanisms are not yet clear, insulin resistance, closely related with obesity, and inflammation are key components <sup>2,3</sup>. Abdominal adipose tissue not only absorbs and stores free fatty acids, but is also able to produce and secrete adipokines such as IL-6, Tumor Necrosis Factor (TNF- $\alpha$ ), Plasminogen Activator Inhibitor-1 (PAI-1) and adiponectin. By producing hormones and inflammatory cytokines, adipose tissue is likely to play an important role in the development of diabetes and atherosclerosis <sup>4-7</sup>.

Adipocyte dysfunction can be the cause but also the consequence of insulin resistance. Causes for adipocyte dysfunction are not well established but include abdominal obesity and elevated free fatty acid plasma concentration 8.

Cytomegalovirus (CMV) en *Chlamydia pneumoniae (Cp)* are thought to be involved in the pathogenesis of atherosclerosis and cardiovascular disorders <sup>9-14</sup>. It has been shown before that adenoviruses are able to infect adipocytes <sup>15,16</sup> and that this is possibly related to the development of obesity <sup>17,18</sup>. Data from studies in animals indicate a possible role for virus infections in the etiology of human obesity <sup>19-21</sup>. Moreover, Adenovirus-36 is associated with increased body weight and paradoxically lower cholesterol and triglyceride levels in Ad-36 antibody-positive vs. -negative subjects <sup>22</sup>. Therefore it can be hypothesized that (chronic) virus infections induce adipocyte dysfunction thus leading to the development of diabetes mellitus and/or cardiovascular diseases.

In the present study we investigated whether several viruses are able to infect adipocytes *in vitro* and whether this leads to adipocyte dysfunction as measured by the altered production of IL-6, TNF- $\alpha$ , adiponectin and PAI-1.

#### Methods

# **Culturing of adipocytes**

Cryopreserved human subcutaneous pre-adipocytes from an obese donor (BMI >30; SP-F3 passage 2, ZenBlo Inc, Research Triangle Park, NC, USA) were cultured in pre-adipocyte medium (PM-1, ZenBio Inc) in 75 cm² flasks (Nunc, Roskilde Denmark) at 37° C with 5% CO<sub>2</sub>. Subsequently, cells were seeded in 24-well plates (Nunc, Roskilde Denmark) and grown to confluence (passage 4). When cells were grown confluent, differentiation was started by replacing pre-adipocyte medium by differentiation

medium (DM-2, ZenBio Inc) which is PM-1 supplemented with 3-Isobutyl-1-methylxanthine (IBMX), dexamethasone, insulin and a PPAR-γ agonist. After one week the differentiation medium was replaced by adipocyte medium (AM-1, ZenBio Inc). Cells were refed with fresh AM-1 every 3-4 days until experiments were performed. Control pre-adipocytes were kept and refreshed with PM all the time. The progression of lipid droplet accumulation and cell morphology during differentiation was monitored using an inverted microscope or Oil Red O staining.

# Infectious and inactivated micro-organisms

Adenovirus subtypes 2 (Ad2), 36 (Ad36) and Respiratory Syncitial Virus (RSV) were propagated on human Hep-2 larynx carcinoma cells (#03-108/ATCC #CCL 23; Flow laboratories/Amstelstad BV, Zwanenburg, The Netherlands). The Ad36 isolate was kindly provided by E. Fries, Erasmus University Rotterdam. Influenza strain A/H1N1/Nederland/300/00 (Inf) was propagated on LLC-MK2 Rhesus monkey kidney cells (#03-200/ATCC #CCL7; Flow laboratories/Amstelstad BV, Zwanenburg, The Netherlands) and Cytomegalovirus (CMV) was cultured on a human embryonic lung (HEL) cell line. The non-adipogenic Ad2 subtype (control virus for Ad36), RSV, Inf and CMV strains were clinical isolates selected from patients with common respiratory infections (Dept. of Virology, Diakonessen Hospital Utrecht, The Netherlands). The intracellular bacterium Chlamydia pneumoniae AR 39 (Cp; ATCC #535920) was grown on Buffalo Green Monkey (BGM) kidney cells (ECACC no 90092601).

Ad2, Ad36, RSV and Inf infections in cell culture were identified by indirect immunofluorescent staining with a pooled specimen screening reagent containing affinity purified mouse monoclonal antibodies directed against adenovirus; influenza A and B; parainfluenza types 1, 2, and 3 and RSV (Bartels VRK Anti-Viral Screening Reagent, #B1029-86A; Trinity Biotech, Bray Ireland). Cytomegalovirus infection was identified by indirect immunofluorescent staining with anti CMV immediate early antigen antibodies (clone E13, #11-003; Argene Varilhes, France). Apple-green fluorescence in the cytoplasm (Ad, RSV, Inf) or nuclei (CMV, Inf) is observed in a positive stain while uninfected cells (counterstained with Evan's blue) appear red. Cp-infection was identified by direct immunofluorescent detection using the Pathfinder\* Chlamydia Culture Confirmation System (#30701 Biorad, Veenendaal, The Netherlands). This assay detects all known serovars of *Chlamydia trachomatis, Chlamydia psittaci* and *Chlamydia pneumoniae*. Cells infected with *Cp* will show a characteristic apple-green fluorescence of cytoplasmic inclusions against a red background. After propagation on

the appropriate cell lines in tissue culture bottles, the upscaled inoculation materials were harvested, aliquoted and subsequently frozen at -80 °C. Infective doses ( $TCID_{50}$ ) of each isolate were determined using corresponding cell lines according to the method of Reed and Muench <sup>23</sup>. As infection controls for the main experiments, aliquots of all isolates were inactivated using one standard procedure: incubation at 56° C for one hour followed by overnight UV-irradiation. Effectiveness of the inactivation procedure was checked by incubating undiluted inactivated isolates on the cell lines on which they were propagated initially for 48 hours.

# **Infection experiments**

Cultures with more than 75% differentiated adipocytes were infected 13-15 days after the start of the differentiation. For each inoculation of cultures with differentiated adipocytes, cultures with pre-adipocytes were inoculated simultaneously. Infection rates were determined by calculating the number of infectious virus per cell. This was expressed as the multiplification of infection (MOI). For each inoculation with infective virus, inactivated virus was inoculated as an infection control. As a negative control, cells were incubated with culture medium only. All incubations were monitored by the corresponding immunofluorescent staining for each virus type. Optimal infection rates for the main experiments were determined in preliminary infection experiments with serial dilutions of each stock (data not shown). The rationale for using fairly low MOIs was to mimic chronic infection *in vivo* and to avoid unwanted cell death. Therefore infection rates of maximum 25% (fluorescence) were chosen.

Prior to infection, adipocytes were refed with fresh AM and pre-adipocytes with PM. Adipocytes and pre-adipocytes were inoculated with 200  $\mu$ L of both infective and inactivated virus; multiplifications of infections (MOI) were: 0.01 and 0.001 for CMV, Ad2 and Ad 36 and MOI 0.02 and 0.002 for *Cp*, Influenza and RSV.

In addition to the infections, the ability of cells to produce adipokines was measured by incubating the cells with 1 or 10 ng/mL IL-1 $\beta$  or IL-6 or 100  $\mu$ g/mL TNF- $\alpha$ .

All experiments were performed 3 times in duplicate on cells at passage 4, grown in 24-well plates. After inoculation, plates were centrifuged 1 hour at 500g. All incubations were kept for 48 hours at 37 °C and 5%  $\rm CO_2$ . After 48 hours supernatants were collected and frozen at -80 °C until adipokines were measured. Infections and incubations with inactivated virus were checked by immunofluorescence.

### Measurement of adipokines

Commercially available ELISA kits were used for measuring IL-6 (Pelikine compact human IL-6 kit, # M1916 Sanguin Reagents, The Netherlands), (Pelikine compact human TNF-α kit, # M1923 Sanguin Reagents, The Netherlands), PAI (PAI-1 antigen Elisa reagent kit, #TC11070, Technoclone GmbH Vienna, Austria) and adiponectin (human Adiponectin/Acrp30 ELISA kit, (#ELH-ACRP30-001, RayBiotech Inc. Norcross GA, USA). All assays were performed three times in duplicate. Optical densities were measured with an automated plate reader (Elx800, Biotek Instruments Inc.) at 450/630 nm and data were processed in analytical curve fitting software (KC Junior, Biotek instruments Inc.).

#### Statistical analysis

ELISA results were processed using the GraphPad Instat<sup>m</sup> statistical software package (V2.05a). After infection, adipokine levels in the supernatants of infected adipocytes were compared with those in pre-adipocytes and in uninfected control cells or cells incubated with inactivated virus. In addition, adipokine production after incubation with IL-1 $\beta$ , IL-6 or TNF- $\alpha$  was compared to the adipokine levels in uninfected cells. An unpaired Student t-test was performed on all results unless standard error of means were significantly different (p <0.05). On those results where standard errors of means did differ, a Welch t-test, correcting for these differences was performed.

#### Results

#### (Pre-) adipocyte culture

After seeding in 24-well plates, cultured pre-adipocytes (passage 4) needed 3-5 days to grow to confluence (**Figure 1a**). During differentiation cells transformed from spindle shaped "fibroblast-like" to more rounded cells which accumulated lipid droplets in their cytoplasm (**Figure 1b**). The differentiation grade was at least 75% after 13 days.

# Virus infection of adipocytes

All viruses (CMV, Ad2, Ad 36, Influenza A, RSV) and Cp were able to infect adipocytes *in vitro* (**Figure 2 a,b,c,d,e,f**). The corresponding MOIs were 0.01 and 0.001 for CMV, Ad2 and Ad 36 and MOI 0.02 and 0.002 for *Cp*, Influenza A and RSV.

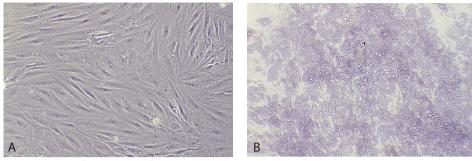
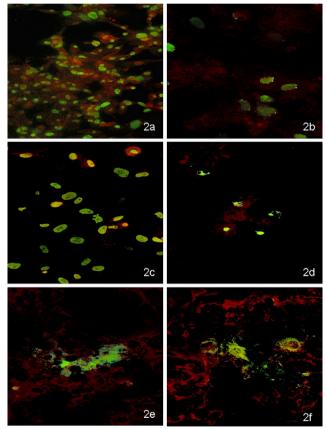


Figure 1 (Pre-) Adipocytes in culture (200x)

- A Pre-adipocytes: spindle-shaped "fibroblast-like" cells grown to confluence.
- B Adipocytes: differentiated cells 13 days after start of differentiation. Cells are rounded and have accumulated lipid droplets in their cytoplasm. (For color figure see page 221)



**Figure 2. Fluorescent images of infected adipocytes** (200x, 90% differentiated) after 48 hours infection with: Ad2, Ad 36, Cp, CMV, Inf, and RS-virus. Background (red): uninfected cells. Infected cells: apple green fluorescence either condensed in the nuclei (CMV;2c) or diffuse in the cytoplasm (Ad 2, Ad 36, Inf, RSV; 2a,b,d,e,f respectively). Cp-infected cells: compact fluorescent vacuoles in the cytoplasm (2d). (*For color figure see page 221*)

### IL-6 production by uninfected pre-adipocytes and adipocytes

During the differentiation process, adiponectin production increased from  $159\pm155$  pg/mL at day 0 (start of differentiation) to  $259\pm110$  pg/mL on day 3 and  $8972\pm3228$  pg/mL on day 7. Pre-adipocytes did not produce adiponectin at all. The IL-6 production by uninfected adipocytes decreased from  $3058\pm550$  pg/mL at day 0 (start of differentiation) to  $1433\pm485$  pg/mL (day 3),  $1146\pm285$  pg/mL (day 7) and  $71\pm32$  pg/mL (day 9) and increased from day 13: ( $201\pm66$  pg/mL) to day 15: ( $177\pm24$  pg/mL) and day 19 ( $257\pm57$  pg/mL). IL-6 production in pre-adipocytes remained at the same level as was measured at the start of the differentiation phase.

# Adipokine production by infected adipocytes

After 48 hours, IL-6 production of adipocytes was 1030±86 pg/mL (p<0.01) after incubation with infective Ad36 and 744±91 pg/mL (p <0.01) after incubation with inactivated Ad36 respectively, both compared to uninfected adipocytes (192±22 pg/mL). However, there was not a significant difference between IL-6 production by heat inactivated an infective Ad36 (p=0.08). After incubation with lower doses of A36, IL-6 levels in adipocytes were not elevated anymore (p<0.05). Therefore the effect was considered to be virus-dose dependent. In contrast to the clearly elevated IL-6 production by Ad 36, IL-6 production in Ad2 infected adipocytes was not affected (171±30 pg/mL; p>0.05).

Compared to uninfected adipocytes, infection of adipocytes with CMV or RSV also induced significant increase in IL-6 production (838±59 pg/mL and 1241±191 pg/mL respectively vs. uninfected adipocytes (192±22 pg/mL; p<0.01) (**Figure 3a**).

Baseline production of IL-6 by pre-adipocytes was considerably higher than in adipocytes (5234±837 pg/mL vs. 192±22 pg/mL) but quite similar to infected adipocytes, infection of pre-adipocytes with Ad36 or RSV resulted in a more than 2-fold increase in IL-6 from baseline (10957±1945 pg/mL (p<0.05) and 12079±2302 pg/mL (p<0.01) respectively vs. 5234±837 pg/mL in uninfected cells) CMV-infection did not lead to elevated IL-6 production by pre-adipocytes (**Figure 3b**).

Mean TNF- $\alpha$  productions of uninfected adipocytes and pre-adipocytes were 52.9  $\pm$  3.9 pg/mL and 55.4  $\pm$  11.3 pg/mL respectively as measured after 15 days incubation. None of the micro-organisms were able to increase TNF- $\alpha$  secretion by adipocytes (**Figure 4a**). In contrast, pre-adipocytes infected with the highest doses of Ad 36, CMV and RS-virus produced approximately 2-fold more TNF- $\alpha$  than uninfected pre-adipocytes

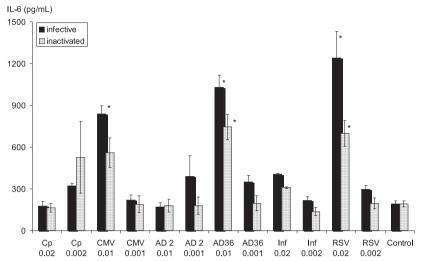


Figure 3a. IL-6 production in infected adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). II-6 production in adipocytes incubated with infective micro-organisms. IL-6 production in adipocytes incubated with inactivated micro-organisms. Y-axis: IL-6 production in pg/mL. X-axis: Micro-organisms used for incubations with corresponding MOI (multiplification of infection) values. Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus; \* p <0.01 (t-test).

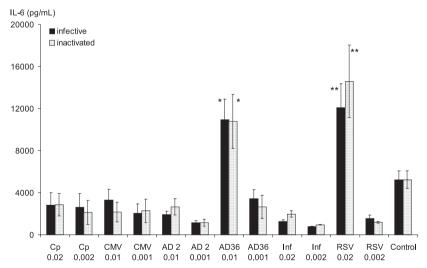


Figure 3b. IL-6 production in infected pre-adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). ■: IL-6 production in pre-adipocytes incubated with infective micro-organisms. 

IL-6 production in pre-adipocytes incubated with inactivated micro-organisms. 

Y-axis: IL-6 production in pg/mL. 

X-axis: Micro-organisms used for incubations with corresponding MOI (multplification of infection) values. 

Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus; \* p<0.01, \*\* p<0.05 (t-test).

(120.3 $\pm$ 38.2 pg/mL, 102.9 $\pm$ 17.2 pg/mL and 122.1 $\pm$ 54.5 pg/mL, respectively vs. 55.4  $\pm$  11.3 pg/mL in uninfected cells), but this increase was not statistically significant (p> 0.05; Welch t-test). There was also a tendency towards increased TNF- $\alpha$  production by CMV, Ad36 and RS-infected pre-adipocytes compared with pre-adipocytes incubated with the same, but inactivated, micro-organisms (102.9 $\pm$ 17.2 vs. 45.8 $\pm$ 18.1 pg/mL for CMV, 120.3 $\pm$ 38.2 vs. 37.1 $\pm$ 18.6 pg/mL for Ad36 and 122.1 $\pm$ 54.5 vs. 47.6 $\pm$ 1.6 pg/mL for RS), but again these differences were not statistically significant (p> 0.05; Welch t-test) (**Figure 4b**).

Only incubation with the highest infective dose of Ad36 led to decreased PAI-1 production in pre-adipocytes (709 $\pm$ 43.3 ng/mL), compared to uninfected pre-adipocytes (1071 $\pm$ 71.8 ng/mL; p<0.01) but not in adipocytes: 285 $\pm$ 26.8 ng/mL; p=0.05 vs. 477 $\pm$ 71.2 ng/mL in uninfected adipocytes. None of the other micro-organisms affected PAI-production (**Figure 5a,b**).

After 48 hours, the adiponectin production by uninfected adipocytes was 9325±3029 pg/mL. Compared to adiponectin production by uninfected adipocytes, the adiponectin production by adipocytes that were incubated with either infective or inactivated micro-organisms, tended to decrease. (Overall -35%; Cp -37%, CMV -52%, Ad2 -30%, Ad-36 -30%, Inf A -26%, RSV -36%). This effect however, was not statistically significant (**Figure 6**). Pre-adipocytes did not produce adiponectin at all (data not shown).

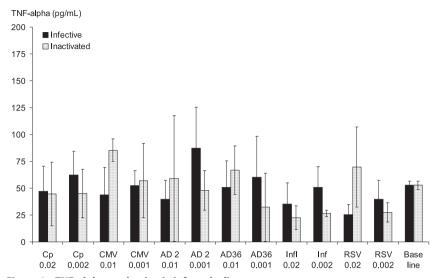


Figure 4a. TNF-alpha production in infected adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). TNF-alpha production in adipocytes incubated with infective micro-organisms. TNF-alpha production in adipocytes incubated with inactivated micro-organisms. Y-axis: TNF-alpha production in pg/mL. X-axis: Micro-organisms used for incubations with corresponding MOI (multiplification of infection) values. Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus.

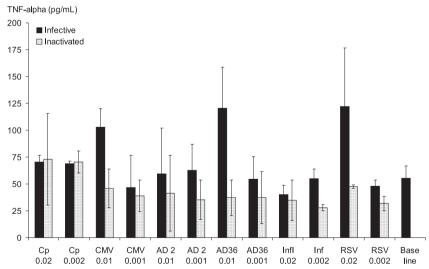


Figure 4b. TNF-alpha production in infected pre-adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). TNF-alpha production in pre-adipocytes incubated with infective micro-organisms. TNF-alpha production in pre-adipocytes incubated with inactivated micro-organisms. Y-axis: TNF-alpha production in pg/mL. X-axis: Micro-organisms used for incubations with corresponding MOI (multplification of infection) values. Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus.

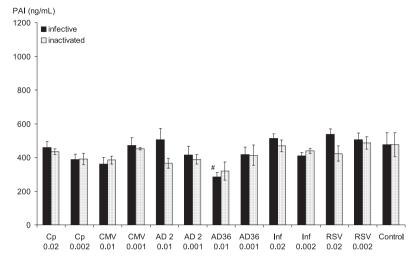


Figure 5a. PAI-1 production in infected adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). : PAI-1 production in adipocytes incubated with infective micro-organisms. PAI-1 production in adipocytes incubated with inactivated micro-organisms. Y-axis: PAI-1 production in ng/mL. X-axis: Micro-organisms used for incubations with corresponding MOI (multiplification of infection) values. Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus; # p=0.05 (t-test).

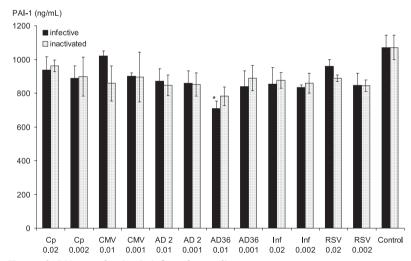


Figure 5b. PAI-1 production in infected pre-adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). PAI-1 production in pre-adipocytes incubated with infective micro-organisms. PAI-1 production in pre-adipocytes incubated with inactivated micro-organisms. Y-axis: PAI-1 production in ng/mL. X-axis: Micro-organisms used for incubations with corresponding MOI (multiplification of infection) values. Baseline: Uninfected cells; Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; Ad: Adenovirus; Inf: Influenza A virus; RSV: Respiratory syncytial virus; \* p <0.05 (t-test).

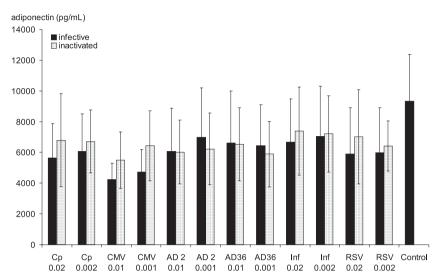


Figure 6. Adiponectin production in infected adipocytes.

Means ± SE values of 3 experiments in duplicate (n=3). ■: Adiponectin production in adipocytes incubated with infective micro-organisms. 

Adiponectin production in adipocytes incubated with inactivated micro-organisms. 

Y-axis: adiponectin production in pg/mL after log transformation. 

X-axis: Micro-organisms used for incubations with corresponding MOI (multiplification of infection) values. 

Baseline: Uninfected cells; 

Cp: Chlamydia pneumoniae; CMV: Cytomegalovirus; 

Ad: Adenovirus; 

Inf: Influenza A virus; 

RSV: Respiratory syncytial virus.

### Discussion

The present study shows that various microorganisms are able to infect adipocytes *in vitro*. Adipocyte infection resulted in an increased production of IL-6. Adenovirus infection of pre-adipocytes leads to altered PAI-1 production. These results can be viewed in the context of involvement of adipocytes in the pathogenesis of cardiovascular diseases and in relation with the development of diabetes mellitus. Compared with endothelial cells <sup>9,13</sup>, monocytes <sup>10</sup> and hepatocytes <sup>24</sup> adipocytes react in a similar way to infection with a variety of microorganisms by producing inflammatory cytokines. Abdominal adipose tissue was thought to be just a storage depot for fat, but is now considered to be an active endocrine organ able to produce several hormones and inflammatory cytokines. These hormones and inflammatory cytokines have systemic effects on the vessel wall, vascular risk factors and on insulin resistance. By the portal circulation, these hormones and inflammatory cytokines are also able to directly influence liver metabolism, potentially leading to accelerated inflammatory response, e.g. C-Reactive Protein (CRP) production, and leading to

changes in hemostasis by the production of coagulant factors and proteins involved in fibrinolysis. Specific causes for changes in adipocyte function are not well known. We propose a role for microorganisms in inducing and maintaining adipocyte dysfunction. At present, it is commonly accepted that (chronic) inflammation plays a key role in atherogenesis and atherosclerosis  $^{25-27}$ . Elevated plasma concentrations of CRP are associated with increased cardiovascular risk and are seen in patients with obesity, metabolic syndrome or type 2 diabetes  $^{28-30}$ . Microorganisms are thought to play a role in the development of atherosclerosis  $^{31,32}$  although treatment of Cp with antibiotics did not lead to a reduction in the occurrence of vascular events  $^{33}$ . The absence of an effect of antibiotic treatment on vascular events may be due to ineffective treatment strategies not affecting Cp in circulating monocytes  $^{34}$ . Also, it can still be argued that viral infections, not affected by macrolide treatment, are involved in atherogenesis, arterial thrombosis and plaque rupture  $^{35,36}$ .

Recently, adenovirus infections were considered to play a role in the development of obesity and are associated with plasma lipid abnormalities <sup>15,17,18,37,38</sup>. In the present study it is shown that a range of different viral and bacterial micro organisms are able to infect human adipocytes and induce an inflammatory response. The production of IL-6 by adipocytes increased in a dose and infection dependent manner.

Circulating TNF- $\alpha$  and IL-6 plasma concentrations are mildly elevated in obesity. Both TNF- $\alpha$  and IL-6 modify insulin sensitivity by interfering with intracellular insulin signaling pathways <sup>39</sup>. The production of TNF- $\alpha$  and IL-6 by infected abdominal adipocytes and their release into the portal circulation, may lead to increased production of CRP by the liver and influence lipid metabolism leading to an increased cardiovascular risk <sup>40,41</sup> and increased risk for the development of type 2 diabetes <sup>42,43</sup>. The inflammatory response of adipocytes upon infection may also have paracrine effects by affecting the function of neighboring adipocytes and may attract macrophages into adipose tissue, increasing the capacity for the production of inflammatory mediators. The production of the proinflammatory cytokines IL-6, TNF- $\alpha$  and MCP-1 is unregulated in a co-culture with 3T3-L1 adipocytes and a macrophage cell line RAW264 <sup>44</sup>. It has been suggested that one third of circulating plasma IL-6 is secreted by adipose tissue <sup>45,46</sup> reaching systemic concentrations high enough to elicit biological effects <sup>47,48</sup>.

Adenoviruses were the first pathogens associated with human obesity. SMAM-1 was the first (avian) adenovirus described to be positively related to obesity <sup>37</sup>, and Ad36 was the major adenovirus subtype reported to enhance adipocyte differentiation and having an association with obesity <sup>15,37</sup>. The human adenovirus types 5, 31 and 37 also appear to

increase pre-adipocyte differentiation, although their association with obesity is still unclear <sup>16,22,38</sup>. In the present study we showed that the Ad36 virus was able to increase IL-6 production by pre-adipocytes and adipocytes. Interestingly, the Ad2 subtype did not exert any functional effects on (pre-) adipocytes, indicating a specific mechanism of action in adipocyte signaling for the Ad36 virus type.

It has been suggested that enhanced differentiation of cultured pre-adipocytes by Ad36, but not Ad2, may contribute to adipogenesis <sup>49,50</sup> This would implicate that Ad-36 is associated with increased body weight. This was confirmed by showing that the antiviral agent cidofovir attenuated differentiation induced by Ad-36 through altering gene expression in adenovirus infected 3T3-L1 cells <sup>49</sup>. In mice, treatment with cidofovir significantly suppressed plasma IL-6 concentrations after intraperitoneal infection with cowpox <sup>51</sup>. Still, it remains unclear whether reduction of IL-6 concentrations by cidofovir treatment also leads to subsequent weight loss.

We acknowledge some limitations of the present study. For all experiments a stable culture of human subcutaneous cells from one obese donor was used. This culture was not continuous. It is not known whether the results from these adipocytes can be extrapolated to cells derived from other donors and from different body depots. Adipocytes beyond passage 4 had a reduced capacity for differentiation and had changed capacity for producing adipokines. This indicates that adipocyte cultures are very sensitive to functional changes. In our experiments a differentiation grade of approximately 75% was reached. Not reaching full differentiation was not considered to be an issue because adipose tissue in vivo consists of 30-60% adipocytes. Others already suggested differences in the capacity of adipokine production by preadipocytes compared to mature adipocytes <sup>52-54</sup>. We demonstrated that adiponectin was mainly produced by adipocytes, while IL-6 and PAI were produced by both adipocytes and pre-adipocytes, but pre-adipocytes produced more IL-6 and PAI than adipocytes.

In conclusion, various microorganisms are able to infect human adipocytes in vitro. Infection of adipocytes elicited increased IL-6 production. This pro-inflammatory effect of adipocyte infection may contribute to the development of cardiovascular diseases and/or the development of diabetes mellitus.

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Intracellular infections
enhance IL-6 and PAI-1
production by co-cultivated
human adipocytes and
THP-1 monocytes

## **Abstract**

#### Introduction

Obesity is associated with a chronic inflammatory state, and adipocyte dysfunction is thought to play a crucial role in this. Infection of adipose tissue may trigger the production of inflammatory cytokines, leading to increased recruitment of macrophages into adipose tissue, which in turn may exacerbate the inflammatory state in obesity.

#### Methods

Low-grade inflammation was mimicked in an *in vitro* co-culture model with human adipocytes and THP-1 monocytes. Adipocytes and monocytes were infected with adenovirus, cytomegalovirus (CMV), or influenza A (inf A) virus. After 48 hours, transinfection was evaluated and interleukin (IL)-6, tumor necrosis factor (TNF)- $\alpha$ , adiponectin and plasminogen activator inhibitor (PAI)-1 were measured.

#### Results

IL-6 production was up regulated in co-cultures of uninfected adipocytes and THP-1 macrophages in a THP-1 cell number-dependent fashion. IL-6 production by CMV-infected adipocytes was increased relative to that of uninfected adipocytes (p<0.01). IL-6 production by CMV-infected co-cultures, was 16-37 fold higher than that of uninfected adipocytes,(p<0.001). IL-6 production in influenza A-infected co-cultures was increased 12- to 20-fold (p<0.05). Only CMV-infection increased levels of PAI-1 in co-cultures (4-fold, p<0.05). Soluble factors produced by THP-1 macrophages rather than by adipocytes were responsible for the increased production of IL-6 in co-cultures.

#### **Conclusions**

Infection of co-cultivated human adipocytes and THP-1 monocytes with CMV or influenza A led to increased production of IL-6 and PAI-1. Thus infection of adipose tissue evokes an inflammatory response, leading to adipose tissue dysfunction and subsequent overproduction of IL-6 and PAI-1. This may further compound the atherogenic effects of obesity.

**Key words:** adipocytes, macrophages, co-culture, inflammation, IL-6, plasminogen activator inhibitor, cytomegalovirus, influenza.

## Introduction

Abdominal obesity is an important risk factor for the development of insulin resistance, metabolic syndrome, diabetes mellitus type 2, and atherosclerosis <sup>1, 2</sup>. Adiposity is also associated with a state of low-grade chronic inflammation. Adipose tissue is composed of different cell types, including adipocytes and macrophages 3, both of which are involved in lipid storage and cytokine secretion. Inflamed adipose tissue that has been invaded by macrophages produces interleukin-6 (IL-6) and tumor necrosis factor (TNF)-α, thereby contributing to the development of insulin resistance and cardiovascular diseases <sup>4-7</sup>. Moreover, the increase in adipose tissue mass may lead to hypoxia, which may stimulate the invasion of T-lymphocytes 8 and macrophages 9, <sup>10</sup>. Although the trigger for adipose tissue inflammation is not yet known, it may be viral infection. The inflammatory response of adipocytes to infection may stimulate the production of inflammatory cytokines, which in turn may exert paracrine effects on neighboring cells and attract more macrophages into adipose tissue. This further increases the capacity of adipose tissue to produce inflammatory mediators 11, 12. In addition, viruses and bacteria can also exert direct atherogenic effects following infection of the vascular wall 13-16.

Although a viral etiology of obesity remains unproven and controversial, adenoviruses may be causally related to obesity, suggesting a role for these pathogens in the etiology of obesity <sup>17-21</sup>. Previously, we showed that human adipocytes could be infected by several microorganisms *in* vitro, and that infection led to an increased production of IL-6. These results suggested that viral infections could contribute to the development of type 2 diabetes and atherosclerosis <sup>22</sup>. In the present study, we investigated the effects of infection of co-cultures of human adipocytes and macrophages by cytomegalovirus, influenza A virus, and adenovirus subtypes 2 and 36 on the production of inflammatory cytokines and plasminogen activator inhibitor-1 (PAI-1), and whether transinfection of the two types of cells occurred.

#### Methods

#### **Preparation of virus stocks**

Adenovirus subtypes 2 and 36 were propagated on human Hep-2 larynx carcinoma cells (#03-108/ATCC #CCL 23; Flow laboratories/Amstelstad BV, Zwanenburg, the Netherlands). The adenovirus 36 isolate was kindly provided by the Department of

Microbiology of the Erasmus University Rotterdam, the Netherlands. Influenza A/H1N1/ Netherlands/300/00 was propagated on LLC-MK2 Rhesus monkey kidney cells (#03-200/ATCC #CCL7; Flow laboratories/Amstelstad BV, Zwanenburg, the Netherlands) and Cytomegalovirus (CMV) was cultured on a human embryonic lung cell line. The nonadipogenic adenovirus 2 subtype (control virus for adenovirus 36), influenza A, and CMV strains were clinical isolates from patients with common respiratory infections (Department of Virology, Diakonessen Hospital Utrecht, the Netherlands). Infection of cells in cultures with adenoviruses 2, 36 and influenza A was established by indirect immunofluorescence staining with a pooled specimen screening reagent containing affinity purified mouse monoclonal antibodies directed against respiratory viruses, among which adenovirus and influenza A/B. (Bartels VRK Anti-Viral Screening Reagent, #B1029-86A; Trinity Biotech, Bray Ireland). CMV infection was identified by indirect immunofluorescent staining with anti-CMV immediate early antigen antibodies (clone E13, #11-003; Argene, Varilhes, France). The cytoplasm (RSV, influenza A) and nuclei (adenoviruses, CMV, influenza A) of infected cells display apple-green fluorescence whereas the cytoplasm of uninfected cells is red (counterstained with Evan's blue). After propagation on the appropriate cell lines in tissue culture bottles, the inoculation materials were harvested, aliquoted, and frozen at -80°C. The 50% tissue culture infective dose (TCID<sub>50</sub>) of each isolate was determined using corresponding cell lines according to the method of Reed and Munch <sup>23</sup>.

#### Cell culture

Adipocyte precursor cells were grown and stimulated to differentiate into mature adipocytes as described previously <sup>22</sup>. Briefly, human subcutaneous pre-adipocytes (ZenBlo Inc. SP-F3, Tebu-Bio BV Heerhugowaard, the Netherlands) were seeded at passage 4 in 24-well plates (Nunc, Roskilde Denmark) and grown to semi-confluence in pre-adipocyte medium (PM-1, ZenBio Inc. Tebu-Bio BV Heerhugowaard, the Netherlands) at 37 °C and 5% CO<sub>2</sub>. Cells were stimulated to differentiate by replacing PM-1 by differentiation medium (DM-2, ZenBio Inc. Tebu-Bio BV Heerhugowaard, the Netherlands), which is PM-1 supplemented with 3-isobutyl-1-methylxanthine (IBMX), dexamethasone, insulin, and a PPAR-γ agonist. After 1 week, the differentiation medium was replaced by adipocyte medium (AM-1, ZenBio Inc. Tebu-Bio BV Heerhugowaard, the Netherlands). The AM-1 medium was replaced every 3-4 days until experiments were performed.

Cells of the human leukemic monocyte cell line THP-1 (ECACC 88081201) were grown in suspension cultures in RPMI 1640-medium (21875 Gibco, Invitrogen Breda, the Netherlands) supplemented with 10% fetal calf serum, 2 mM L-glutamine, and antibiotics. Cell densities were kept between 2\*10<sup>5</sup> cells/mL and 8\*10<sup>5</sup> cells/mL. Five days prior to infection, monocytes were driven to differentiate into macrophage-like cells by addition of 50 ng/mL phorbol-12-myristate-13-acetate (PMA; Fluka 79346, Sigma Zwijndrecht, the Netherlands).

# Infection of adipocytes

Cultures containing more than 75% differentiated adipocytes were infected 13-15 days after the start of differentiation. All infections were monitored by fluorescent immunostaining, and the severity or grade of infection was determined by calculating the number of cells infected, expressed as a proportion of the total cell count. Optimal infection grades, expressed as the multiplication of infection (MOI), were determined in preliminary experiments with serial dilutions of each stock (data not shown). The rationale for using fairly low MOI values was to mimic chronic infection in vivo and to avoid causing cell death. One day prior to infection, the AM-1 medium was replaced and adipocytes and macrophages were incubated with 100 µl of each virus at MOI 0.01. As a negative control, cells were incubated with culture medium only. In addition to infection, the ability of cells to produce adipokines was determined by incubating the cells with 1 ng/mL IL-1\u00ed. All experiments were performed three times in triplicate on cells at passage 4, grown in 24-well plates. After the addition of virus, the plates were centrifuged for 1 hour at 500g and then incubated at 37 °C and 5% CO<sub>3</sub> for 48 hours, after which the culture medium was collected and frozen at -80 °C until adipokines were measured.

#### Infection of co-cultivated adipocytes and THP-1

Twenty-four hours before the adipocytes and THP-1 cells were infected, the AM-1 medium was replaced. Adipocytes and THP-1 monocytes were co-cultivated in fixed monocyte:adipocyte ratios of 1:2.5, 1:12.5, and 1:50, achieved by incubating 50,000, 10,000 and 2,500 THP-1 monocytes with 125,000 adipocytes per well. In contrast to the method of cell infection described above, in this experiment adipocytes or monocytes were infected separately by pre-incubating the cells with virus for 4 hours at 37 °C, after which the cells were washed and 1 mL of fresh AM-1 medium was added, followed by

the other cell type. Controls were uninfected adipocytes, THP-1 cells, and co-cultivated cell cultures. After 48 hours, the culture medium was harvested and frozen at -80 °C. Adipocytes and THP-1 monocytes were separated and fixed in acetone/methanol (1:1 vol/vol). To assess whether viruses are able to migrate from adipocytes to monocytes and vice versa (transinfection), viral infections in fixed cells were monitored with the appropriate fluorescent antibodies as described above. The number of fluorescent cells in four fields per view was counted and expressed as a percentage of the total cell count

# IL-6 production after incubation with conditioned medium

Incubations of fresh adipocytes and THP-1 cells with medium recovered from previously infected THP-1 or adipocytes were conducted to determine the relative contribution of THP-1 cells vs. adipocytes to increased IL-6 levels in co cultures. Thus 250  $\mu$ L of conditioned medium was added to 750  $\mu$ L of fresh AM-1 medium and added to freshly cultured cells, and 48 hours later IL-6 was measured in the culture medium, as an indicator of adipokine production. To rule out the possibility of virus being present in the conditioned media we performed immunofluorescence on the cells that were inoculated with the conditioned media.

### **Adipokine assays**

Commercially available ELISA kits were used for measuring IL-6 (Pelikine compact human IL-6 kit, # M1916 Sanguin Reagents, the Netherlands), TNF-α (Pelikine compact human TNF-alpha kit, # M1923 Sanguin Reagents, the Netherlands), PAI (PAI-1 antigen Elisa reagent kit, #TC11070, Technoclone GmbH Vienna, Austria) and adiponectin (human Adiponectin/Acrp30 ELISA kit, (#ELH-ACRP30-001, RayBiotech Inc. Norcross GA, USA / Tebu-Bio BV Heerhugowaard, the Netherlands). All assays were performed three times in duplicate. Optical density was measured with an automated plate reader (Elx800, Biotek Instruments Inc.) at 450/630 nm and data were processed with analytical curve fitting software (KC Junior, Biotek instruments Inc.).

# **Cytotoxicity testing**

Lactate dehydrogenase (LDH) was measured in culture medium to monitor cytotoxicity in infected and uninfected (control) cultures of adipocytes and/or THP-1 macrophages, using a colorimetric cytotoxicity detection assay (Cytotoxicity Detection Kit PLUS (LDH)

#04744926001, Roche Diagnostics BV, Almere, the Netherlands). The amount of color formed is proportional to the number of lysed cells and is presented as a percentage of maximal LDH release following cell lysis.

# **Data analysis**

Means and corresponding standard errors of adipokine levels in the culture medium of infected adipocytes and monocytes were compared with those in the culture medium of uninfected co-cultures. One-way ANOVA with Tukey-Kramer Multiple Comparison post tests were performed using GraphPad Prism version 5.01 for Windows (GraphPad Software, San Diego California USA). Two-way ANOVA with Bonferroni post tests on co-cultivation results were performed to evaluate differences in adipokine concentrations.

# Results

# Infection of adipocytes and THP-1 cells

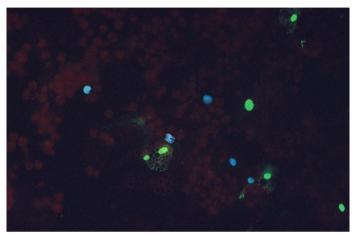
The percent of infected adipocytes cells after 48 hours incubation with adenoviruses 2 and 36, CMV, and influenza A was  $1.4\pm0.2\%$ ,  $2.8\pm0.3\%$ ,  $2.5\pm0.1\%$ , and  $0.2\pm0.1\%$ , respectively. The percent of infected THP-1 cells after 48 hours incubation with adenovirus 36, CMV, and influenza A was  $4.8\pm0.8\%$ ,  $5.2\pm0.4\%$ , and  $4.0\pm0.6\%$  respectively. THP-1 cells were refractory to adenovirus 2 [**Table 1**]. Transinfection occurred from adipocytes to THP-1 cells and vice versa. Addition of THP-1 cells to infected adipocytes resulted in infection of THP-1 cells by adenovirus 36 ( $4.6\pm1.0\%$ ),

Table 1 Transinfection in co-cultivated adipocytes and THP-1 monocytic cells

			pre-incubated adipocytes		pre-incubated THP-1	
	Adipocytes	THP-1	adipocytes	THP-1	adipocytes	THP-1
Ad 2	1.4 ± 0.2	nd	1.5 ± 0.1	nd	0.1 ± 0.0	nd
Ad 36	$2.8 \pm 0.3$	$4.8 \pm 0.8$	$2.8 \pm 0.5$	$4.6 \pm 1.0$	$1.4 \pm 0.3$	$2.6 \pm 0.4$
CMV	$2.5 \pm 0.1$	$5.2 \pm 0.4$	$2.6 \pm 0.2$	$1.8 \pm 0.7$	$1.4 \pm 0.5$	$2.3 \pm 0.3$
Inf A	$0.2 \pm 0.1$	$4.0 \pm 0.6$	$0.3 \pm 0.1$	$0.2 \pm 0.1$	$0.4 \pm 0.1$	2.5 ± 1.1

Percentages of positive cells from total in adipocytes and THP-1 cells as determined by specific immunofluoresence for each virus type. For co-cultivation either adipocytes or THP-1 were pre-incubated during 4 hours with virus before adding the other cells. All percentages derived from 4 fields per view, determined in 3 different infection experiments. MOI for each virus type: 0.01. Ad2: Adenovirus type 2; Ad36 Adenovirus type 36, CMV: Cytomegalovirus, Inf A: Influenza A virus, nd: not detected

CMV (1.8 $\pm$ 0.7%), and influenza A (0.2 $\pm$ 0.1%); adenovirus 2 infection did not proceed to infect THP-1 cells. Addition of uninfected adipocytes to infected THP-1 cells led to subsequent infection of adipocytes by adenovirus 36 (1.4 $\pm$ 0.3%), CMV (1.4 $\pm$ 0.5%), and influenza A (0.2 $\pm$ 0.1%).



**Figure 1** Dual color immunofluorescence image of transinfection of influenza A in a co-culture of adipocytes and THP-1 cells. Influenza infected THP-1 cells were co-cultured with uninfected adipocytes for 48 hours. THP-1 cells infected with influenza A demonstrate blue fluorescence (AMCA label). After 48 hours, influenza infection was transmitted from THP-1 cells to adipocytes, as indicated by apple green fluorescence (FITC label). Background: uninfected adipocytes and THP-1 cells (round) counterstained with Evans Blue (red). Lipid droplets in adipocytes are clearly visible as dark inclusions in the cytoplasm of adipocytes. (*For color figure see page 222*)

#### **Cytotoxicity**

Viral infection of co-cultures did not evoke a substantial cytotoxic reaction, as determined by the release of LDH into the culture medium. Overall cytotoxicity in infected cell cultures relative to uninfected (co-) cultures varied between 0 % and 7.4%. In cell cultures incubated for 48 hours with 1 ng/mL IL-1, cytotoxicity was maximally 11.3%.

# PAI-1 production in adipocyte/THP-1 co-cultures

After 48 hours, PAI-1 production by uninfected adipocytes was 199.8  $\pm$  82.6 ng/mL. Neither uninfected THP-1 monocytes nor virus infected THP-1 cells alone, produced PAI-1 (<2 pg/mL; data not shown). In uninfected co-cultures with high THP-1 numbers, PAI-1 production increased, but was not significant (406 $\pm$ 74,9 ng/mL; p>0.05). There were no significant effects of infection on PAI-1 production by adipocytes, THP-1 cells

or co-cultures except in the co-cultures infected with CMV and a high THP-1 cell/adipocyte ratio ( $788\pm167.3$  ng/mL; p<0.05) [**Figure 3**]. Considering the fact that THP-1 cells did not produce PAI-1, it seems obvious that adipocytes were accountable for the increased PAI-1 production, induced by CMV-infection in combination with a high number of THP-1 cells.

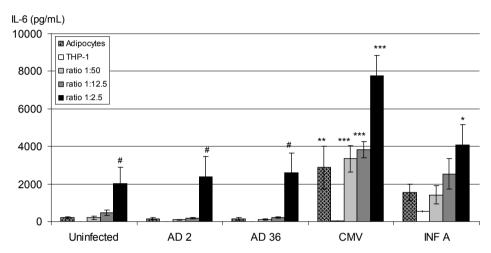


Figure 2 IL-6 production by infected adipocytes and THP-1 cell co-cultures.

Y-axis: IL-6 production in pg/mL. Bars represent control adipocytes without THP-1 cells (), control THP-1 cells without adipocytes ( $\Box$ ) and adipocytes co-cultured with THP-1 cells. Number of THP-1 cells added to 125,000 adipocytes per well and THP-1 cells/adipocytes ratio (between brackets): 2500 (ratio 1:50), 10,000 (ratio 1:12.5), 50,000 (ratio 1:2.5). X-axis: Adipocytes co-cultivated with THP-1 cells and infected with adenovirus type 2 (AD2), adenovirus type 36 (AD36), Cytomegalovirus (CMV) and influenza A (INF A) and uninfected adipocytes (uninfected). # p <0.05 compared to uninfected adipocytes. # p <0.05, # p <0.01, # p <0.01 compared to uninfected (co-)cultures.

# Adiponectin and TNF-α production in adipocyte/THP-1 co-cultures

Adiponectin production by uninfected adipocytes was 13917 $\pm$ 850 pg/mL after 48 hours. As expected, THP-1 monocytes did not produce adiponectin. Adiponectin production by adipocytes and monocytes alone or in co-culture was not affected by infection (data not shown). TNF- $\alpha$  production by uninfected adipocytes was 24 $\pm$ 6.3 pg/mL; it was 5 $\pm$ 1.9 pg/mL by uninfected THP-1 monocytes. TNF- $\alpha$  production by adipocytes and monocytes alone or in co-culture was not affected by infection (data not shown).

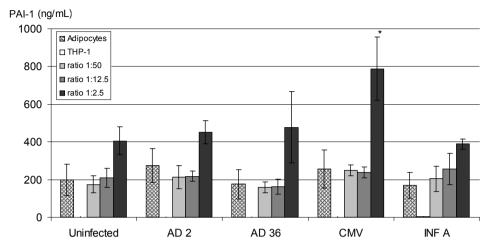


Figure 3 PAI-1 production by infected adipocytes and THP-1 cell co-cultures.

Y-axis: PAI-1 production in ng/mL. Bars represent control adipocytes without THP-1 cells (■), control THP-1 cells without adipocytes (□) and adipocytes co-cultured with THP-1 cells. Number of THP-1 cells added to 125,000 adipocytes per well and THP-1 cells/adipocytes ratio (between brackets): ■ 2500 (ratio 1:50), ■ 10,000 (ratio 1: 12.5), ■ 50,000 (ratio 1:2.5). X-axis: Adipocytes co-cultivated with THP-1 cells and infected with adenovirus type 2 (AD2), adenovirus type 36 (AD36), Cytomegalovirus (CMV) and influenza A (INF A) and uninfected adipocytes (uninfected). \*\*: p <0.05.

# IL-6 production in THP-1 cultures

After 48 hours, uninfected THP-1 cells did not produce IL-6, and neither did adenovirus 2 and 36 infected THP-1. CMV-infected THP only marginally produced IL-6:  $40\pm6$  pg/mL,  $45\pm1$  pg/mL and  $43\pm1$  pg/mL by 2,500, 12,500 and 50,000 THP-1 cells, respectively. IL-6 production by influenza infected THP-1 was  $558\pm128$  pg/mL,  $627\pm2$  pg/mL and  $551\pm43$  pg/mL by 2,500, 12,500 and 50,000 THP-1 cells, respectively (**Figure 2**).

# IL-6 production in adipocyte cultures

IL-6 production by adipocytes infected with adenoviruses 2 ( $148\pm59$  pg/mL, p>0.05) and 36 ( $133\pm67$  pg/mL, p>0.05) was similar to that of uninfected adipocytes ( $210\pm49$  pg/mL). Although IL-6 production by CMV- and influenza infected adipocytes increased to  $2877\pm1130$  pg/mL and  $1559\pm442$  pg/mL this tendency was not statistically significant (both p>0.05 compared to uninfected adipocytes) (**Figure 2**).

# IL-6 production in adipocyte/THP-1 co-cultures

IL-6 production in uninfected co-cultures with high THP-1 cells/adipocytes ratio's (1:2.5) was 2004 $\pm$ 882 pg/mL (p<0.05) compared to IL-6 production in uninfected adipocytes

(210±49 pg/mL). IL-6 production by adenovirus-infected co-cultures with high THP-1 cell numbers (THP cell/adipocyte ratio 1:2.5) was higher, but not significantly so, than in uninfected co-cultures: 2400±1075 pg/mL (adeno 2), and 2608±1056 pg/mL (adeno 36) vs. 2004±882 pg/mL (uninfected); both p>0.05). In contrast, IL-6 production was significantly increased in CMV-infected co-cultures of adipocytes and THP-1 cells (3345±715 pg/mL; p<0.01 (ratio 1:50), 3838±430 pg/mL; p<0.01 (ratio 1:12.5) and 7768±1073 pg/mL; p<0.001 (ratio 1:2.5). Influenza A infection only increased IL-6 production in co-cultures with high THP-1 numbers (4081±1073 pg/mL; p<0.05) [**Figure 2**].

The increased IL-6 production in these co cultures could be attributable to either adipocytes or THP-1 cells. To determine the relative contribution of THP-1 cells vs. adipocytes to increased IL-6 levels in co cultures, fresh adipocytes and THP-1 cells were incubated with medium recovered from previously infected THP-1 or adipocytes (conditioned media).

# IL-6 production in co-cultures after incubation with conditioned media

In **table 2** ratios are given for the relative contribution of THP-1 cells vs. adipocytes to increased IL-6 levels in co cultures. Each ratio was calculated by dividing the measured IL-6 concentration after 48 hours incubation with conditioned media by the input concentrations of the conditioned media themselves. Because the IL-6 contents of the conditioned media were very different, we corrected for these differences by expressing the quotients as a ratio indicative for the relative "de novo IL-6 production". Therefore the ratio's given in table 2 must be interpreted as the active increase of IL-6 after incubation of fresh adipocytes or THP-1 (alone or in co-culture) with conditioned media from previously infected cells [**Table 2**]. The IL-6 concentration in the conditioned medium of CMV-infected adipocytes was 870±238 pg/mL and 370±95 pg/mL in influenza A infected adipocyte conditioned medium. After a 48-hour incubation of THP-1 cells with adipocyte CMV conditioned medium, the measured IL-6 concentration in the culture medium was 1762±1042 pg/mL (p=0.36, ratio 2.0); it was 550±177 pg/mL (p=0.33; ratio 1.4) after incubation of THP-1 cells with adipocyte influenza A conditioned medium [**Table 2**].

The IL-6 concentration in the conditioned medium of CMV-infected THP-1 cells was  $11\pm2$  pg/mL and  $117\pm11$ pg/mL in influenza A infected THP-1 conditioned medium. After a 48-hour incubation of adipocytes with THP-1 CMV conditioned medium, the measured IL-6 concentration in the culture medium was  $1038\pm204$  pg/mL (p=0.01;

ratio 76.5); it was  $639\pm149$  pg/mL (p=0.03; ratio 3.8) after incubation of THP-1 cells with adipocyte influenza A conditioned medium [**Table 2**]. Conditioned medium from adenovirus 2 or 36 infected adipocytes and THP-1 did not induce any changes in IL-6 production.

Immunofluorescence on cells that were incubated with the respective conditioned media, did not demonstrate any viable virus, ruling out the possibility that viral presence could have accounted for the observed effects.

Table 2 IL-6 production after incubation with conditioned media

	uninfected	Ad 2	Ad 36	CMV	inf A
AD-CM THP	59 ± 12	66 ± 14	73 ± 11	1762 ± 1042	550 ± 177
Ratio	0.6	1.3	1.6	2.0	1.4
AD-CM co-culture	$361 \pm 4$	$310 \pm 27$	$364 \pm 31$	1490 ± 583	772 ± 79
Ratio	0.4	-1.0	0.8	1.3	1.2
THP-CM adipocytes	229 ± 38	$266 \pm 62$	242 ± 41	$1038 \pm 204$	639 ± 149
Ratio	2.4	4.7	8.6	76.5 *	3.8 *
THP-CM co-culture	$382 \pm 65$	$388 \pm 71$	365 ± 22	1224 ± 509	802 ± 191
Ratio	2.5	2.9	4.0	79.5 *	3.9 *

IL-6 concentrations (mean ± SE in pg/mL, n=2) after incubation with conditioned medium from adipocytes and THP-1 cells. Ad2: adenovirus subtype, Ad36: adenovirus subtype 36, CMV: cytomegalovirus, inf A: influenza A. AD-CM THP: adipocyte conditioned medium incubated with THP-1 cells, AD-CM co-culture: adipocyte conditioned medium added to co-cultures, THP-CM adipocytes: THP-1 cell conditioned medium added to adipocytes, THP-CM co-culture: THP-1 cell conditioned medium added to co-cultures of adipocytes and THP-1 cells. *Ratio*: de novo IL-6 synthesis expressed as ratio of measured IL-6 concentrations against IL-6 present in the conditioned medium used for incubation after subtraction of baseline values. \* indicates statistically significant (P<0.05) increase in de novo IL-6 production.

# **Discussion**

Obesity induces an inflammatory state in adipose tissue <sup>6, 10, 24</sup>. Elevated plasma concentrations of inflammatory cytokines are associated with an increased risk of metabolic syndrome, type 2 diabetes, and cardiovascular diseases <sup>25-27</sup>. The production of various (pro-atherogenic) chemokines and other factors, including angiotensin, free fatty acids, leptin, and PAI-1, by adipose tissue may contribute to the pathogenesis of these diseases. In the present study, we showed that both human adipocytes and THP-1 monocytes can be infected by adenoviruses 36, CMV, and influenza A viruses.

Adipocytes are also susceptible to adenovirus type 2, but THP-1 monocytes are refractory to this virus type. The percent of infection in monocytes by adenovirus 36 and CMV was 2-fold higher than that in adipocytes. This difference was even more pronounced for influenza A, with virus infecting more THP-1 cells than adipocytes  $(4.0\pm0.6\% \text{ in THP-1 vs. } 0.2\pm0.1\% \text{ in adipocytes})$ . Infected adipocytes transmitted the infection to monocytes and vice versa (transinfection). In general, the rate of transinfection was 50% of that of direct infection.

In this study, infections with CMV and influenza evoke an inflammatory response in co-cultivated adipocytes and monocytes, as measured by enhanced IL-6 and PAIproduction.. Circulating TNF-α and IL-6 plasma concentrations are mildly elevated in obesity <sup>28</sup>. IL-6 not only predisposes cells to become insulin resistant but also enhances the hepatic production of acute-phase proteins, such as C-reactive protein or fibrinogen <sup>29, 30</sup>. Therefore, obesity may represent a state of chronic low-grade inflammation accompanied by adipocyte dysfunction, as reflected by enhanced production of IL-6. This may also provide a link between obesity and the development of associated vascular complications such as atherosclerosis 31-33. Previously we showed that viral infection adipocytes stimulated IL-6 production <sup>22</sup>. In the present study, infection with CMV and influenza further accelerates IL-6 production in co-cultures of adipocytes and THP-1 cells. This extensive IL-6 production may be explained by the ability of infected adipocytes as wells as monocytes to produce IL-6 and that both infection and monocyte numbers significantly contributed to this. Monocytes do not constitutively produce PAI-1. Nevertheless, IL-6 and other soluble factors produced by monocytes may exert paracrine actions on adipocytes, suggesting a pathway by which adipocytes in vitro can be stimulated to produce elevated PAI-1 levels.

Our results are consistent with those of recent studies with animal cell models. In a co-culture model with murine 3T3-L1 adipocytes and murine macrophage-like cell line RAW264 (peritoneal macrophages), the production of IL-6, TNF- $\alpha$ , and monocyte chemotactic protein-1 (MCP-1) was up regulated whereas that of adiponectin was down regulated <sup>34</sup>. Also, the duration of co-culture influences the increase in TNF- $\alpha$  mRNA-expression in co-cultures of 3T3-L1 and THP-1 <sup>35</sup>. Incubation with bacterial lipopolysaccharide in a co-culture of 3T3-L1 and RAW264 did not influence TNF- $\alpha$  production, but increased IL-6 production 100-fold, suggesting an exaggerated biological interaction between macrophages and adipocytes in persistent low-grade

infection by gram-negative bacteria 36.

The increased production of IL-6 in co-cultures may be mediated by cell-cell contact, because the production of IL-6 increased with the number of THP-1 monocytes added to the co-cultures. However, it is more likely that locally produced inflammatory substances produced by dysfunctional adipocytes and adipose tissue macrophages exerted autocrine/paracrine effects on neighboring cells. Conditioned media from uninfected or virus-infected adipocytes did not stimulate IL-6 production by THP-1 cells or co-cultures, whereas incubation of adipocytes and co-cultures with conditioned medium from CMV-infected THP-1 led to a nearly 80-fold increase in IL-6 production. Infective virus was not detected in the conditioned media, indicating that there was no lytic infection and that de novo IL-6 synthesis was not a direct effect of infection. Moreover, this suggests that the observed effects depended on cross-talk between cells that was orchestrated by soluble factors produced by infected cells in the culture environment. This is consistent with the observation that the release of inflammatory cytokines in adipose tissue samples from obese subjects is mainly due to the production of cytokines by cells other than adipocytes <sup>37</sup>. Adipose tissue macrophages are able to produce extensive amounts of pro-inflammatory mediators, which could contribute to the development of insulin resistance 12. Here, we conclude that the paracrine effects of soluble factors (such as IL-6 or TNF-α) released by infected macrophages, rather than by infected adipocytes, were responsible for the increased production of IL-6 in co-cultures. The transinfection phenomenon we demonstrated may explain how infectious agents invade several parts of the body and infect cells of different organ systems, such as vessel walls and adipose tissue, thus contributing to local and systemic inflammatory states.

Microorganisms, such as CMV or influenza A, that infect adipose tissue may initiate and perpetuate a chronic inflammatory response, attracting more macrophages into the already inflamed tissue and exacerbating the total inflammatory response to the infection. The results of the present study further support the concept that both adipocytes and adipose tissue macrophages are involved in adipokine production and that their concerted actions synergistically contribute to increased production of IL-6 and PAI-1, reflecting the inflammatory state associated with obesity. Although adipocytes are the most abundant cell type in adipose tissue, the extent or recruitment of (infected) macrophages is likely to be decisive for the inflammatory state of adipose tissue. It is also conceivable that infected monocytes invade adipose tissue.

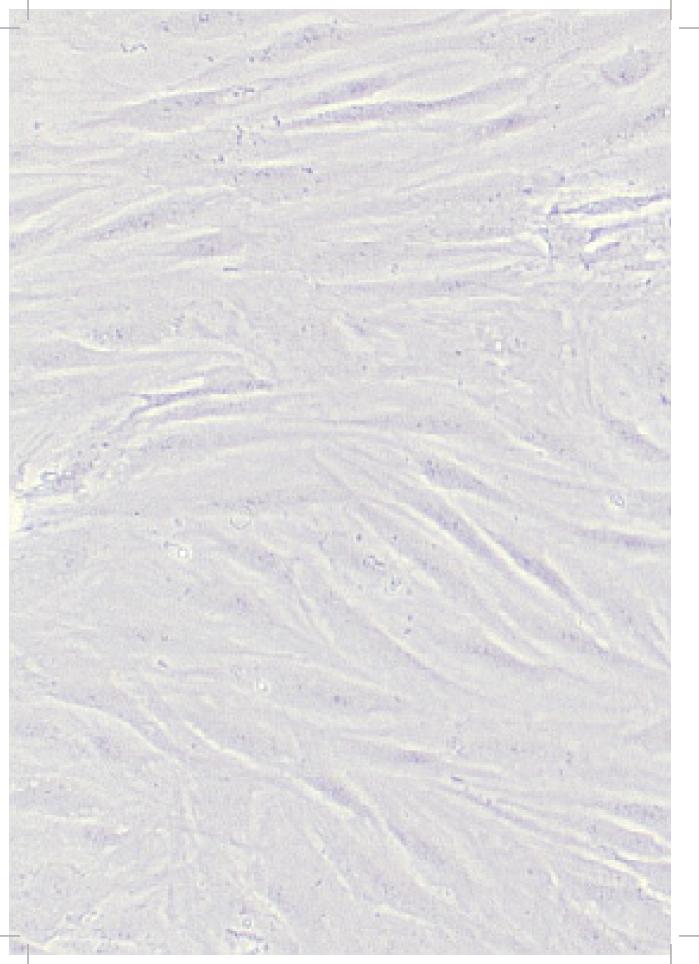
In conclusion, adipocytes and monocytes can become infected by several viruses *in vitro*, leading to an inflammatory response. Transinfection occurs from adipocytes to monocytes and vice versa. The relative contribution of monocytes to the production of IL-6 in co-cultures is likely to be higher than that of adipocytes.

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Discussion
Summary
Samenvatting
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Color figures

# Inflammation, infection, and atherosclerosis

Chronic low-grade inflammation, as is often seen in atherosclerosis, obesity, diabetes and the metabolic syndrome, is a component of many diseases. The presence of inflammation in atherosclerosis is evidenced by elevated plasma concentrations of a variety of markers, including C-reactive protein (CRP), tumor necrosis factor (TNF)-α, and several interleukins and adhesion molecules. Inflammation is critically involved in the destabilization of atheroma in the vascular wall, with the stability of arterial plaques being determined by the severity of inflammation. The origin of the inflammation, however, is unclear. Vascular damage may trigger inflammation, but most likely inflammation originates from (excessive) abdominal adipose tissue. The release of cytokines exacerbates the existing insulin resistance and endothelial dysfunction, which play a central role in the pathogenesis and development of atherosclerosis and thus promote the atherogenic process. Potential triggers for local vascular inflammation are traditional risk factors for artery disease, such as hypercholesterolemia, hypertension, diabetes, free radicals, low levels of highdensity lipoprotein cholesterol, and elevated or modified LDL. Chronic or repetitive infections may be triggers for chronic inflammation<sup>1,2</sup>. However, atherosclerosis is a multifactorial process and cannot be explained solely in terms of the traditional cardiovascular risk factors, and intracellular infections have been suggested as a direct cause of atherosclerosis. Several clinical and experimental studies suggest that various infectious agents such as Chlamydia pneumoniae (Cp) or herpes viruses may be the cause of chronic inflammation in lesions, but to date data about a direct causal relation between infections with these pathogens and atherosclerosis are conflicting. In the in vitro studies presented in this thesis, we investigated the mechanisms by which pathogens may function as etiological factors for atherosclerosis.

# Pathogens involved in atherogenesis

Cytomegalovirus (CMV) and *Cp*, in particular, have been associated with the presence and development of cardiovascular diseases<sup>3-5</sup>. Results from animal studies and *in vitro* studies show that *Cp* can modulate atheroma biology, including lipid- and inflammatory-related processes<sup>6</sup>. CMV infection of endothelial cells and macrophages has an important role in the establishment of latency and persistence, which are critical for the maintenance of CMV in the host. Infection with CMV can stimulate processes related to the development of atherosclerosis<sup>7</sup>. The replication of CMV,

however, appears to be dependent on the specific origin and tropism of endothelial cells and macrophages<sup>8</sup>. In patients with documented coronary artery disease and elevated levels of interleukin 6 (IL-6), CMV seropositivity was independently associated with future cardiac mortality. These data support the hypothesis that the atherosclerotic effects of CMV are mediated through an underlying inflammatory response<sup>9</sup>. In contrast to CMV and *Cp*, influenza virus causes more acute infections with severe (systemic) complications. Influenza induces substantial inflammation of the vessel wall and may trigger plaque destabilization, which leads to acute coronary syndromes<sup>10</sup>. In the studies presented in this thesis, we infected endothelial cells, hepatocytes, adipocytes and monocytes with *Cp*, CMV, and influenza A *in vitro*. We investigated which of the effects caused by these infections could account for the impairment of vascular function and atherogenesis.

It has been proposed that obesity represents a chronic inflammatory state that has multiple implications for cardiovascular morbidity and mortality<sup>11,12</sup>. Because adenoviruses have been associated with obesity<sup>13</sup> and may indirectly contribute to the development of cardiovascular diseases, we also infected adipocytes with adenovirus. We found that infection with adenovirus 36 increased the production of pro-inflammatory IL-6.

# Properties of intracellular pathogens

The microorganisms we used in most of our studies (CMV, *Cp*, and influenza virus) have repeatedly been shown to be associated with cardiovascular disorders. Nevertheless, these microorganisms have their own distinct properties. Although most of these viruses cause infection, we found that the effects of intracellular infections differed by virus type, indicating that infection-dependent mechanisms are virus specific. To test the specificity of the association of these viruses with cardiovascular disease, we also tested other viruses not associated with cardiovascular disease, such as parainfluenza (chapter 2), RSV (chapters 2,7), and adenovirus type 2 (chapters 7,8). We also used heat-inactivated viruses to determine whether the measured effects really were infection dependent. In some studies we found striking differences. For instance, infection of human umbilical vein endothelial cells (HUVEC) and monocytes with CMV and *Cp*, but not with influenza virus, significantly reduced the clotting time. In contrast, interleukin production was higher after infection with influenza virus than after infection with CMV or *Cp*. (chapters 2,3). In studies with adipocytes,

we demonstrated that infection with Ad36, but not Ad2, increased IL-6 production by pre-adipocytes and adipocytes (chapter 7). We also demonstrated that Ad36 readily infected monocytes whereas Ad2 was not capable of infecting monocytes (chapter 8). This may explain why Ad2 is associated with more superficial (respiratory) infections and not with systemic infections, which supports our choice of using Ad2 as a 'control virus' in this study. These differences show that infecting microorganisms adversely affect cell function via distinct mechanisms in a tissue-dependent manner.

Infective progeny virus may invade cells involved in atherogenesis and be disseminated to other cells. Intracellular infections *in vivo* may become latent or even persistent if the viral genome integrates into a host chromosome and only few viral proteins are synthesized because viral genes are not, or poorly, expressed. Even in such situations, viruses may still be capable of having a deleterious effect on cells. For example, CMV mimics host proteins and may behave as a molecular pirate, using clever strategies to gain cell entry, manipulate host gene regulation, and cause immune evasion<sup>14</sup>. Highjacking the cellular machinery alters cellular function and may result in cellular dysfunction and inflammatory reactions, which are a hallmark of atherosclerosis.

# Infection of various cells by various pathogens leading to different cellular responses: implications for atherogenesis

Infections with the microorganisms used in the experiments described in this thesis are common. One of the ports of entry is the respiratory tract, and community-acquired respiratory infections with multiple microorganisms are common. Even though respiratory infections may be seasonal, clinical studies suggest that acute respiratory tract infection may be a risk factor for acute coronary syndromes <sup>15,16</sup>. During influenza epidemics, the mortality rate for cardiovascular syndromes increases <sup>17,18</sup>. The number of deaths due to ischemic heart disease, hypertension, and cerebrovascular disease peaks 2 weeks after the influenza infection peaks <sup>10,19</sup>. Atherosclerosis however, is a chronic disease that begins in early life with the formation of fatty streaks and may progress to cause severe adverse effects in adult life. Individuals are infected with a variety of microorganisms during their lifetime. Viruses replicate in the epithelial cells of the respiratory tract and release infective virus in the vicinity of alveoli. In this way, neighboring endothelial cells in vessel walls aligning lung tissue may become infected. It is highly likely that virus can spread hematogenously, either as free virus (viremia) or in macrophages. In this way, a virus can be transported to remote tissues

and directly infect healthy endothelium and atherosclerotic plaques. It is not known whether viruses infect plaques but several studies have shown the presence of viral sequences in cells of atherosclerotic plaques<sup>20-25</sup>. An important question is whether microorganisms need to enter the vessel wall in order to elicit immune responses. Local inflammatory effects could induce arterial wall inflammation and thereby trigger plaque destabilization or even rupture of vulnerable plaques. In this way, what starts as local vessel wall inflammation could develop into systemic (low grade) inflammation. In our studies, we demonstrated that various microorganisms are able to infect endothelial cells (chapters 2, 5 and 6), monocytes (chapters 3 and 8), hepatocytes (chapter 4), and (pre-)adipocytes (chapters 7, 8). Intracellular infection of these cells in vitro leads to cellular dysfunction and initiates procoagulant activity, hypofibrinolysis, impaired cellular function, and inflammation, as reflected by increased expression of pro-inflammatory cytokines. This may contribute to local and/ or systemic inflammation. There was a remarkable resemblance in how monocytes, hepatocytes, endothelial cells, and adipocytes reacted to infection, with the activation of processes that could be involved in atherogenesis. This suggests that not only direct infection of the vessel wall but also indirect infection by transmission of viruses to tissues elsewhere in the body may initiate a systemic inflammatory and procoagulant state.

In a co-culture model, we demonstrated that viruses were able to disseminate from infected adipocytes to monocytic cells and vice versa (chapter 8), a phenomenon that we termed 'transinfection'. This phenomenon may be quite similar to the ability of viruses to infect other cell types after a primary infection. This is in line with the demonstration that monocytes were able to transmit infectious CMV in a co-culture model of peripheral blood mononuclear cells and CMV-infected endothelial cells. Thus the endothelium and circulating monocytes have an interactive role in the dissemination of virus<sup>26</sup>. Taken together, these findings provide a mechanism by which viruses are able to reach various tissues. Moreover, they show that infectious agents can invade several parts of the body and infect cells of different organs, such as the vessel wall and adipose tissue, contributing to local and systemic inflammatory states (**Figure 1**).

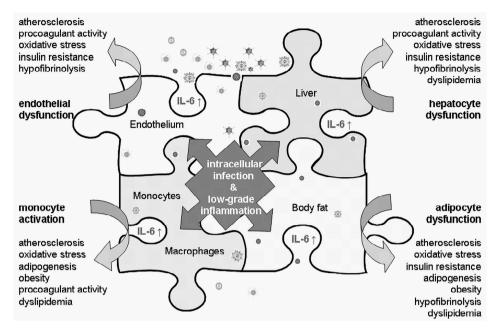


Figure 1 The concept of intracellular infections in the etiology of vascular diseases.

A variety of microorganisms are able to infect cells and organs *in vitro*. Intracellular infection of endothelial cells, liver cells, fat cells and monocytes/macrophages leads to cellular dysfunction, as reflected by procoagulant activity, hypofibrinolysis, and (chronic) local or systemic low-grade inflammation. Impaired cellular function with subsequent inflammation is reflected by increased expression of pro-inflammatory cytokines. Increased IL-6 production is a predominant key feature of an altered secretion profile in infected cells and may be considered as a marker for cellular dysfunction. Endothelial, monocyte, hepatocyte, and adipocyte dysfunction caused by intracellular infections may contribute to local and systemic inflammation, with subsequent effects on vascular function that may contribute to the development of cardiovascular diseases. In addition, cross-talk between cells of the immune system, cells of remote tissues (liver, fat), and the vessel wall, mediated by cytokines and other pro-inflammatory substances, is responsible for the cascade effects leading to exacerbation of vascular disorders.

# Aspects of in vitro culture experiments

The methods we used to infect cells *in vitro* may not mimic the mechanisms by which cells become infected *in vivo*. We acknowledge this as a limitation of our studies but our aim was to mimic chronic low-grade intracellular infection, consistent with the concept of atherosclerosis being a (low-grade) chronic process. In this respect, it was important to avoid lytic infection, which could overrule the specific effects initiated by infection. We always carefully monitored cytotoxicity, and the severity of infection was calculated and monitored by immunofluorescence. Obviously, infections rates

were dependent on the nature and strain of the microorganism involved and on the susceptibility of cells to become infected. This was also illustrated in a study in which CMV infection resulted in a lytic infection in brain microvascular endothelial cells but not in aortic macrovascular cells. This could indicate that aortic macrovascular cells act as a reservoir of virus in the host<sup>8</sup>. Other experimental data show that RNA from influenza virus is present in atherosclerotic plaques but not in undamaged vessel<sup>27</sup>. This tropism may also explain how pathogens become internalized, especially into atherosclerotic plaques. It is difficult to develop an experimental model of persistent infection because, for example, while chemicals can be used to suppress the replication of intracellular organisms, they will also influence or stop cellular metabolism.

# Detection of microorganisms involved in atherosclerosis and vascular disease

Given the nature of persistency, it is conceivable that the role of intracellular pathogens in atherogenesis is underestimated in clinical studies. A lack of standardized methods and inconsistent use of diagnostic procedures may give rise to discrepant results<sup>28</sup>. There is no "gold standard" for diagnostic methods to detect infection. Viral cultures, immunoassays, serologic and PCR methods may be insensitive, and performance and criteria are not always well defined. These considerations are important because inadequate detection may lead to ineffective treatment. Moreover, treatment may be inadequate if a single infection is targeted, especially because the effectiveness of treatment is probably determined by the pathogen burden. The importance of pathogen burden was clearly demonstrated in a study involving consecutive patients undergoing elective carotid endarterectomy. The *Cp* burden was associated with plaque expression of IL-6, which in turn was associated with serum levels of IL-6 and CRP<sup>29</sup>. Moreover, multifactorial disease processes, such as those involved in atherosclerosis, are unlikely to be activated by a single microbe or vascular risk factor. Several studies indicate that infections with multiple pathogens are common<sup>30</sup> and could contribute to coronary artery disease by precipitating inflammation and endothelial cell injury<sup>31,32</sup>. Increasing pathogen burden is also associated with an increased risk of myocardial infarction and death in patients with known coronary heart disease<sup>33-35</sup>. Therefore, the cumulative effects of simultaneous infections at various sites and/or repetitive infections may contribute to the pathogenesis and progression of atherosclerosis<sup>36,37</sup>.

# Evidence for a causal role of infections in atherogenesis?

The role of infections in endothelial injury and vascular wall inflammation has been studied extensively but evidence for a causal relation is still lacking<sup>2,4,38,39</sup>. Several infectious agents have been investigated but none has proved to play a causative and specific role. Cp has been studied the most extensively, but the results of large clinical trials of antibiotics against this disease have on the whole been disappointing<sup>40,41</sup>. Nevertheless, certain infections are now considered to be a risk factor for atherosclerosis development, but current data do not allow us to determine whether infection is a cause or a co-factor in atherogenesis. Various criteria have been proposed to help discriminate between causal and non-causal associations. Atherosclerosis has been proposed as an infectious disease, but there is little evidence for this according to Koch's postulates<sup>42</sup>. Koch stated that the host infection must lead to disease and that the pathogen must be isolated from the infected host and should be present in all infected persons. However, it may not be correct to apply these criteria to chronic diseases. Apart from the fact that atherosclerosis is multifactorial disorder, the main characteristics of chronic infections are precisely that microorganisms are latent or persistent and may be hard to culture, if viable at all. It is possible that viral proteins are synthesized de novo, but that there is no or little productive infection. Therefore, some of Hill's criteria for causality may be more applicable for determining the causality of infection in atherosclerosis<sup>43,44</sup>. The results of the studies presented in this thesis meet most of these criteria, namely, strength of association, consistency, coherence, biologic plausibility, and doseresponse relationship. An overview of the results from our studies shows that infection of various cell types consistently resulted in pro-inflammatory and procoagulant activity, hypofibrinolysis, and cytokine production, and that these effects were time and dose dependent. In experiments with Cp, treatment of hepatocytes (chapter 4) and endothelial cells (chapter 5) with macrolide antibiotics markedly decreased the inflammatory effects of the infection. As for biologic plausibility, the procoagulant effects of influenza infection in vitro are consistent with the clinical observation of coagulation disorders as complications of influenza. Causality between infections and occurrence of vascular diseases is difficult to establish in clinical studies. In the present thesis, several in vitro studies are presented showing that various infections are able to trigger cellular responses involved in atherogenesis. We therefore propose a causal role for intracellular pathogens in the etiology of vascular diseases.

## Involvement of immune cells in atherogenesis

Macrophages are central mediators of cellular innate and adaptive immunity and are essential to the initiation and progression of vascular lesions. Once activated, they initiate the oxidation of LDL and rapidly take up oxidized LDL through specific scavenger receptors, leading to foam-cell formation. Activated macrophages also secrete a variety of pro-inflammatory products that affect lesion progression and plaque stability<sup>45</sup>. Although immune responses may rely mainly on tissue-resident macrophages, the infiltration of activated monocytes (and their differentiation into macrophages) into tissues is markedly increased under inflammatory conditions. Oxidized LDL and locally secreted chemokines affect the chemotactic migration of monocytes and T-cells, but not neutrophils, into the subendothelial space. In atherosclerosis, the recruitment of monocytes is essential for lesion formation. T- and B-cells may also have proather ogenic effects but are probably not obligatory for lesion initiation or progression. In the studies described in chapter 3 and chapter 9, we showed that monocytes and macrophages are susceptible to infection and can be infected directly or via transinfection. The observation that monocytes may also function as vehicles for delivering infectious agents to tissues (vessel wall, liver, abdominal fat) contributes to our understanding of the increased tissue inflammation seen in obesity. We and others have demonstrated that obesity is accompanied by inflammation, as evidenced by the overproduction of several adipokines<sup>46</sup>. Increased macrophage recruitment into various tissues may exacerbate the total inflammatory response to the infection and may have potent proatherogenic effects in the vasculature. Crosstalk between cells of the immune system, cells of other tissues (liver, fat), and the vessel wall, mediated by cytokines and other proinflammatory substances, is responsible for initiating a cascade of effects leading to vascular disorders. We conclude that monocyte and/or macrophage activation plays a central role in infection-induced inflammation, potentially contributing to atherogenesis.

# Can *in vitro* experimental results be extrapolated to the *in vivo* situation?

While most suspected infectious agents have an atherogenic effect by initiating or aggravating a chronic vascular or systemic inflammatory process, influenza may have a rather different effect by triggering destabilization of already vulnerable plaques<sup>47</sup>. However, the current body of evidence is not conclusive about the causal relationship

between pathogens and cardiovascular disease. Animal and pathological studies probably provide the best evidence that infections have effects *in vivo* similar to those seen in atherosclerosis. Several studies have shown that *Cp* induces atherosclerosis in animals<sup>48,49</sup>. Moreover, inoculation of chicken with Ad 36 virus induces adiposity<sup>50,51</sup>. This may be associated with inflammation and subsequent vascular complications. Evidence that CMV causes atherosclerotic disease comes from studies of the development of atherosclerosis in transplant recipients<sup>52-54</sup>. CMV-negative recipients of CMV-positive donor hearts have an impaired epicardial endothelial function and an increased incidence of cardiovascular-related events and death during follow-up. CMV infection may contribute to allograft failure by accelerating coronary endothelial dysfunction<sup>55</sup>.

In clinical studies, it is difficult or even impossible to distinguish between cause and effect. Even intervention trials do not confirm the role of pathogens in vascular disease. It is possible that pathogens are not associated with atherosclerosis at all, but the overall results of clinical and laboratory studies suggest that there is such a relation. The inconclusive results of intervention trials are probably due to the methodological shortcomings, such as underpowering, transient short-term effects, and low-dosage and low-duration regimens. Moreover, treatment of patients with cardiovascular diseases with conventional antibiotics may be ineffective because microorganisms become persistent or less susceptible to antibiotics.

The validity of using *in vitro* study results to explain *in vivo* events could be questioned. Cultured cells may not respond to infection in the same way as cells cultured from tissue explants or primary cells. Moreover, conclusions based on *in vitro* findings cannot necessarily be extrapolated directly to the *in vivo* situation, and clinical implications should be interpreted with caution. While the methods currently used for the culture and propagation of *C. pneumoniae* may not be analogous to the mechanisms by which the infection is propagated *in vivo*, the results of our *in vitro* studies suggest that antibiotic treatment of infection-induced atherosclerosis may improve endothelial function by restoring normal cell function. This may be the case not only *in vitro* but also *in vivo*. Another *in vitro* study indicated that continuous *C. pneumoniae* infections may more closely resemble the events that occur *in vivo* and, therefore, may be a good model for the *in vitro* study of *C. pneumoniae* infection. In the same study macrolides did not completely eliminate the organism. This may be important because the failure of antibiotic therapy against *C. pneumoniae* infection in humans has been reported<sup>56</sup>. *In vitro* studies, such as those presented in this thesis, remain essential for providing

insight into the mechanisms by which microorganisms exert potential adverse effects on the vasculature

# **Concluding remarks**

In conclusion, the results of the studies presented in this thesis suggest that intracellular infections have a causal role in the etiology of vascular diseases. We suggest that intracellular microorganisms should be added to the list of factors that induce endothelial dysfunction and contribute to the development of atherosclerosis. We identified inflammation, procoagulant activity, and hypofibrinolysis to be major outcomes of intracellular infection of different types of cells. The *in vitro* studies presented in this thesis identified at least some of the mechanisms by which intracellular infections may contribute to the development of cardiovascular diseases.

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9

Discussion
Summary
Samenvatting
Dankwoord
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Color figures

Chronic disorders, such as atherosclerosis, are currently considered to be associated with viral infections. Influenza virus, Cytomegalovirus (CMV), and Chlamydia pneumoniae (Cp), in particular, have been associated with cardiovascular disorders. However, it is unclear whether these infections cause these disorders or merely exacerbate existing processes. Severe complications can occur in acute infectious diseases such as influenza. For example, in the Netherlands about 2000 people die annually of mainly cardiovascular complications occurring secondary to influenza. Various factors, including viral infections, are thought to initiate or accelerate the process of atherosclerosis. When plaques rupture and their contents pass into the bloodstream, various systemic processes are evoked, such as enhanced coagulation and inflammatory responses. Chronic infections may also have a role in vessel wall damage and loss of the constitutionally protective properties of endothelial cells. Although the hypothesis that infections may underlie cardiovascular disorders is widely accepted, definite evidence is still lacking. In the studies described in this thesis, we performed several in vitro studies to investigate the hypothesis that cardiovascular disorders have an infectious etiology.

In the study described in **chapter 2**, we demonstrated that respiratory viruses (influenza A, influenza B, parainfluenza-1, respiratory syncytial virus, adenovirus, and CMV) infect lung fibroblasts and human umbilical vein endothelial cells *in vitro*. As determined in a one-stage clotting assay, all viral pathogens induced procoagulant activity in infected endothelial cells, decreasing the clotting time by a mean of 55%. When factor VII-deficient plasma was used, the clotting time was not reduced. These results indicated that the reduction in clotting time was factor VII dependent and therefore mediated by the extrinsic coagulation pathway. The induction of procoagulant activity was associated with a 4- to 5-fold increase in the expression of tissue factor (TF), as measured by the generation of factor Xa. Thus viral infection is accompanied by an increased expression of TF on endothelial cells. We concluded that both enveloped and non-enveloped respiratory viruses are capable of infecting cultured human endothelial cells, causing a shift from anticoagulant to procoagulant activity associated with the induction of TF expression.

The production of cytokines by monocytes may accelerate the process of atherosclerosis and may contribute to vascular disorders by eliciting plaque instability. Whether virus-infected monocytes initiate coagulation and produce (pro-) inflammatory cytokines

was investigated in the study described in chapter 3. For this study, we designed an improved method for isolating platelet-free monocytes. Freshly isolated human monocytes were incubated with CMV, Cp, and influenza A. The clotting time of CMVand Cp-infected monocytes was reduced by 60%. Addition of factor VII-deficient plasma normalized the clotting time. As with endothelial cells, the clotting time of monocytes was factor VII dependent and therefore mediated by the extrinsic coagulation pathway. The initiation of coagulation by virus-infected monocytes was a result of the expression of TF. By using factor VII-deficient plasma and anti-TF antibodies, we showed that TF plays an important role in virus-induced monocytic procoagulant activity. This could be a direct effect of virus but could also be triggered by IL-6 produced by monocytes. By producing IL-6, infected monocytes could stimulate uninfected cells to express TF. Although infection with influenza A only marginally influenced clotting times, it strongly stimulated the production of IL-6 and IL-8. In contrast, the infection of monocytes with CMV and Cp only modestly stimulated the production of IL-6 and IL-8. During viral infections, monocytes predominantly produce the inflammatory cytokines IL-6 and IL-8. Interleukin-10 is detected predominantly in atherosclerotic lesions, but monocytes may produce this cytokine transiently. In this study, we did not find virus-infected monocytes to produce IL-10. The balance between proinflammatory and anti-inflammatory cytokines in the atherosclerotic plaque may be decisive for the progression of the lesion. Our results indicated that only small amounts of an infectious virus are needed to stimulate monocytes to exert considerable immunological effects. We concluded that the infection of monocytes by viruses triggers proatherosclerotic and prothrombotic processes in vitro.

The liver has multiple metabolic, endocrine, and secretory functions and is an important source of a variety of proteins, among which acute-phase proteins, coagulation factors, complement factors, cholesterol, and triglycerides. Inflammation caused by infection or tissue damage stimulates the secretion of inflammation-associated cytokines, including IL-1, IL-6 and tumor necrosis factor (TNF)-alpha. These cytokines stimulate hepatocytes to increase the synthesis and release of positive acute-phase proteins, including C-reactive protein (CRP). IL-6, also called hepatocyte-stimulating factor, is the major stimulus for CRP production. Fibrinogen is involved in hemostasis by stabilizing the fibrin matrix and is synthesized by hepatocytes under the control of inflammatory cytokines. In the study described in **chapter 4**, we reported that human HepG2 hepatocytes could be infected with the intracellular pathogens *Cp and* 

CMV. We also investigated whether azithromycin was able to inhibit the infection of human hepatocytes by *Cp* and reduce the inflammatory response of hepatocytes. We showed that *in vitro* infection of hepatocytes with *Cp* led to time and dose-dependent production of IL-6 and that this inflammatory response was strongly and significantly inhibited by azithromycin, confirming the antibacterial properties of azithromycin against *Cp*. Fibrinogen production was increased by CMV and *Cp*, but azithromycin was not able to suppress this CMV- or *Cp*-induced production of fibrinogen. Apparently, the effects of azithromycin on *Cp*-infected hepatocytes are due to its antimicrobial properties and not to its anti-inflammatory properties.

Chronic infection could contribute to atherogenesis by a number of mechanisms, such as direct vascular injury and induction of a systemic inflammatory state, both of which lead to endothelial dysfunction. Nitric oxide (NO) is a key regulator of endothelial function. Under pathological conditions, uncoupling of endothelial nitric oxide synthase (eNOS) leads to vessel damage and endothelial dysfunction as a result of the production of oxygen radicals instead of NO. Excessive formation of reactive oxygen species (ROS) leads to inflammation, endothelial dysfunction, and increased adhesion of leukocytes. In the study described in **chapter 5**, we measured the *in vitro* production of eNOS, cyclic quanosine monophosphate (cGMP, as a surrogate for NO), and ROS by human umbilical vein endothelial cells (HUVEC) infected with Cp or CMV. We hypothesized that infection-induced atherosclerosis could be initiated by changes in NO metabolism and that this effect could be reversed by azithromycin treatment. eNOS, cGMP, and ROS production were attenuated in Cp-infected HUVEC, whereas infection with CMV did not change cGMP or ROS production and only marginally decreased eNOS production. Co-incubation with azithromycin restored eNOS, cGMP, and ROS production by Cp-infected HUVEC to levels comparable with those of uninfected HUVEC. These results suggested that antibiotic treatment for infectioninduced atherosclerosis improves endothelial function.

Increased expression of endothelial adhesion molecules, such as vascular cell adhesion molecule-1 (VCAM-1), intercellular adhesion molecule-1 (ICAM-1), and endothelial selectin (E-selectin), are markers of endothelial dysfunction. Given the role of iron in a wide range of infection processes, the presence of iron may complicate the infection-mediated activation of endothelial function. Iron has been found to accumulate in human atheroma, and several experimental studies have shown that iron chelation

or iron-deficient diets lead to a reduced formation of early atherosclerotic lesions. In the study described in **chapter 6**, we showed that Cp and CMV infection induced the expression of ICAM-1 and VCAM-1 and up-regulated the expression of E-selectin. The expression of adhesion molecules in HUVEC infected with Cp was increased by incubating the cells in iron-rich medium. Thus the endothelial response to chronic infections depends on the intracellular level of iron. However, the modulating effect of iron is not due to an increase in the infectivity or replication of the microorganism because the endothelial response was observed before Cp infection or replication was noticed. We also observed that both chelation and radical scavengers counteracted the effects of iron. These findings suggested that the main effect of iron is to prime endothelial cells by causing oxidative stress. Iron-primed endothelial cells may be more responsive to the paracrine effects of endothelial cell infection. In contrast, the level of intracellular oxygen-derived radicals in infected HUVEC was unchanged, indicating that the induction of ICAM-1 and VCAM-1 was not driven by infection-related radical formation. This suggested that infections exert their effects on endothelial activation through a pathway that is different from that for the formation of oxygen-derived radicals.

Abdominal adipose tissue not only absorbs and stores free fatty acids but also produces and secretes adipokines such as IL-6, TNF-alpha, PAI-1, and adiponectin. These factors may have effects on the arterial wall, on vascular risk factors, and on insulin resistance. By producing hormones and inflammatory cytokines, adipose tissue may have an important role in the development of diabetes and atherosclerosis. It has been suggested that obesity is a chronic inflammatory state in which adipocyte dysfunction has a crucial role. In the study described in chapter 7, we showed that several microorganisms were able to infect human adipocytes and pre-adipocytes in vitro. Infection of adipocytes with Adenovirus type 36, but not with Adenovirus type 2, led to increased production of IL-6. This may contribute to chronic low-grade inflammation, a process known to be involved in the development of cardiovascular diseases and type 2 diabetes. These findings support a potential role for microorganisms in inducing and maintaining adipocyte dysfunction. We also demonstrated that adiponectin was exclusively produced by adipocytes and not by pre-adipocytes and that IL-6 and PAI-1 were mainly produced by pre-adipocytes. Thus during their passage through the portal circulation, inflammatory substances are able to directly influence liver metabolism, potentially leading to an accelerated inflammatory response, e.g. CRP production, and to changes in hemostasis as a result of the production of coagulant factors and proteins involved in fibrinolysis.

The inflammatory response of adipocytes to infection may also have paracrine effects by influencing the function of neighboring adipocytes and may attract macrophages into adipose tissue, increasing the capacity for the production of inflammatory mediators. In the study described in **chapter 8**, we tested this concept in an *in vitro* coculture model of human adipocytes and macrophage-like differentiated human THP-1 cells. We first showed that transinfection occurred, namely, that infected adipocytes infected macrophages and vice versa. This phenomenon may explain how infectious agents can infect cells of various organ systems, such as vessel walls and adipose tissue, thus contributing to local and systemic inflammatory responses. Levels of IL-6 and PAI-1 were increased in co-cultures of human adipocytes and THP-1 macrophages infected with CMV or influenza A virus. The invasion of adipose tissue by microorganisms such as CMV and influenza A may initiate and perpetuate a chronic inflammatory response, attracting more macrophages into the already inflamed tissue, thereby exacerbating the total inflammatory response to the infection. Although adipocytes seem to be the major source of bioactive cytokines, the extent of macrophage recruitment may be decisive for the severity of inflammation. However, it is conceivable that infected monocytes invading adipose tissue trigger an inflammatory response. In addition, we showed that primarily macrophages were responsible for the increased levels of IL-6 measured in infected co-cultures. This response provides insight into how adipose tissue signals to remote tissues (endothelium, liver, muscle) and may provide a pathological link between obesity and its associated cardiovascular complications.

#### From the results presented in this thesis, we conclude:

- Intracellular infections with various pathogens, such as respiratory viruses, *Chlamydia pneumoniae (CP)*, and cytomegalovirus (CMV), induce procoagulant activity in endothelial cells and monocytes *in vitro*. This procoagulant activity is TF dependent.
- Infection of endothelial cells with *Cp in vitro* attenuates eNOS, cGMP, and ROS production, and azithromycin reverses this attenuation.
- Infection of HepG2 human hepatocytes *in vitro* leads to an increased production of IL-6 and fibrinogen. Azithromycin dose dependently reduces *Cp*-induced IL-6, but not fibrinogen, production. We could not confirm that azithromycin has an anti-inflammatory effect.
- Infection of endothelial cells leads to increased expression of adhesion molecules.
   The endothelial response to chronic infections depends on the intracellular iron level.
- Adipocytes can be infected by several microorganisms *invitro*. Infection of adipocytes with CMV, Ad 36, and influenza A leads to increased production of IL-6.
- Infection of co-cultures of human adipocytes and THP-1 macrophages with CMV and influenza A induces the overproduction of IL-6 and PAI-1.



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Er bestaan aanwijzingen dat infecties, met influenzavirus, Cytomegalovirus (CMV) en *Chlamydia pneumoniae* (*Cp*) in het bijzonder, het proces van atherosclerose ('aderverkalking') in gang kunnen zetten en/of versnellen. Dit kan gebeuren via directe infectie van de cellen in de binnenbekleding van de bloedvaten (endotheelcellen) en indirect door infectie van andere cellen en organen, waardoor een gegeneraliseerde ontstekingsreactie ontstaat. Hierdoor kunnen uiteindelijk aandoeningen aan bloedvaten ontstaan zoals een hartinfarct, herseninfarct of perifeer arterieel vaatlijden. In de laboratoriumstudies die deel uitmaken van dit proefschrift wordt onderzocht of bepaalde micro-organismen in staat zijn om verschillende cellen van verschillende organen te infecteren en wordt onderzocht wat het effect is op de functie van die cellen. Vooral wordt gekeken naar processen die een rol spelen bij het ontstaan van atherosclerose.

In **hoofdstuk 2** laten we zien dat luchtwegvirussen (influenza A en B, parainfluenza-1, respiratoir syncytieel virus (RSV), adenovirus en CMV) uit navelstrengen verkregen endotheelcellen kunnen infecteren. Als reactie op infectie ontstond verhoogde stollingsactiviteit op geïnfecteerde endotheelcellen. De factoren van het extrinsieke stollingssysteem, zoals tissue factor (weefselfactor, factor III) en factor VII, zijn eiwitten die op endotheelcellen aanwezig zijn. Tissue factor staat aan het begin van het stollingssysteem en wordt geactiveerd door beschadiging van de bloedvatwand. Uit een stollingsproef bleek dat de gemiddelde stoltijd met 55% verkort werd. Deze versnelde stolling was afhankelijk van stollingsfactor VII. Daarmee werd aangetoond dat de stolling via de tissue factor afhankelijke (extrinsieke) stollingsroute verliep. Naast de verkorte stollingstijden, nam ook de expressie van het weefselfactor eiwit op het oppervlak van geïnfecteerde endotheelcellen met een factor 4 tot 5 toe. De conclusie is dan ook dat respiratoire virussen in staat zijn om menselijke endotheelcellen te infecteren en dat de toegenomen expressie van tissue factor de endotheelcellen in een staat van verhoogde stollingsactiviteit brengt.

Monocyten behoren tot een fractie van witte bloedcellen die een rol spelen bij het opruimen van indringers zoals bacteriën. De productie van ontstekingseiwitten (cytokines) door monocyten kan het atherosclerotisch proces versnellen en bijdragen aan het ontstaan van vaatproblemen als gevolg van vorming van instabiele atherosclerotische plaques in de vaatwand. In **hoofdstuk 3** werd onderzocht of de met virus geïnfecteerde monocyten ook de bloedstolling in gang zetten en

ontstekingsbevorderende cytokines kunnen produceren. Door middel van een geoptimaliseerde isolatiemethode werden monocyten verkregen die geïnfecteerd werden met CMV, *Cp* en influenzavirus. De stollingstijden van CMV- en *Cp*-geinfecteerde monocyten bleken met ca. 60% verkort te zijn. Net als bij endotheelcellen bleek de stolling factor VII- afhankelijk te verlopen via de extrinsieke stollingsroute. De door geïnfecteerde monocyten op gang gebrachte stolling werd veroorzaakt door expressie van tissue factor. Tevens produceerden geïnfecteerde monocyten de ontstekingsbevorderende cytokines interleukine- 6 (IL-6) en interleukine-8 (IL-8). Onze resultaten geven aan dat slechts kleine hoeveelheden infectieus virus in staat zijn om monocyten aan te zetten tot een merkbare immunologische reactie. Geconcludeerd wordt dat infectie van monocyten leidt tot ontstekingsreacties en stollingsbevorderende processen die tot verergering van atherosclerose kunnen leiden.

De lever is een belangrijke producent van acute fase eiwitten, stollingsfactoren, cholesterol en triglyceriden. De aanmaak van acute fase eiwitten is een eerste normale reactie van het lichaam op ontstekingen. Ontsteking als gevolg van een infectie of weefselbeschadiging versterkt de productie van ontstekingsbevorderende cytokines als IL-1, IL-6 en tumor necrosis factor (TNF)-alfa. Deze cytokines stimuleren op hun beurt weer levercellen tot sterk verhoogde aanmaak van acute fase eiwitten zoals C-reactief proteïne (CRP). IL-6 is de belangrijkste cytokine die de lever aan kan zetten tot productie van CRP. De stollingsfactor fibrinogeen wordt aangemaakt door levercellen en is betrokken bij het stabiliseren van bloedstolsels. Onder abnormale omstandigheden, zoals chronische ontstekingsprocessen bij atherosclerose, kunnen plasmaconcentraties van acute fase eiwitten en stollingsfactoren langdurig verhoogd zijn. In hoofdstuk 4 beschrijven wij dat menselijke levercellen in het laboratorium geïnfecteerd kunnen worden door de intracellulaire pathogenen Cp and CMV. Het bleek dat infectie van levercellen met Cp leidt tot een tijd- en dosisafhankelijke productie van IL-6. Ook hebben wij onderzocht of azitromycine in staat zou zijn om Cp-infectie en de ontstekingsreactie in levercellen af te remmen. Bepaalde antibiotica zoals azithromycine en roxithromycine (macroliden) zijn namelijk effectief voor de behandeling van intracellulaire infecties omdat langdurig hoge antibioticumconcentraties in de cellen bereikt kunnen worden. Toevoeging van azitromycine remde de ontstekingsreactie van geïnfecteerde levercellen en onderdrukte de infectie van Cp in vitro. Wij concluderen dat de waargenomen afname van de ontstekingsreacties na azithromycine gebruik het rechtstreeks gevolg is van de antibacteriële werking van het antibioticum en niet een ontstekingsremmend effect van dit middel is.

Chronische infecties kunnen directe vaatschade maar ook systemische ontstekingsreacties tot gevolg hebben. In beide gevallen kan dit leiden tot een verstoorde functie van endotheelcellen (endotheeldisfunctie). Stikstofoxide (NO) is een belangrijke regulator van endotheelfunctie en zorgt o.a. voor vaatverwijding. Onder pathologische omstandigheden wordt minder NO gevormd en juist meer schadelijke reactieve zuurstofradicalen (reactive oxygen species, ROS), hetgeen leidt tot vaatbeschadiging en daardoor endotheeldisfunctie en ontstekingreacties. In hoofdstuk 5 werd, na infectie van endotheelcellen met Cp- en CMV, de productie gemeten van NO, ROS en het enzym endotheliaal nitric oxide synthase (eNOS), wat zorgt voor aanmaak van NO. Omdat NO lastig te meten is werd in plaats daarvan cyclisch guanosine monofosfaat (cGMP) gemeten, als maat voor de beschikbaarheid van NO. Wij veronderstelden dat infecties atherosclerose op gang kunnen brengen door onderdrukking van NOproductie en dat toevoeging van azitromycine dit effect weer zou kunnen herstellen. Cp-geinfecteerde endotheelcellen bleken inderdaad minder eNOS, cGMP en ROS te produceren. Toevoeging van azitromycine herstelde de eNOS, cGMP en ROSproductie. De resultaten van deze studie suggereren dat behandeling met antibiotica de verstoorde endotheelfunctie ten gevolge van infecties mogelijk zou kunnen verbeteren.

Bij endotheeldisfunctie kan de expressie van de adhesiemoleculen vasculair cel adhesiemolecuul-1 (VCAM-1), intercellulair adhesiemolecuul-1 (ICAM-1) en endotheliaal selectine (E-selectin) verhoogd zijn. Infecties kunnen de aanleiding zijn voor de activatie van endotheel en de gelijktijdige aanwezigheid van ijzer kan dit proces versterken. Diverse studies hebben aangetoond dat vorming van atherosclerose afnam door ijzerchelatietherapie (wegvangen van ijzer) of ijzerarme diëten. In **hoofdstuk 6**, laten we zien dat *Cp*- en CMV-infecties de ICAM-1 en VCAM-1 expressie in endotheelcellen op gang brengen en de productie van E-selectin versterken. In *Cp*-geinfecteerde endotheelcellen werden deze effecten nog eens verder versterkt wanneer ijzerrijk medium werd toegevoegd. De aanwezigheid van ijzer beïnvloedt weliswaar de reactie van endotheelcellen op infectie, maar dit effect werd al gezien voordat infectie en vermenigvuldiging van *Cp* kon worden aangetoond. Omdat na infectie de concentratie van zuurstofradicalen in endotheelcellen niet veranderde, is

het ook niet waarschijnlijk dat de expressie van ICAM-1 en VCAM-1 werd aangestuurd door productie van radicalen. Kennelijk verloopt endotheelactivatie door infectie via een andere route. Dit geeft aan dat ijzer endotheelcellen eigenlijk vooral prikkelt door het opwekken van oxidatieve stress. Op deze wijze kunnen endotheelcellen echter wel gevoeliger worden voor de effecten van infecties en bijdragen aan het ontstaan van atherosclerose.

Buikvet zorgt niet alleen voor de opname en opslag van vrije vetzuren, maar is ook in staat tot productie en uitscheiding van adipokines (cytokines geproduceerd door vet). Vetzucht wordt ook wel voorgesteld als een staat van chronische ontsteking waarvan bekend is dat die betrokken is bij het ontstaan van hart- en vaatziekten en diabetes mellitus. Adipocytendisfunctie is daarbij cruciaal. Uit **hoofdstuk 7** blijkt dat ook adipocyten (vetcellen) en pre-adipocyten (voorloper vetcellen) geïnfecteerd kunnen worden met verschillende virussen. Infectie van adipocyten met adenovirus type 36, maar niet met adenovirus type 2, leidt tot verhoogde productie van IL-6 door adipocyten. Dit ondersteunt de hypothese dat micro-organismen mogelijk een rol spelen in het ontstaan en onderhouden van een chronisch laaggradig ontstekingsproces dat gepaard gaat met adipocytendisfunctie. Het feit dat ongeveer een derde deel van de totale IL-6 plasmaconcentratie afkomstig is van vetweefsel geeft aan dat ontstekingreacties in vetweefsel tot belangrijke biologische effecten kunnen leiden.

Micro-organismen die het vetweefsel infecteren kunnen een chronische ontstekingsreactie in gang zetten. De hierbij geproduceerde adipokines kunnen vervolgens de functie van naastgelegen cellen beïnvloeden (paracrien effect) en er voor zorgen dat meer macrofagen het vetweefsel binnendringen. Hierdoor kan de totale productiecapaciteit van ontstekingsbevorderende stoffen nog meer stijgen. Het is ook denkbaar dat toegenomen infiltratie van geïnfecteerde macrofagen in vetweefsel de aanzet geeft tot een ontstekingsreactie. In **hoofdstuk 8** werden menselijke vetcellen (adipocyten) en macrofagen samen gekweekt (co-cultuur) en geïnfecteerd. Wij konden laten zien dat infectie van adipocyten ook daadwerkelijk overgaat naar macrofagen en vice versa (transinfectie). CMV- en influenza A infectie in co-culturen van adipocyten en macrofagen leidt tot overproductie van IL-6 en plasminogeen activator inhibitor-1 (PAI-1). Directe communicatie tussen deze celtypen leidt kennelijk tot de aanzienlijk verhoogde staat van ontsteking die gezien wordt bij obesitas. Dit ondersteunt de

gedachte dat zowel adipocyten als macrofagen in vetweefsel betrokken zijn bij adipokine productie. Hoewel adipocyten de belangrijkste bron van cytokines lijken te zijn, is de mate van infiltratie door macrofagen waarschijnlijk bepalend voor de ernst van de ontstekingsreactie. Uiteindelijk bleek ook dat de stijging van IL-6 in geïnfecteerde co-culturen vooral aan macrofagen toegeschreven kon worden. Deze bevindingen verschaffen inzicht in de manier waarop infectie van vetweefsel op afstand betrokken kan zijn bij het activeren van andere weefsels (endotheel, lever, spieren) en zodoende een pathologische link zou kunnen zijn tussen obesitas en de verhoogde kans op het ontstaan van hart- en vaat aandoeningen.

## **Conclusies van dit proefschrift:**

- Intracellulaire infecties met micro-organismen zoals respiratoire virussen, Chlamydia pneumoniae (Cp) en cytomegalovirus (CMV) leiden tot verhoogde stollings-activiteit van endotheelcellen en monocyten in vitro die geassocieerd is met verhoogde expressie van tissue factor.
- *In vitro* infectie van endotheelcellen met *Cp* onderdrukt eNOS, cGMP en ROS productie. Incubatie met azitromycine voorkomt deze effecten.
- Infectie van levercellen *in vitro* leidt tot een toename van IL-6 en fibrinogeen productie. Toevoeging van azitromycine verlaagt weliswaar de door *Cp* geïnduceerde IL-6 productie, maar niet de fibrinogeen productie. Azithromycine heeft geen directe anti-inflammatoire effecten op levercellen.
- Infectie van endotheelcellen leidt tot toegenomen expressie van adhesiemoleculen op endotheelcellen. De mate waarin endotheelcellen op infecties reageren, hangt ook af van de ijzer concentratie in het kweekmedium van de cellen.
- Vetcellen zijn met verscheidene micro-organismen *in vitro* te infecteren. Infectie van vetcellen met CMV, Ad 36 and influenza A leidt tot verhoogde IL-6 productie.
- CMV- en influenza A infecties induceren toegenomen productie van IL-6 en PAI-1 in co-culturen van vetcellen en monocyten.





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Game, set en match: het boekje ligt er en ik ben trots op het resultaat!

Toen mij paar jaar geleden werd voorgesteld om te gaan promoveren moest ik daar eens rustig over nadenken. Later besefte ik dat ik er gewoon voor moest gaan. Uiteindelijk ben ik toch jarenlang met veel plezier aan het experimenteren geweest en het is mooi dat die inspanningen nu met dit proefschrift bekroond zijn.

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幸運はライデン大学で楽しい時を過し、en 日本で会おう!!

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John

Juni 2009.



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Johannes Jacobus Maria Bouwman was born on April 28th 1959 in Utrecht, The Netherlands.

After graduation from secondary school (college De Klop, Utrecht) in 1976, he started his HBO-A education medical microbiology at the Ir. L.W. Gijsen Instituut in Utrecht. He fulfilled his training period at the CBSL Laboratories Hilversum and successfully completed his study in 1978. His career as a lab technician started at the Streeklaboratorium Deventer. In 1979 the Military Hospital Dr. A. Mathijsen in Utrecht became his next employer. From 1988 until 1990 he was a lab technician at the virology department of the GG&GD Amsterdam. At this department he acquired the knowledge and proficiency for the cell and viral culture techniques that eventually have been the basis for the fundamental research work presented in the studies of this thesis.

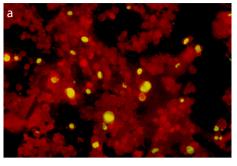
The Department of Medical Microbiology of the Diakonessen Hospital Utrecht has been the employer since December 1990. At first, all obligatory tasks as a lab technician in medical microbiology were performed. Starting from 1992 he was involved in development and implementation of a virology unit and started introducing molecular diagnostics in 1996. Next to these activities, in 1994 he became a laboratory attendant for HLO-trainees and medical students during their research- and graduation stage. In 2001 the research described in this thesis started, alternating part time and full time, under supervision of dr. RJ.A. Diepersloot (Department of Medical Microbiology; Diakonessen Hospital Utrecht) and dr. F.L.J. Visseren (Deptartment of Vascular Medicine; University Medical Center Utrecht). 2008 and 2009 were largely dedicated to research and the completion of this thesis.

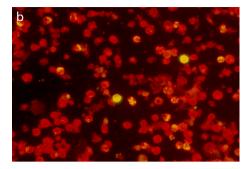
John Bouwman married Annelies Hornstra in 1982. They live in Nieuwegein and have two daughters: Monique (1987) and Rianne (1990).

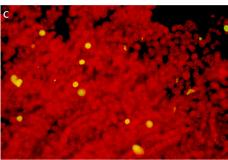


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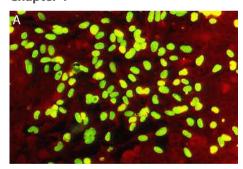
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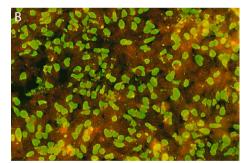




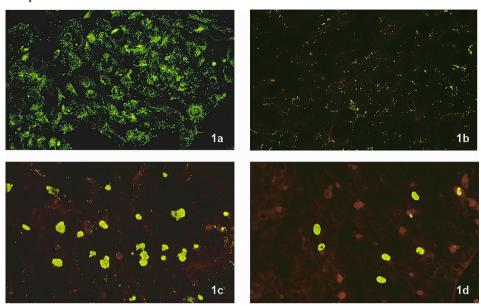


**Figure 1** Fluorescence microscopy images of virus-infected monocytes. Magnification 200x after overnight incubation; Red background: uninfected monocytes. Green areas: virus-infected monocytes stained with FITC-labeled anti-virus antibodies. (a) CMV, specific staining of CMV in the nuclei of the infected cells. (b) Influenza A, smooth staining pattern of influenza A in monocytes. (c) Cp, denser staining pattern of Cp in the cytoplasm of monocytes.

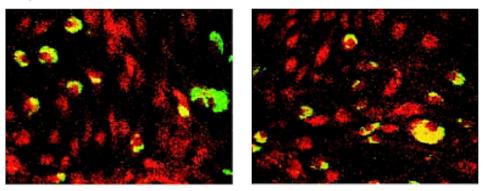




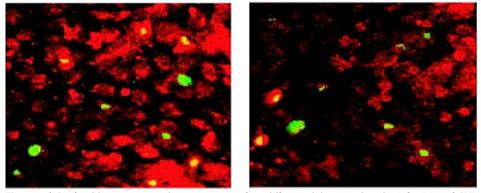
**Figure 1** Infection of HepG2 cells. **A**: Fluorescence staining of CMV-infected HepG2 cells. Note apple-green fluorescence of CMV immediate early antigen in the nuclei of infected cells. **B**: Fluorescence staining of C. pneumoniae–infected HepG2 cells. Apple-green fluorescence (genus-specific) denotes Chlamydia antigens in the cytoplasm of infected cells. Background of each photo contains uninfected cells counterstained with Evans blue (red). Original magnification 100x



**Figure 1**. Fluorescence microscopy images ( $\times$ 200) of human umbilical vein endothelial cells (HUVECs) 48 h after inoculation. Indirect immunostaining with antihuman von Willebrand factor (a), anti-Cp (b,c) and anti-CMV (d) antibodies. Secondary antibody: FITC-labelled rabbit antimouse conjugate; counterstaining: Evans Blue. Red background: uninfected cells. Apple green fluorescence: positive identification of von Willebrand factor (a), inactivated Cp (b), Cp (MOI = 1) (c) and CMV (MOI = 1) (d).



**Figure 4d** Confocal laser micrographs, representing four different slides, visualize the infectivity of Cp (green) on HUVECs (red) in the absence and presence of iron.



**Figure 5d** Confocal laser micrographs, representing four different slides, visualize the infectivity of CMV (green) on HUVECs (red) in the absence and presence of iron.

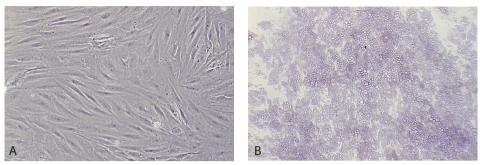
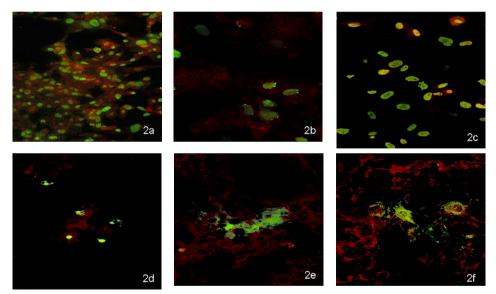
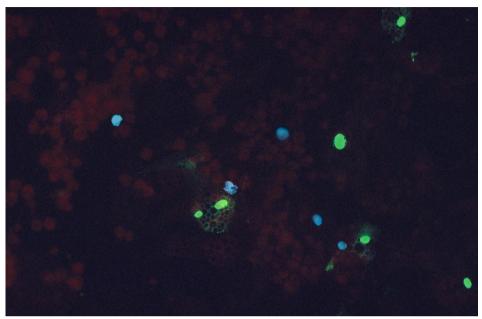


Figure 1 (Pre-) Adipocytes in culture (200x)

- A Pre-adipocytes: spindle-shaped "fibroblast-like" cells grown to confluence.
- B Adipocytes: differentiated cells 13 days after start of differentiation. Cells are rounded and have accumulated lipid droplets in their cytoplasm



**Figure 2. Fluorescent images of infected adipocytes** (200x, 90% differentiated) after 48 hours infection with: Ad2, Ad 36, Cp, CMV, Inf, and RS-virus. Background (red): uninfected cells. Infected cells: apple green fluorescence either condensed in the nuclei (CMV;2c) or diffuse in the cytoplasm (Ad 2, Ad 36, Inf, RSV; 2a,b,d,e,f respectively). Cp-infected cells: compact fluorescent vacuoles in the cytoplasm (2d)



**Figure 1** Dual color immunofluorescence image of transinfection of influenza A in a co-culture of adipocytes and THP-1 cells. Influenza infected THP-1 cells were co-cultured with uninfected adipocytes for 48 hours. THP-1 cells infected with influenza A demonstrate blue fluorescence (AMCA label). After 48 hours, influenza infection was transmitted from THP-1 cells to adipocytes, as indicated by apple green fluorescence (FITC label). Background: uninfected adipocytes and THP-1 cells (round) counterstained with Evans Blue (red). Lipid droplets in adipocytes are clearly visible as dark inclusions in the cytoplasm of adipocytes.



