

**Low-Density Lipoprotein Receptor-Related Protein  
Novel Aspects Revealed**

**Patricia Spijkers**

Cover illustration: Sabine Spijkers  
Photograph taken at lab day 2004 at Austerlitz

Printed by Febodruk B.V., Enschede

ISBN: 90-393-4001-3

# **Low-Density Lipoprotein Receptor-Related Protein Novel Aspects Revealed**

Low-density Lipoprotein Receptor-Related Protein – nieuwe aspecten onthuld  
(met een samenvatting in het Nederlands)

## **Proefschrift**

ter verkrijging van de graad van doctor aan de  
Universiteit van Utrecht op gezag van de Rector  
Magnificus, Prof. Dr. W.H. Gispen, in gevolge het  
besluit van het College voor Promoties in het  
openbaar te verdedigen op dinsdag 20  
september 2005 des middags te 14.30 uur

door

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Geboren op 6 september 1978 te 's-Hertogenbosch

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The study described in this thesis was supported by a grant of the Dutch Organization of Scientific Research (NWO) (no. 902-26-236).

Financial support by NWO and the Stichting voor Afweerstoornissen Nederland for the publication of this thesis is gratefully acknowledged.

Additional financial support by the University of Utrecht, Dr. Ir. van der Laar Stichting, Pfizer B.V., Internationale Stichting voor Alzheimer Onderzoek, and Tebu-Bio is also gratefully acknowledged.

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## Abbreviations

$\alpha_2$ M	$\alpha_2$ -macroglobulin	NIF	neutrophil inhibitory factor
A $\beta$	amyloid $\beta$ -peptide	NK	natural killer
AD	Alzheimer's Disease	NMDA	N-methyl-D-aspartate
APC	antigen presenting cell	PBMC	peripheral blood mononuclear cell
ApoE	Apolipoprotein E	PDGF	platelet-derived growth factor
ApoER2	ApoE receptor 2	PFA	paraformaldehyde
APP	A $\beta$ precursor protein	PHA	phytohaemagglutinin
BBB	blood-brain barrier	PKA	protein kinase A
BEC	brain endothelial cells	PMA	phorbol-12-myristate-13 acetate
C4BP	C4b-binding protein	PMN	polymorphonuclear cell
CCP	complement control protein	PNP	p-nitrophenyl phosphate
CD	cluster of differentiation	RAGE	receptor for advanced glycation end products
DIA $\beta$ 40	A $\beta$ 40 peptide with Dutch/Iowa mutation	RAP	receptor-associated protein
EGF	epidermal growth factor	rC4BP	recombinant C4BP
FITC	fluorescein isothiocyanate	RU	resonance units
GST	glutathione-S-transferase	SPR	surface plasmon resonance
GVHD	graft-versus-host disease	TCA	trichloroacetic acid
HSPG	heparan sulfate proteoglycan	TFPI	tissue-factor pathway inhibitor
HUVEC	human umbilical vein endothelial cells	TNF- $\alpha$	tumor necrosis factor- $\alpha$
ICAM	intercellular adhesion molecule	uPA	urokinase-type plasminogen activator
IL	interleukin	uPAR	uPA receptor
LDL	low-density lipoprotein	VLDL	very low-density lipoprotein
LRP	LDL receptor-related protein	VWF	Von Willebrand Factor
MLR	mixed lymphocyte reaction		
NGF	nerve growth factor		

Een reis van duizend mijl begint  
met de eerste stap  
(*Lao-Tse*)



**chapter**  
**1**

**General Introduction**

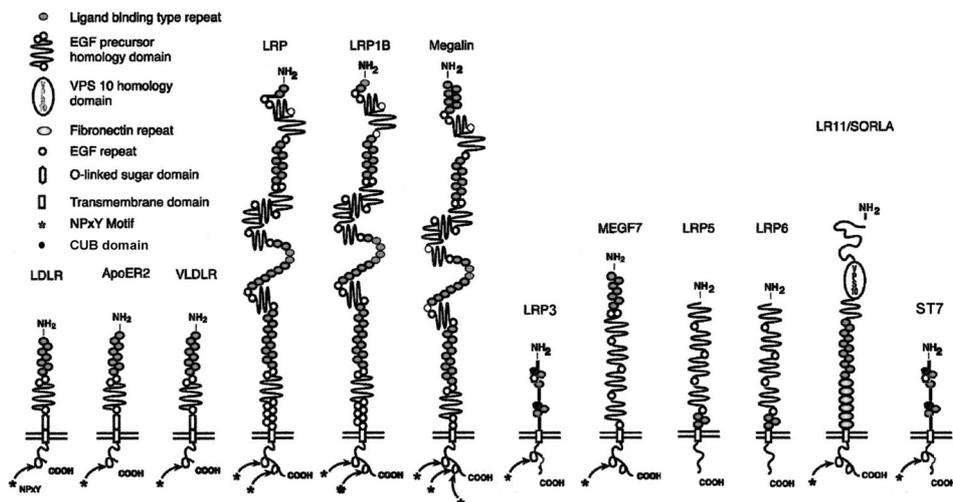
## Multiligand Receptors

Cell surface receptors play a crucial role in maintaining cellular homeostasis by linking the extracellular environment and intracellular compartments. In addition, they can act as an anchor in the adhesion process to other cells or the extracellular matrix. Whereas receptors have long been thought to recognize one specific ligand, a broad range of receptors has now been identified that are able to bind multiple ligands. Moreover, these ligands may even belong to structurally and functionally unrelated families. As a result of this complexity, the receptor must therefore be accompanied by a tightly organized regulatory system to ensure that the appropriate cellular responses are initiated. For example, multiligand receptors may be specialized in distinct functions, like the removal of obsolete proteins, or may be differentially expressed by particular cell types. Also, the multiligand receptor could be part of a heterologous protein complex that controls a specific physiological process<sup>1;2</sup>.

## Low-Density Lipoprotein (LDL)-Receptor Family

The LDL-receptor family is a classical group of multiligand cell-surface receptors that share structural and functional homology (*Fig 1*). The first member that has been characterized was the LDL receptor, which plays a key role in cholesterol homeostasis<sup>3</sup>. At present, 12 members are described in humans: LDL receptor, Apolipoprotein E receptor 2 (ApoER2), very low density lipoprotein (VLDL) receptor, LDL receptor-related protein (LRP), LRP1B, megalin, LRP3, LRP4/protein 7 with multiple epidermal growth factor (EGF)-like repeats (MEGF7), LRP5/7, LRP6, LDL receptor relative (LR) 11/ sorting protein-related receptor containing LDL receptor class A repeats (SorLA) and ST7/LRP12 (reviewed in Li et al<sup>4</sup> and Herz<sup>5;6</sup>). In general, members contain the following structural domains (*Fig 1*). Their extracellular domain contains ligand-binding regions that consist of clusters of cysteine-rich complement-type repeats. These repeats are separated by EGF precursor repeats and YWTD modules. The latter two domains are thought to be involved in pH-dependent ligand dissociation from the receptors in the endosomes after internalization<sup>7;8</sup>. The receptors are anchored in the cell membrane via a single-chain transmembrane domain, which is followed by a relatively short intracellular tail that harbors motifs for endocytosis and signal transduction.

The statement that all members of the LDL receptor family are multiligand receptors, does not imply that they recognize identical ligands and provoke the same cellular responses. In contrast, ligand binding and cellular reaction may be highly regulated, *e.g.* by differential cellular expression patterns or variable splicing. An overview of tissue-specific expression, ligands and function of the members is summarized in table 1.



**Figure 1. Schematic representation of the human LDL receptor family.** Currently, 12 members are known with structurally homologous domains (depicted in the upper left box). Figure adapted and modified from Herz et al<sup>6</sup>.

## Structure of LRP

In the late 80's, three groups independently described a new protein that displays homology to the LDL receptor. Herz et al<sup>9</sup> searched for cDNA clones containing the highly conserved cysteine-rich complement-type repeats that are present in LDL receptor. This resulted in the discovery of a ~15 kb cDNA stretch, encoding for a 4525 amino acid protein, designated LRP. Simultaneously, Gliemann et al<sup>10;11</sup> and Ashcom et al<sup>12</sup> purified a high affinity receptor for activated  $\alpha_2$ -macroglobulin ( $\alpha_2$ M) from human hepatocytes, which was therefore called  $\alpha_2$ M receptor. Protein sequence analysis revealed that the  $\alpha_2$ M receptor and LRP were the same receptor<sup>13</sup>.

Table 1. Characteristics of the human LDL receptor family.

Receptor	Examples of expression	Examples of ligands	Examples of functions	References
LDL receptor	liver	ApoE, ApoB	lipid metabolism in liver	3;14
ApoER2/LRP8	brain, platelets	reelin, ApoE, $\beta$ 2-GPI*	brain development, platelet activation	15
VLDLR	heart, muscle, adipose tissue	lipoproteins, reelin	lipid metabolism in extrahepatic tissues, brain development	16
LRP	see table 2	see table 3	see text	11;9
LRP1B/LRP-DIT*	brain, thyroid, salivary glands	uPA* / PAI-1*, APP*	cell migration	17;18
Megalin	kidney	vitamin D binding protein, ApoE, uPA / PAI-1	protein absorption from the urine, vitamin D catabolism	19
LRP3	skeletal muscle, ovary	?	?	20
LRP4/MEGF7	brain	?	VLDL receptor-like signaling?	21
LRP5/7	brain, aorta, bone, liver	Wnt	Wnt signaling, osteogenesis	22;23
LRP6	brain, embryonic tissues	Wnt	Wnt signaling	22;24
LR11/SorLA	brain, liver	ApoE, head activator peptide	protein catabolism, cell proliferation	25
ST7/LRP12	heart, skeletal muscle	?	transformation?	26;27

\* APP, amyloid precursor protein;  $\beta$ 2-GPI,  $\beta$ 2-glycoprotein-1; DIT, deleted in tumors; PAI-1, plasminogen-activator inhibitor-1; uPA, urokinase-type plasminogen activator.

Whereas LDL receptor has a molecular weight of 130 kDa, LRP is a considerably larger protein. It is processed as a 600-kDa single-chain protein that is cleaved by furin in the *trans* Golgi apparatus<sup>28;29</sup>. The two chains, a 515-kDa heavy or  $\alpha$ -subunit, and an 85-kDa light or  $\beta$ -subunit, are expressed at the cell surface as a non-covalently linked heterodimer. Both  $\alpha$ - and  $\beta$ -chains are heavily N-linked glycosylated<sup>30</sup>.

The 515-kDa extracellular domain harbors four clusters of 2, 8, 10 and 11 complement-type repeats, designated as cluster I to IV. Cluster II and IV have been shown to dominate ligand binding<sup>31;32</sup>. The LRP 85-kDa cytoplasmic tail contains two copies of NPXY motifs, one YXXL motif and two di-leucine motifs. These motifs are generally known as internalization signals, which are also present in the LDL receptor<sup>33</sup>. In addition, they can serve as docking sites for cytosolic adaptor proteins involved in cellular signaling<sup>34</sup>.

Besides its localization at the cell surface, a low amount of LRP is also present in plasma<sup>35;36</sup>. This soluble form consists of the full  $\alpha$ -chain and the extracellular part of the  $\beta$ -chain, as a result of metalloprotease activity at the cell surface. Furthermore, the cytoplasmic tail also contains a cleavage site, which is sensitive to  $\gamma$ -secretase activity. Cleavage may allow transportation of the disconnected tail to other intracellular compartments and subsequent participation in intracellular processes<sup>37;38</sup>.

## Tissue Distribution of LRP and Regulation of Expression

LRP protein is expressed by a broad number of cells<sup>39</sup>, which are summarized in table 2.

One of the aspects of LRP biology that has been poorly explored so far, relates to the molecular regulation of its expression. In view of its multifunctionality, it seems reasonable to assume that LRP requires strict regulation of its cellular expression. Unlike the LDL-receptor gene, the promoter region of the LRP gene does not contain a sterol-regulatory region<sup>40</sup>. In placenta, cyclic AMP (cAMP) prevents LRP expression during differentiation of cytotrophoblasts into syncytiotrophoblasts<sup>41</sup>. In reverse, maturation of monocytes to macrophages results in an increase of LRP protein expression<sup>42</sup>. In retinal pigment epithelial cells, LRP mRNA and protein expression is increased in response to transforming growth factor (TGF)- $\beta$ 1, TGF- $\beta$ 2 or vascular endothelial growth factor<sup>43</sup>. An increase of cell surface expression may also be a result of mobilization of LRP from an endosomal storage pool to the cell surface, as is the case for insulin-treated adipocytes<sup>44</sup>. Also nerve growth factor (NGF) stimulates rapid translocation of LRP to

the cell surface in neurons<sup>45</sup>. This differentially regulated expression may be a key in specifying LRP's function.

**Table 2. Tissue and cell distribution of LRP protein<sup>39</sup>.**

<i>Gastro-Intestinal Tract</i> hepatic parenchymal cells intestinal epithelial cells kupffer cells	<i>Blood and other Bone Marrow-Derived Cells</i> monocytes macrophages polymorphonuclear cells langerhans cells in skin dendritic cells erythroblasts
<i>Reproductive System</i> syncytiotrophoblasts granulosa cells in ovary leydig cells in testis sertoli cells in testis stromal cells in ovary decidua cells in placenta hofbauer cells in placenta mammary epithelial cells	<i>Central Nervous System</i> astrocytes neurons retinal muller cells retinal pigment epithelial cells trabecular meshwork cells in eye cerebral microvessel endothelial cells purkinje cells
<i>Glandular System</i> medullary cells in adrenal gland epithelial cells of parathyroid gland	<i>Other</i> adipocytes bone marrow reticulum cells chondrocytes pericytes in blood vessels myofibroblasts smooth muscle Cells fibroblasts mesothelial cells keratinocytes
<i>Urinary Tract</i> dendritic interstitial cells of kidney glomerular mesangial cells in kidney	
<i>Respiratory Tract</i> lung macrophages laryngeal epithelium bronchial and alveolar epithelium	

## LRP Ligands

Following activated  $\alpha_2M$ , ApoE-enriched  $\beta$ -VLDL was the second protein that was identified as a ligand for LRP<sup>46;47</sup>. Rapidly, more and more proteins were found to bind to LRP and this number is still increasing. These ligands are structurally and functionally unrelated, although part of them can be categorized based on functional characteristics (Table 3)<sup>48;49</sup>. The ligands mainly bind to the clusters II and IV and the isolated recombinant clusters are sufficient to recognize ligands<sup>50</sup>. Further molecular dissection of these clusters showed that a certain ligand binds to only a few repeats and that ligands show different or overlapping binding sites<sup>32;51</sup>. Binding competition between ligands could occur via direct competition or sterical hindrance of adjacent bound ligands.

**Table 3. LRP ligands<sup>48;49</sup>.**

<i>Lipoproteins and Lipases</i>	<i>Proteases and Inhibitor Complexes</i>
ApoE	activated $\alpha_2$ -macroglobulin
lipoprotein lipase	pregnancy zone protein
hepatic lipase	$\alpha_1$ -antitrypsin
chylomicron remnants	elastase. $\alpha_1$ -antitrypsin
sphingolipid activator protein	protein C inhibitor
<i>Haemostatic Components</i>	<i>Complement Factors</i>
factor VIII/ factor VIIIa	C3
factor IXa	C4BP*
TFPI* (/factor VIIa/TF*(/factor Xa))	
tPA*	<i>Growth Factors</i>
uPA*	platelet-derived growth factor
	midkine
<i>Combination of Protease...</i>	connective tissue growth factor
thrombin	
factor IXa	<i>Toxic Compounds</i>
factor XIa	pseudomonas exotoxin A
tPA	rhinovirus
uPA	HIV* tat-protein
<i>and Inhibitors</i>	circumsporozoite protein
protease nexin-1	ricin A
protease nexin-2	saporin
heparin cofactor II	trichosanthin
PAI-1*	gentamycin
antithrombin III	polymycin B
C1-inhibitor	
activated $\alpha_2$ -macroglobulin	<i>Chaperones and Intracellular Proteins</i>
$\alpha_1$ -antitrypsin	HSP96*
	RAP*
<i>Matrix Proteins</i>	Calreticulin
thrombospondin-1	
thrombospondin-2	<i>Other</i>
matrix metalloprotease-2	lactoferrin
matrix metalloprotease-9	APP* / A $\beta$ * peptides
matrix metalloprotease-13	melanotransferrin
fibronectin	

\* A $\beta$ , amyloid  $\beta$  peptide; APP, amyloid precursor protein; C4BP, C4b-binding protein; HIV, human immunodeficiency virus; HSP, heat shock protein; PAI-1, plasminogen-activator inhibitor-1; RAP, receptor-associated protein; TF, tissue factor; TFPI, tissue factor pathway inhibitor; tPA, tissue-type plasminogen activator; uPA, urokinase-type plasminogen activator.

Whereas some ligands bind to LRP via hydrophobic interactions, ligand-receptor interaction occurs predominantly via electrostatic interactions. LRP displays negative charges on its surface, whereas ligands are mainly positively charged<sup>1</sup>. All ligand-LRP interactions are calcium-dependent<sup>1</sup>. As in LDL receptor, each EGF precursor and complement-type repeat contain a single calcium ion<sup>52;53</sup>. Incorporation of calcium ions is important in the maintenance of the conformation of LRP and, thereby, for ligand binding<sup>54;55</sup>.

A ligand of special interest is receptor-associated protein (RAP). During purification of LRP by ligand-affinity chromatography, a  $\sim$  39 kDa protein was co-

purified<sup>12</sup>. This protein was called receptor-associated protein and binds to LRP with high affinity. It functions as an intracellular chaperone by facilitating protein folding in the endoplasmic reticulum and preventing premature ligand binding during the transport to the cell membrane. RAP is the only protein known to bind to cluster III as well<sup>56</sup>. Later, RAP was also found to show high affinity for the other LDL receptor family members, except LDL receptor itself, which has much lower affinity for RAP. Because of RAP's high affinity for the receptors, it is commonly used as a receptor antagonist.

Other ligands relevant to the scope of this thesis will be discussed later in this chapter.

## **The Physiological Role of LRP**

One intriguing aspect of LRP concerns its role in embryonic development. Whereas deficiency of LDL receptor homologues in general results in vital offspring<sup>57-59</sup>, deficiency of LRP is associated with early embryonic death at day 10 of gestation<sup>60</sup>. Apparently, LRP's most homologous relatives, LRP1B and megalin, are unable to compensate for the loss of LRP. This may be caused by a highly restricted cellular expression pattern of LRP1B and megalin, or by an LRP-specific function. This critical role of LRP during embryogenesis may point to the involvement of this receptor in various processes. To facilitate studying LRP function *in vivo*, however, conditional LRP knock-out mice have been constructed using the Cre/Lox system<sup>61</sup>.

### **LRP as an endocytic receptor**

One of the first functions that could be assigned to LRP was its role in lipoprotein metabolism. Kowal et al<sup>46</sup> found that LRP mediates the uptake and lysosomal delivery of ApoE-enriched  $\beta$ -VLDL in hepatocytes. LRP's endocytic capacity was not limited to this ligand, but is applied to many other proteins as well. Therefore, LRP is considered mainly as a multiligand clearance receptor. LRP continuously shuttles between the cell surface and endosomes, independent of the presence of ligands<sup>62-64</sup>. Li et al, however, suggest that endocytosis is facilitated by protein kinase A (PKA)-mediated phosphorylation of the intracellular tail of LRP upon ligand binding<sup>65</sup>. Based on characteristics of other endocytic receptors, like the LDL receptor, the two NPXY motifs in the cytoplasmic tail of LRP were thought to be the internalization signals<sup>33</sup>. However, more recent studies on LRP showed that the dominant endocytosis signals are a YXXL and a di-leucine motif that are located in the intracellular tail<sup>63</sup>. Receptor-mediated endocytosis starts

at clathrin-coated pits in the cell membrane. The growing vesicle pinches off from the cell membrane and is transported to endosomes. Because of an acidic environment in endosomes, ligands dissociate from LRP and are transported to lysosomes for degradation. In contrast, LRP efficiently recycles back to the plasma membrane for reuse<sup>4;66;67</sup>. During each cycle, a substantial part of the total LRP pool is targeted to the proteasome for degradation<sup>68</sup>.

A considerable number of ligands use other membrane-anchored proteins than LRP as primary docking sites for cellular binding. A major group comprises heparan sulfate proteoglycans (HSPGs), which are abundantly expressed at the cell surface. They consist of a core protein, termed proteoglycan, and heterogeneous heparan sulfate side-chains. Two proteoglycan families are known that can form HSPGs, called syndecans and glypicans. They are inserted in the plasma membrane via a transmembrane domain or a glycosyl phosphatidyl inositol moiety, respectively. Via the heparan sulfate side chains, HSPGs bind to a wide variety of positively charged proteins with relatively high affinity ( $K_d = 1-100$  nM). Because of their abundance, HSPGs function as important ligand capturing proteins, thereby increasing local protein concentration and directing it to its specific receptor (reviewed in ref. 69). One of such a specific receptor is LRP. In the binding and catabolism of many heparin-binding ligands LRP efficiently cooperates with HSPGs. Examples of heparin-binding LRP ligands are: ApoE<sup>70;71</sup>, factor VIII<sup>72</sup>, factor IXa<sup>73</sup> and thrombospondin-1<sup>74</sup>.

### **LRP and phagocytosis**

Macrophages are specialized cell types that have the ability to internalize large particles by a process known as phagocytosis. Binding of a particle triggers a signaling cascade leading to its engulfment and cytoskeletal reorganization. A variety of receptors have been demonstrated to mediate phagocytosis. LRP has been implicated in phagocytosis of apoptotic cells by macrophages<sup>75;76</sup>. Apoptotic cells can be recognized by calreticulin. Since calreticulin also interacts with LRP, it can serve as a physical connection between LRP and apoptotic cells during phagocytosis<sup>77</sup>. Studies using chimeras containing the intracellular tail of LRP demonstrated that this domain of LRP plays a dominant role in the phagocytic process, since mutations in the di-leucine motifs or in the NPXYXXL motif abolish phagocytosis of sheep red blood cells<sup>75</sup>. This implies that an adaptor protein is required that interacts with the cytoplasmic tail of LRP. The group of Ravichandran indeed found that the intracellular adaptor protein GULP binds to LRP in the phagocytic process<sup>76</sup>.

### **LRP and transcytosis**

Transcytosis is an intracellular pathway that mediates transport of proteins from the apical cell membrane to the basolateral cell membrane or vice versa. One specialized transcytosis pathway is transendothelial transport across the blood-brain barrier (BBB). This pathway allows tightly regulated transfer of proteins from blood to the brain tissue via brain endothelial cells to prevent entry of toxic compounds into the brain. Therefore, it has been suggested that transcytosis across the BBB predominantly involves specific receptors. Recently, Demeule et al found that LRP is most probably the candidate receptor for mediating transcytosis of melanotransferrin from the blood across the BBB<sup>78</sup>. Later, also RAP is found to cross the BBB via LRP<sup>79</sup>. However, in contrast to melanotransferrin, RAP is predominantly transported by LRP from the basolateral to the apical membrane, whereas megalin is responsible for transport the other way around. All studies on LRP-mediated transcytosis were unable to provide evidence on receptor-mediated intracellular trafficking of the ligands. While little lysosomal degradation occurs, it is not known whether the ligands remain associated with LRP during transcytosis, or whether ligands are transferred to another receptor that transports them to the other site of the cell surface.

### **LRP and nuclear targeting**

Growth factors employ various mechanisms to exert their effect on cellular proliferation, including their translocation to the nucleus after receptor-mediated internalization. A number of growth factors require their nuclear localization to carry out their mitogenic activity<sup>80-83</sup>. Midkine is a growth factor that is involved in various biological functions, such as neuronal survival, carcinogenesis and tissue remodeling<sup>84</sup>. LRP is the receptor for midkine that mediates its internalization<sup>85</sup>. However, LRP delivers midkine not only to the endosomal/lysosomal degradation pathway, but also to the nucleus. In this latter pathway, nucleolin activity is required to take over the trafficking of midkine to ensure nuclear localization of midkine, which is required for full activity of midkine<sup>86</sup>. Midkine is the only growth factor described to be dependent on LRP for its nuclear localization. However, since platelet-derived growth factor (PDGF) is also transported to the nucleus<sup>87</sup> and is a ligand for LRP as well<sup>88</sup>, it is tempting to speculate that LRP is also able to mediate nuclear targeting of PDGF.

### **LRP as a signaling receptor**

In recent years, it has become apparent that LRP not only functions in intracellular transport pathways, but also contributes to cellular signaling processes. Whereas endocytosis requires LRP's localization in clathrin-coated pits, LRP was also found to be present in caveolae<sup>89</sup>, a cell membrane domain well known to

be involved in cellular signaling. In contrast to the LDL receptor, the cytoplasmic NPXY motif of LRP plays a minor role in endocytosis. However, this motif has an important function in docking of a wide variety of intracellular adaptor proteins, including Disabled (Dab)-1, Dab-2, Fe65, Shc, c-Jun kinase-interacting protein (JIP), PSD-95, CAPON, SEMCAP-1 and ced-6/GULP via their phosphotyrosine binding or PSD-95/Desclarge/ZO-1 domains<sup>76;90;91</sup>. For interaction with these adaptor proteins, LRP's tail needs to be tyrosine phosphorylated at the NPXY motifs by members of the Src kinase family upon activation by diverse stimuli<sup>92</sup>. Most of these adaptor proteins function in cell adhesion, vesicle trafficking, neurotransmission, migration and proliferation<sup>90</sup>. Besides phosphorylation of tyrosine residues, serine and threonine residues in LRP's tail may also be subjected to phosphorylation<sup>65;93</sup>. The responsible kinase is PKA, which in its turn is activated by cAMP<sup>65</sup>. A guanosine triphosphate-binding protein is able to associate with the cytoplasmic tail of LRP after binding of lactoferrin or urokinase-type plasminogen activator (uPA)<sup>94</sup>, causing the observed elevation of cAMP levels. Phosphorylation by PKA results in an increase in the endocytosis rate<sup>65</sup>. Opposing results were obtained by Ranganathan et al<sup>93</sup>. In this study, serine/threonine phosphorylation by protein kinase C causes a decrease in endocytosis velocity. In conclusion, serine/threonine phosphorylation may be involved in regulation of endocytosis rate<sup>65;93</sup>, whereas tyrosine phosphorylation may play a role in other signal transduction pathways.

#### **LRP as a component of heterologous cell surface complexes**

In the previous paragraphs, LRP has been discussed in relation to its function as a single, non-complexed receptor. However, several studies reported that LRP may also participate in heterologous receptor complexes. Within these complexes, LRP and its co-receptor may affect their functions interdependently.

One example of a heterologous complex is the LRP / PDGF-receptor complex. PDGF is able to bind to both the PDGF receptor and LRP, resulting in PDGF receptor-induced phosphorylation of the intracellular NPXY motif of LRP<sup>88</sup>. This reaction results in association of the adaptor protein Shc to the intracellular tail of LRP. Phosphorylation does not occur when phosphoinositide 3-kinase, a protein downstream of PDGF receptor, was blocked by wortmannin, indicating that the PDGF receptor directs phosphorylation of LRP<sup>89</sup>. The authors also demonstrated that LRP phosphorylation occurs in the caveolae of the cell membrane. PDGF enhances smooth muscle cell migration and proliferation. LRP knock-down experiments showed that in the absence of LRP, PDGF-mediated smooth muscle cell migration is increased<sup>95</sup>. Thus, it is conceivable that LRP prevents overstimulation by PDGF, thereby decreasing migration and proliferation of smooth muscle cells<sup>96</sup>. In this way, LRP can prevent the formation of atherosclerotic plaques<sup>97</sup>.

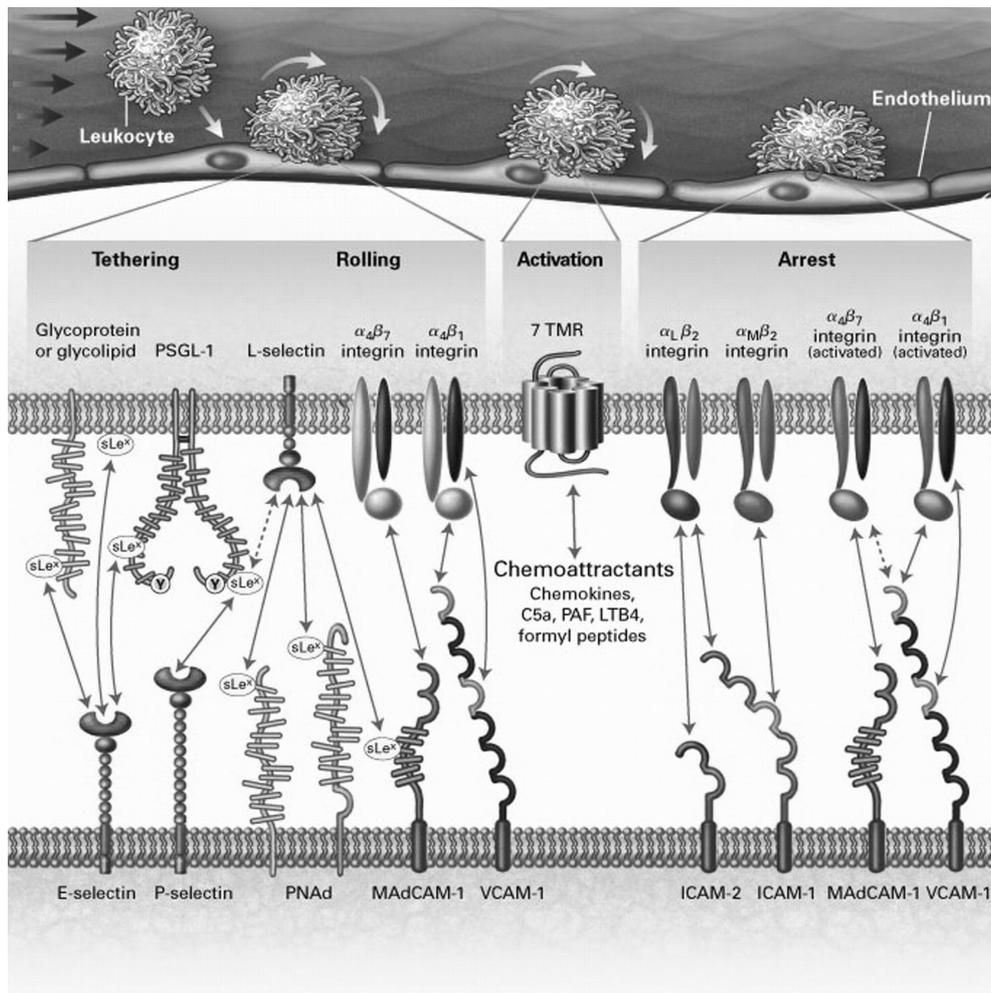
The PDGF receptor is not the only receptor that may cooperate with LRP. Qiu et al<sup>98</sup> described that activated  $\alpha_2$ M influences intracellular calcium signaling in neurons. Pretreatment with activated  $\alpha_2$ M significantly decreases calcium signals in response to N-methyl-D-aspartate (NMDA) through an LRP-mediated pathway. The authors hypothesize that  $\alpha_2$ M ligation to LRP causes downregulation of the NMDA receptor, resulting in an attenuation of extracellular calcium influx through the NMDA receptor. In addition, Bacskai et al found that binding of activated  $\alpha_2$ M to LRP in neuronal cells immediately increases calcium influx from extracellular pools through NMDA receptors<sup>99</sup>. This requires dimerization of LRP, and it is also suggested that both receptors are interlinked via the scaffold protein PSD-95<sup>90;100</sup>. Apparently, LRP has a dual function in modulating NMDA-induced calcium influx. It is responsible for the acute elevated calcium levels and for long-term downregulation of the NMDA receptor, thereby causing desensitization.

## **$\beta$ 2-Integrins**

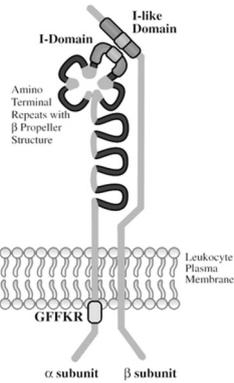
Leukocyte recruitment from the blood is a key event in the regulation of immune responses. Activation of leukocytes by chemokines results in rolling of the cells over the vessel wall, mediated by selectins. Finally, leukocytes arrest, spread and emigrate through to endothelial layer to the subendothelial tissues in a  $\beta$ 2-integrin-dependent manner<sup>101;102</sup> (Fig 2).

$\beta$ 2-Integrins are cell surface receptors that are specifically expressed by leukocytes<sup>103</sup>. They are believed to be involved in processes, such as inflammation, wound healing, and cell growth and survival.  $\beta$ 2-integrins are heterodimeric proteins, consisting of a  $\beta$ 2-subunit (CD18) and one of the following  $\alpha$ -subunits (CD11):  $\alpha_L$  (or CD11a or LFA-1),  $\alpha_M$  (or CD11b or MAC-1),  $\alpha_X$  (or CD11c or p150,95) and  $\alpha_D$  (CD11d) (Fig 3 and Table 4)<sup>104;105</sup>.

For firm adhesion, a tight interaction between  $\beta$ 2-integrins and molecules at the cell surface of endothelial cells is required. To do so,  $\beta$ 2-integrins are able to interact with a wide subset of endothelial cell (-associated) and soluble proteins. The main counter receptors on the endothelial-cell surface are intercellular adhesion molecules (ICAM)-1 and -2<sup>106-108</sup>. Other ligands for  $\beta$ 2-integrins include fibrinogen<sup>109-111</sup>, factor X<sup>112</sup>, heparin<sup>113</sup>, neutrophil inhibitory factor (NIF)<sup>114</sup> and iC3b<sup>115</sup>. Ligand recognition occurs mainly via the I (*inserted*) domain located in the  $\alpha$ -subunit, while  $\beta$ 2-subunits are indispensable for proper integrin function. They are involved in correct subcellular localization, activation of signaling pathways, and direct binding and regulation of the affinity for ligands<sup>116;117</sup>. Whereas  $\alpha_L\beta_2$  is constitutively expressed at the cell surface,  $\alpha_M\beta_2$  cell surface expression



**Figure 2. Mechanism of leukocyte adhesion after activation.** Leukocytes tether to endothelial cells and start rolling. Tethering is mediated by leukocyte cell surface receptors L-selectin, P-selectin, P-selectin glycoprotein ligand 1 (PSGL-1) and  $\alpha_4$ -integrins. The selectins counter receptors at the endothelial cell surface are sialyl-Lewis<sup>x</sup> (sLe<sup>x</sup>) bearing molecules. Rolling leukocytes become activated by chemokines secreted by endothelial cells via receptors with 7 transmembrane domains (7 TMR). The consequential rapid  $\beta_2$ -integrin activation results in firm adhesion and subsequent transmigration to the subendothelial tissues. Figure adapted from Von Andrian et al, 2000<sup>102</sup>.



**Figure 3. Structure of the  $\beta 2$ -integrin heterodimer.** Figure adapted from Harris et al, 2000<sup>103</sup>.

is increased upon stimulation via a mechanism that involves translocation of  $\alpha_M\beta 2$  from intracellular storage vesicles to the cell membrane<sup>118</sup>.

Normally, leukocytes circulate in blood in a non-adhesive state. However, they must be rapidly transformed into active cells, which are able to bind to their counter receptors via  $\beta 2$ -integrins. The precise mechanisms on how integrins are rapidly activated are still not completely understood<sup>119</sup>. In general, it is thought that activation occurs via an inside-out mechanism in which endothelial chemokines are involved. This results in a conformational change in the extracellular domain of the  $\alpha$ -subunit<sup>120</sup>. In addition, activation is also associated with lateral movements of  $\beta 2$ -integrins to particular regions of the cell membrane (=clustering), thereby increasing their avidity<sup>121</sup>. This process requires reorganization of the cytoskeleton<sup>122</sup>, since  $\beta 2$ -integrins are tightly associated with it. Thus, both inside-out signaling and clustering are required for optimal ligand binding<sup>123</sup>.

**Table 4. Main characteristics of  $\beta 2$ -integrins.** Overview of distribution and ligands of the  $\beta 2$ -integrin family members<sup>104;105</sup>.

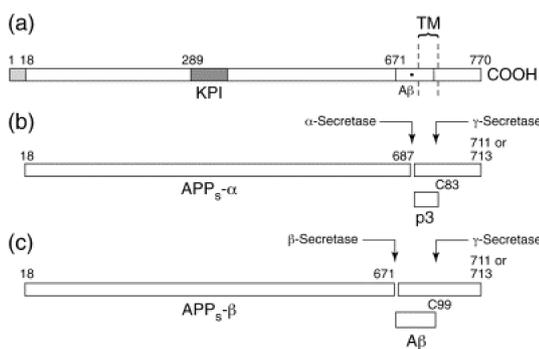
Integrin	Other names	Distribution	Ligands
$\alpha_L\beta 2$	CD11a/CD18, LFA-1	B- and T-lymphocytes, monocytes, neutrophils	ICAM-1, ICAM-2, ICAM-3, ICAM-4, ICAM-5, E-selectin, collagen I, JAM-A*
$\alpha_M\beta 2$	CD11b/CD18, Mac-1, Mo-1, CR3	monocytes, neutrophils, subset of cytotoxic T-lymphocytes	ICAM-1, ICAM-2, ICAM-4, iC3b, GPIIb $\alpha$ *, fibrinogen, VWF*, factor X, heparin, CNTF*, NIF, LPS*, CD23, uPAR*, CD14, CD16, Cyr61, GP63, JAM-C
$\alpha_X\beta 2$	CD11c/CD18, p150,95, CR4	monocytes, neutrophils, eosinophils, subset of cytotoxic T-lymphocytes	iC3b, fibrinogen, LPS, CD23, collagen I, uPAR, JAM-C
$\alpha_D\beta 2$	CD11d/CD18	monocytes	ICAM-3, VCAM*

\* CNTF, ciliary neurotrophic factor; GPIIb $\alpha$ , glycoprotein IIb; JAM, junction adhesion molecule; LPS, lipopolysaccharide; uPAR, urokinase-type plasminogen activator receptor; VCAM, vascular cell adhesion molecule; VWF, Von Willebrand Factor.

## Amyloid $\beta$ (A $\beta$ ) Peptides in Brain

Amyloid Precursor Protein (APP) is a type I transmembrane protein present in *e.g.* neurons. Due to alternative splicing three isoforms are produced that comprise 770, 751, or 695 amino acids. The largest two isoforms contain a Kunitz-type domain at the C-terminus. At the cell surface, APP can be proteolytically cleaved by  $\alpha$ - or  $\beta$ -secretases (*Fig 4*), resulting in release of a fragment (called protease nexin-2) from the cell surface into the cerebrospinal fluid. The carboxy-terminal fragment that remains associated with the membrane, can be further processed to generate A $\beta$  peptides, which are constitutively secreted into the extracellular space<sup>124</sup>. Deposition of A $\beta$  peptides in the brain is central to the pathogenesis of Alzheimer's Disease (AD). Therefore, it is important that A $\beta$  peptides are eliminated from brain tissue to prevent their neurotoxic accumulation<sup>124;125</sup>.

LRP is responsible for binding and internalization of both transmembrane and soluble APP<sup>126;127</sup>. When the transmembrane form that contains a Kunitz-type domain, is internalized by LRP, intracellular processing of APP can result in increased secretion of A $\beta$  peptides<sup>126</sup>. The intracellular adaptor protein FE65 is thought to bridge the intracellular tails of LRP and APP<sup>34</sup>. In contrast to this promotional effect of LRP on the progression of AD, LRP can also serve as a protector. By using a murine model, Shibata et al<sup>128</sup> showed that LRP plays an important role in the clearance of A $\beta$  peptide from the brain into plasma. Although LRP is also present in neurons, LRP does not seem to be involved in clearance of

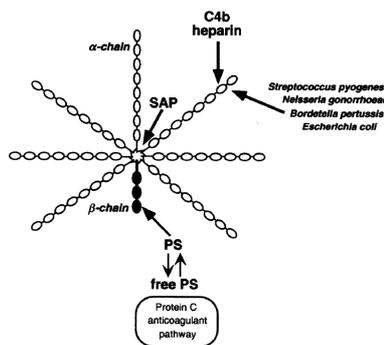


**Figure 4. Intracellular processing of APP.** A. The largest isoform is 770 amino acids (aa), which contains an alternatively spliced exon of 56 aa at position 286. Alternatively, this exon can also comprise 19 aa, resulting in the 751 aa isoform. The A $\beta$  peptide includes 28 aa of the extracellular domain and 12-14 aa of the transmembrane domain (TM). B.  $\alpha$ -secretase activity results in cleavage at position 687. The remaining transmembrane and intracellular part can be further processed at position 711 or 713 by  $\gamma$ -secretases, resulting in the formation of p3 peptides. C. Alternative cleavage at position 671 by  $\beta$ -secretases results in retention of a 99 aa fragment, which can undergo subsequent cleavage by  $\gamma$ -secretases. This action releases the A $\beta$  peptides. Figure adapted from Selkoe<sup>124</sup>.

A $\beta$  peptides in the prevention of A $\beta$  peptide deposition in these cells<sup>129</sup>. A $\beta$  peptides can also be internalized as part of a complex with  $\alpha_2$ M or ApoE<sup>130</sup> – two LRP ligands that are also associated with AD<sup>131</sup>. The relation between LRP and the development of AD is further exemplified by the observation that reduced LRP expression has been noted in the brains of humans and mice suffering from AD<sup>128;132</sup>.

## C4b-binding Protein (C4BP)

C4BP is a plasma glycoprotein (150 mg/L) of approximately 570 kDa that is expressed in the liver<sup>133-135</sup>. C4BP has a heterogeneous structure: it contains 6 or 7 identical  $\alpha$ -chains and 80% also contains a unique  $\beta$ -chain<sup>136;137</sup> (Fig 5). Each  $\alpha$ -chain has a molecular weight of 70 kDa, whereas the  $\beta$ -chain is 45 kDa. The chains are interconnected via disulfide bridges, resulting in a spider-like structure<sup>138</sup>. The  $\alpha$ - and  $\beta$ -chains are composed of 8 or 3 so-called complement control protein (CCP)- or Sushi domains, respectively.



**Figure 5. Structure of C4BP $\alpha\beta$ .** C4BP contains 6 or 7  $\alpha$ -chains and 0 or 1  $\beta$ -chains (as depicted in this figure). Also shown are ligands of both  $\alpha$ - and  $\beta$ -chain. Figure adapted from Blom et al<sup>137</sup>.

C4BP has a dual function, since it is associated with both complement and coagulation pathway. In the complement pathway, it controls C4b-mediated reactions in several ways. First, it acts as a cofactor for factor I in the degradation of C4b and C3b<sup>139;140</sup>. Second, C4BP prevents the assembly of C3-convertase by binding C4b, and accelerates the natural decay of the complex<sup>141</sup>. The  $\alpha$ -chains bind several proteins, like C4b<sup>138;142</sup>, heparin<sup>139;143</sup>, serum amyloid P<sup>144</sup> and components of pathogenic bacteria<sup>145</sup>. The  $\beta$ -chain harbors a high affinity binding site for protein S, which acts as a cofactor for anticoagulant activated protein C in the hemostatic process<sup>146;147</sup>.

C4BP is an acute phase protein. Upon inflammation its concentration increases 7-fold. Mainly the C4BP $\alpha$ -isoform is raised, which has the consequence that due to stable C4BP $\alpha\beta$  levels, the free protein S levels do not decrease during inflammation. To this end, inflammation and coagulation are regulated independently<sup>148</sup>.

Recently, LRP was found to be a receptor for C4BP, thereby mediating its cellular uptake *in vitro* and *in vivo*<sup>149</sup>. In addition, the interaction could be inhibited by heparin, suggesting overlapping binding sites for heparin and LRP.

## Outline of this Thesis

From historical perspective, LRP has been considered as an endocytic receptor that recognizes a multitude of ligands. More recent studies have demonstrated that LRP exerts other (often cell-type restricted) functions as well. The aim of the present thesis was to further explore the various aspects of LRP biology.

- 1 LRP has been found to be expressed in granulocytes and monocytes/macrophages. Although the contribution of LRP to macrophage function has been studied to some extent, little is known concerning the role of LRP in monocytes and granulocytes. In chapter 2, we describe studies in which we searched for potential cell-surface-located components in granulocytes and monocytes that interact with LRP. We were able to identify  $\beta$ 2-integrins as partners for LRP, and showed that complex formation between both receptors is a prerequisite for optimal leukocyte adhesion to endothelial cells.
- 2 Despite the abundant literature on the expression patterns of LRP, the presence of LRP in the various leukocyte subsets has remained a poorly investigated issue. In view of its functional importance in cells of the myeloid lineage, it was of interest to elucidate to what extent LRP is expressed in lymphocytic cells. The studies are described in chapter 3, and demonstrated the presence of LRP in CD4+ and CD8+ T-lymphocytes, B-lymphocytes and Natural Killer cells. One remarkable observation in our studies was that LRP is maintained intracellularly in T-lymphocytes, whereas in B-lymphocytes and Natural Killer cells LRP is constitutively present at the cell surface. In addition, we found that LRP is instantly redistributed to the cell surface upon incubation of the T-lymphocytes with allogenic antigen-presenting cells. The functional consequence of this redistribution awaits further studies.
- 3 There is both biochemical and genetic evidence that LRP is linked to the pathogenesis of Alzheimer's Disease. However, the functional mechanism underlying this link is still unclear. Studies in this regard are complicated by the fact that not only LRP, but also some of its ligands play a role in the development of Alzheimer's Disease. Moreover, LRP is present in various cell types (*ie* neurons and brain capillary endothelium) that contribute to amyloid plaque formation and clearance of amyloid  $\beta$ -peptides. In chapter 4,

studies that focus on the role of LRP present in brain endothelium cells in the clearance of amyloid  $\beta$ -peptides from brain to the circulation are presented. These studies show that high concentrations of amyloid  $\beta$ -peptides direct a downregulation of LRP in brain endothelium cells, resulting in decreased transport of such peptides through the blood-brain barrier and their accumulation in neurotoxic plaques.

- 4 Recently, we have identified C4b-binding protein (C4BP) as a ligand for LRP *in vitro* and *in vivo*. Since binding of C4BP to LRP is inhibited in the presence of heparin, we considered the possibility that patches of positively charged amino acids within the C4BP  $\alpha$ -chain contribute to this interaction. In chapter 5, we describe our studies in this regard. It is shown that mutations within the heparin-binding site of C4BP  $\alpha$ -chains result in reduced binding to LRP. In addition, we have found that the cellular uptake of C4BP by LRP is preceded by binding of C4BP to cell-surface-exposed heparan sulfate proteoglycans.

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chapter **2**

**LDL receptor-related protein regulates  $\beta$ 2-integrin-mediated leukocyte adhesion**

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*Blood*, 2005, 105(1):170-177

## Abstract

$\beta$ 2-Integrin clustering upon activation is a key event in leukocyte adhesion to the endothelium during the inflammatory response. In the search for molecular mechanisms leading to this clustering, we have identified low-density lipoprotein (LDL) receptor-related protein (LRP) as a new partner for  $\beta$ 2-integrins at the leukocyte surface. Immobilized recombinant LRP fragments served as an adhesive surface for blood-derived leukocytes and the U937 cell line. This adhesion was decreased up to 95% in the presence of antibodies against  $\beta$ 2-integrins, pointing to these integrins as potential partners for LRP. Using purified proteins, LRP indeed associated with the  $\alpha_M\beta_2$  complex and the  $\alpha_M$  and  $\alpha_L$  I-domains ( $K_{d,app} \approx 0.5 \mu\text{M}$ ). Immunoprecipitation experiments and confocal microscopy revealed that endogenously expressed LRP and  $\alpha_L\beta_2$  colocalized in monocytes and U937 cells. Furthermore, activation of U937 cells resulted in clustering of  $\alpha_L\beta_2$  and LRP to similar regions at the cell surface, indicating potential cooperation between both proteins. This was confirmed by the lack of  $\alpha_L\beta_2$  clustering in U937 cells treated by antisense oligonucleotides to downregulate LRP. In addition, the absence of LRP resulted in complete abrogation of  $\beta$ 2-integrin-dependent adhesion to endothelial cells in a perfusion system, demonstrating the presence of a previously unrecognized link between LRP and leukocyte function.

## Introduction

Low-density lipoprotein (LDL) receptor-related protein (LRP), also known as  $\alpha$ <sub>2</sub>-macroglobulin receptor or CD91<sup>1</sup>, is a member of the LDL-receptor family. It consists of an 85-kDa intracellular and transmembrane domain that is non-covalently linked to a 515-kDa extracellular domain<sup>2</sup>. The extracellular domain comprises 4 clusters of complement-type repeats, 2 of which (clusters II and IV) play a dominant role in ligand binding<sup>3;4</sup>. At present, more than 30 structurally and functionally unrelated ligands have been identified for this receptor (for a review, see Herz and Strickland<sup>5</sup>), suggesting that LRP is involved in a diverse range of (patho)physiological processes.

The intracellular domain of LRP harbors 2 NPXY-motifs and 1 YXXL motif; the latter is the main motif that controls internalization of LRP<sup>6</sup>. In its function as an endocytic receptor, LRP mediates the cellular uptake of both circulating and membrane-associated proteins, which are subsequently degraded in lysosomes<sup>7</sup>. Alternatively, ligands can be transcytosed<sup>8</sup> or transported to the nucleus<sup>9</sup>. Apart from its endocytic function, LRP has been shown to be involved in signaling pathways<sup>10</sup>. In this respect, various intracellular adaptor proteins have been identified that interact with the cytoplasmic tail of LRP through its NPXY motif<sup>11</sup>. Moreover, binding of adaptor proteins to the cytoplasmic tail inhibits the internalization of LRP<sup>11</sup>. This allows LRP to form heterodimeric complexes with other receptors at the cell surface, such as N-methyl-D-aspartate receptor<sup>12</sup> and platelet-derived growth factor receptor<sup>13;14</sup>. In these cases, the presence of LRP is essential for appropriate function of the co-receptor.

LRP is expressed in a variety of cell types, including hepatocytes, fibroblasts, neurons and smooth muscle cells<sup>15</sup>. Among blood cells, LRP is expressed in leukocytes, including polymorphonuclear cells (PMNs) and monocytes, but not in erythrocytes or platelets<sup>15;16</sup>. PMNs and monocytes are essential in the inflammatory process. Leukocytes circulate in blood in a resting state and become activated in response to inflammatory stimuli. Leukocyte activation may induce adhesion to the vascular wall and subsequent migration to inflammatory sites.

Firm adhesion and subsequent transmigration involves adhesion molecules, such as  $\beta$ 2-integrins.  $\beta$ 2-Integrins are specifically expressed in leukocytes, and 4 isotypes are known<sup>17</sup>.  $\beta$ 2-integrins consist of a common  $\beta$ 2-chain (CD18), which is non-covalently linked to an  $\alpha$ -chain, namely  $\alpha$ <sub>L</sub> (CD11a),  $\alpha$ <sub>M</sub> (CD11b),  $\alpha$ <sub>X</sub> (CD11c), or  $\alpha$ <sub>D</sub> (CD11d). The  $\alpha$ -chains contain an I-domain, which harbors the main ligand-binding site<sup>18-20</sup>. Although all  $\beta$ 2-integrin isoforms interact with a wide subset of proteins, ligand recognition appears to be specific among the isoforms. The main ligands of  $\beta$ 2-integrins expressed at the cell surface of other cell types are the intercellular adhesion molecules (ICAMs). At present, 5 ICAMs

have been described with slightly different binding specificities<sup>21</sup>. For instance, ICAM-1, -2, and -4 associate with  $\alpha_L$  and  $\alpha_M$ , whereas ICAM-3 and -5 are only recognized by  $\alpha_L$ <sup>21-23</sup>. Furthermore,  $\alpha_L$  specifically binds to junction adhesion molecule-1<sup>24</sup> and  $\alpha_M$  binds to junction adhesion molecule-3 and glycoprotein Ib<sup>25-28</sup>. Other  $\beta_2$ -integrin ligands are fibrinogen, collagen type I, iC3b, and neutrophil inhibitory factor<sup>20;29;30</sup>.

Whereas the role of  $\beta_2$ -integrins in leukocyte function has been well studied, little is known about the role of LRP in this connection. The present study focused on the identification of leukocyte-surface proteins that associate with LRP. Our results show that LRP is able to interact with  $\beta_2$ -integrins. Moreover, LRP appears to regulate  $\alpha_L\beta_2$ -integrin clustering and, as such,  $\beta_2$ -integrin-mediated adhesion to endothelial cells. These observations identify a previously unrecognized link between LRP and the inflammatory system.

## Experimental Procedures

### Materials

Cell culture medium RPMI 1640, Dulbecco modified Eagle medium (DMEM)/F-12, penicillin, streptomycin, and L-glutamine were obtained from Gibco Life Technologies (Paisley, United Kingdom). Fetal bovine serum was from Cambrex Bio Science (Verviers, Belgium). Microtiter plates were from Costar (New York, NY) or Nunc (Roskilde, Denmark). The Biacore2000 system and required reagents were from Biacore AB (Uppsala, Sweden). Phorbol-12-myristate-13-acetate (PMA), methotrexate, P-nitrophenyl phosphate (PNP), polyvinylpyrrolidone-360 (PVP-360), and bovine serum albumin (BSA) fraction V were purchased from Sigma (St Louis, MO).

### Antibodies and proteins

The following antibodies were used: anti- $\alpha_L$  clone 38 (R&D Systems, Minneapolis, MN), anti- $\alpha_M$  clone 44 (BD PharMingen, San Diego, CA) and M1/70 (R&D Systems), anti- $\beta_2$  clone 68-5A5 (Cymbus, Hants, United Kingdom) and R2E7B<sup>31</sup>, goat anti-LRP A-18 (Santa Cruz Biotechnology, Santa Cruz, CA), anti-LRP clones 2-M-R-II2C7, 2-M-R-II4/8 and anti-LRP-fluorescein isothiocyanate (FITC) clone 2-M-R-I4C2 (Biomac, Leipzig, Germany), goat anti-mouse-FITC (Becton Dickinson, San Jose, CA), anti-urokinase-type plasminogen activator receptor (uPAR) clone 62022 (R&D Systems), anti-4 clone HP2/1 (Immunotech, Westbrook, ME), donkey anti-goat-FITC and donkey anti-mouse-TexasRed (both from Jackson ImmunoResearch, West Grove, PA). Receptor-associated protein<sup>32</sup> was purified as a glutathione-S-transferase fusion protein (GST-RAP), as described<sup>33</sup>. Purified

full-length LRP and stable cell lines expressing recombinant LRP fragments (LRP clusters II and IV) were kindly provided by Dr H. Pannekoek (Department of Biochemistry, University of Amsterdam, The Netherlands). Clusters II and IV were purified from the cell culture supernatant using a GST-RAP sepharose column, as reported previously<sup>4</sup>. The  $\alpha_M\beta_2$  complex and the I-domains of the  $\alpha_M$  and  $\alpha_L$  subunit fused to GST were purified as described<sup>34</sup>. Human multimeric vitronectin was purified as described<sup>35</sup>.

#### **Cell lines and culture conditions**

Human umbilical vein endothelial cells (HUVECs) were isolated according to the method of Jaffe et al<sup>36</sup> with some modifications<sup>37</sup>. Only first and second passages were used for experiments. The monocytic line U937<sup>38</sup> was obtained from the American Type Culture Collection (Manassas, VA) (CRL-1593.2) and was maintained in RPMI 1640, 10% fetal bovine serum, 50 U/mL penicillin, 50  $\mu$ g/mL streptomycin, and 50  $\mu$ M  $\beta$ -mercaptoethanol in a humidified CO<sub>2</sub> (5%) incubator at 37°C. To downregulate LRP protein expression, U937 cells (5.0  $\times$  10<sup>5</sup> cells/mL) were incubated with phosphorothioate antisense oligodeoxynucleotides at a concentration of 20  $\mu$ M for 2 days. Fresh oligonucleotides (10  $\mu$ M) were added every 24 hours. The sequence of the antisense oligonucleotide was 5'-CGGCGGGGTCAGCAT-3', which is complementary to the initiation site on LRP mRNA<sup>39</sup>. As a control, oligonucleotides having the corresponding sense sequence 5'-ATGCTGACCCCGCCG-3' were used<sup>39</sup>. LRP expression was examined by confocal scanning fluorescence microscopy (see "Confocal scanning fluorescence microscopy"). PMNs were freshly isolated from blood obtained from healthy volunteers by Ficoll-Paque (Amersham-Pharmacia, Uppsala, Sweden) density centrifugation. Erythrocytes were removed from the granulocyte fraction by ice-cold erythrocyte lysis buffer (0.155 mM NH<sub>4</sub>Cl, 7.4 mM KHCO<sub>3</sub>, and 0.1 mM EDTA [ethylenediaminetetraacetic acid], pH 7.4). Peripheral blood monocytes were isolated from the mononuclear cell fraction using CD14 microbeads and AutoMACS (Miltenyi-Biotec, Bergisch-Gladbach, Germany). After isolation, cells were directly used for experiments. Cell purity was greater than 95% for PMNs and 90% for monocytes, as examined by CD15 and CD14 detection by flow cytometric analysis, respectively.

#### **Static cell adhesion**

In static cell adhesion experiments, LRP clusters II and IV (50  $\mu$ g/mL) were immobilized in microtiter wells in Tris-buffered saline (pH 7.4) for 16 hours at 4°C. Alternatively, HUVECs were grown in microtiter wells until confluence. HUVECs were stimulated with tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ) (100 U/mL) for 4 hours and were subsequently fixed for 15 minutes with 4% paraformaldehyde. Wells

were blocked with either 5% BSA (U937 cells) or 0.5% PVP-360 (PMNs/monocytes) for 1 hour at 37°C. Cells ( $2 \times 10^6$ /mL) were washed twice with phosphate-buffered saline (PBS) and were activated with 100 nM PMA in DMEM/F-12 supplemented with 0.1% BSA and 1 mM  $\text{MnCl}_2$  for 15 minutes. Where indicated, cells were preincubated for 15 minutes with specific blocking antibodies (20  $\mu\text{g}/\text{mL}$ ) against different integrin subunits, or wells coated with LRP cluster II or IV were incubated with 50  $\mu\text{g}/\text{mL}$  GST-RAP in the presence of 3 mM  $\text{CaCl}_2$  for 15 minutes before the addition of cell suspensions. Cells ( $1.5 \times 10^5$ /well) were incubated in the microtiter plates for either 60 minutes at 37°C (U937 cells to clusters II and IV) or 30 minutes at room temperature (PMNs/monocytes to clusters II and IV; U937 cells to HUVECs). Non-bound cells were removed by gently washing wells with PBS. Adherent cells were determined by endogenous alkaline phosphatase activity using PNP as a substrate (3 mg/mL in 1% Triton-X100/50 mM acetic acid (pH 5)). Optical density was measured at 405 nm. Alternatively, cells were visualized using light microscopy (Leitz Diaplan; Leica, Rijswijk, The Netherlands) and computer-assisted analysis with OPTIMAS 6.0 software (DVS, Breda, The Netherlands).

### **Surface plasmon resonance analysis**

Surface plasmon resonance (SPR) binding assays were performed using a Biacore2000 biosensor system. LRP was immobilized on a CM5-sensorchip at a density of 7.7 fmol/ $\text{mm}^2$  using the amine-coupling kit, as described by the manufacturer. A control channel was routinely activated and blocked in the absence of protein. Binding of GST/I-domain fusion proteins to LRP-coated channels was corrected for binding to noncoated channels (less than 5% of binding to coated channels). SPR analysis was performed in 100 mM NaCl, 0.005% Tween-20, 2 mM  $\text{CaCl}_2$ , 2 mM  $\text{MnCl}_2$ , 25 mM HEPES (N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid) (pH 7.4) at 25°C with a flow rate of 5  $\mu\text{L}/\text{min}$ . Regeneration of the sensorchip surface was performed by incubating with 0.1 M  $\text{H}_3\text{PO}_4$  for 2 minutes at a flow rate of 5  $\mu\text{L}/\text{min}$ . Data obtained from steady state SPR analysis were used for the calculation of the apparent affinity constants ( $K_{d,app}$ ), as described<sup>40</sup>.

### **Immunoprecipitations**

U937 cells ( $6 \times 10^6$ ) were stimulated for 15 minutes with PMA and were lysed in PBS containing 1% Nonidet-40 and 0.5% DOC at 4°C for 1 hour. Lysates were clarified by centrifugation and precleared by incubation with Protein G-Sepharose. The lysate was immunoprecipitated with anti-LRP or anti- $\alpha_M$  antibodies (1  $\mu\text{g}/\text{mL}$ ) and Protein A- or Protein G-Sepharose, respectively, for 16 hours at 4°C. Immunocomplexes were pelleted, washed, and resolved by sodium do-

decyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and Western blotting. Detergent-resistant membrane fractions were isolated as described<sup>41</sup> and immunocomplexes were prepared similarly.

#### **Confocal scanning fluorescence microscopy**

U937 cells or blood monocytes were collected on object glasses by cytospin centrifugation ( $10^5$  cells/spin). Cells were fixed in 4% paraformaldehyde/PBS and were blocked in 2% BSA/0.1% saponin/PBS. Localization of the  $\alpha_L$  subunit and LRP was detected using mouse anti- $\alpha_L$  (1:10) and goat anti-LRP (1:20) antibodies, followed by incubation with donkey anti-mouse (1:100) and donkey anti-goat (1:200) antibodies, which were labeled with TexasRed and FITC, respectively. Cells were mounted in mowiol containing 2.5% 1,4-diazabicyclo[2.2.2]octane. Cells were visualized using Leica DMIRB confocal scanning laser microscope equipped with a 63x/1.40 Plan APO objective lens and a TCS 4D system (Leica, Voorburg, The Netherlands).

#### **Cell adhesion under flow conditions**

HUVECs were coated on glass coverslips, grown until confluence, and stimulated with TNF- $\alpha$  (100 U/mL) for 4 hours before perfusion. Cells ( $2 \times 10^6$  cells/mL) were perfused over HUVECs for 10 minutes at a flow rate of 100  $\mu$ L/min. Wall shear stress was calculated to be 0.8 dyne/cm<sup>2</sup>. During perfusion, the flow chamber<sup>42;43</sup> was mounted on a microscope stage (Axiovert 25; Zeiss, Oberkochen, Germany), which was equipped with a black-and-white charge-coupled device videocamera (Sanyo, Osaka, Japan) and was coupled to a VHS videorecorder. Video images were evaluated for the number of adherent cells, with dedicated routines made in the image analysis software OPTIMAS 6. U937 cells (non-transfected and transfected with sense or antisense LRP-oligonucleotides) that were in contact with the surface appeared as bright white-centered cells after proper adjustment of the microscope during recording.

#### **Statistical analysis**

All data are expressed as mean  $\pm$  SD, unless stated otherwise. Between-group variations were examined using the Student's t test. A P value of less than .05 was considered statistically significant.

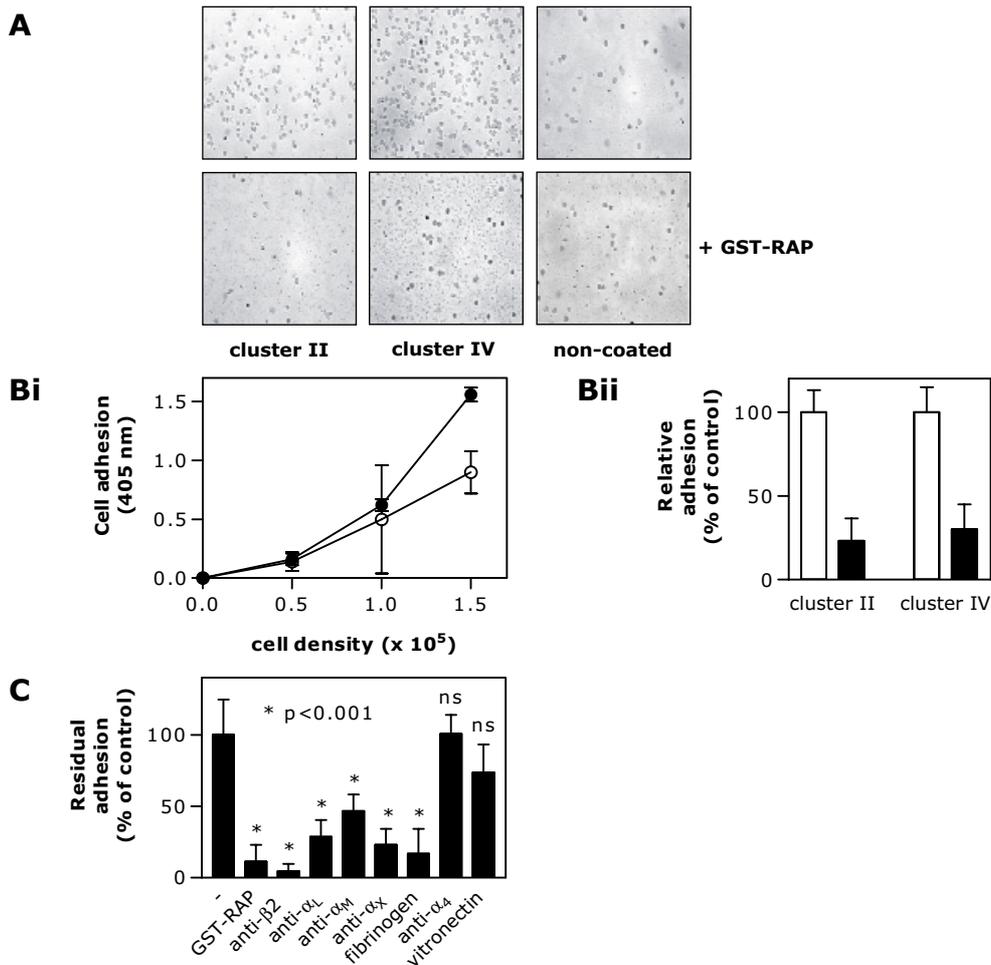
## Results

### Leukocytes adhere to immobilized LRP fragments

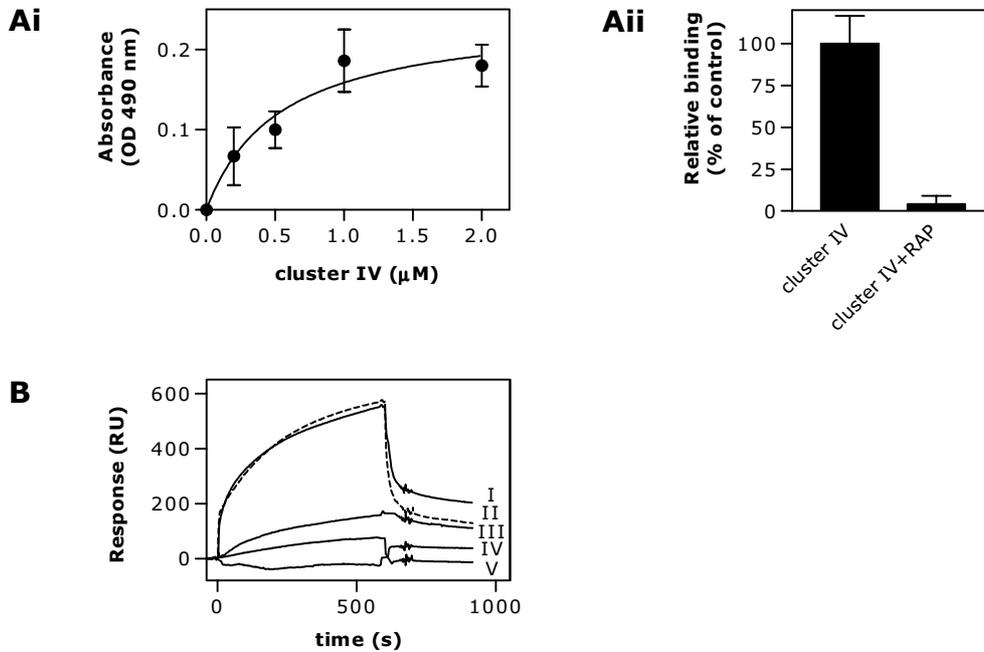
To investigate whether leukocytes express surface proteins that interact with LRP, adhesion of blood monocytes and PMNs, and of the monocytic line U937 to immobilized ligand-binding LRP fragments (ie, clusters II and IV) was determined. Freshly isolated monocytes were stimulated with PMA and were added to wells coated with cluster II or IV. Each recombinant fragment appeared to provide an adhesive surface for monocytes, as visualized by light microscopy (*Fig 1A*). Dose-dependence was subsequently tested by measuring endogenous phosphatase activity of adhered cells, which revealed that adhesion to clusters II and IV was cell number dependent (*Fig 1Bi*). In addition, adhesion to these LRP fragments was decreased by 74% and 71% in the presence of GST-RAP, respectively (*Fig 1Bii*). Similar data were obtained for PMNs (data not shown) and U937 cells (*Fig 1C*), indicating that leukocytes express LRP-binding elements at the surface.

### Binding of U937 cells to LRP fragments involves $\beta$ 2-integrins

Binding of leukocytes to LRP cluster IV was examined in more detail using U937 cells. First, GST-RAP was observed to diminish the adhesion of U937 cells to cluster IV by 89% (*Fig 1C*). Activated leukocytes are characterized by the presence of active adhesion molecules, such as  $\alpha_4\beta_1$ ,  $\alpha_v\beta_3$ , and  $\beta$ 2-integrins. To assess the contribution of these integrins in the binding of U937 cells to LRP fragments, adhesion was examined in the presence of potential inhibitors. Anti- $\alpha_4$ -integrin antibodies and the  $\alpha_v\beta_3$ -ligand vitronectin were unable to reduce adhesion (*Fig 1C*). In contrast, a  $\beta$ 2-integrin-directed antibody inhibited the binding of stimulated U937 cells to cluster IV by 95% (*Fig 1C*). Given that  $\beta$ 2-integrins consist of a heterodimeric complex with distinct  $\alpha$ -subunits, we also examined the effect of antibodies directed against various  $\alpha$ -subunits on cell adhesion. These antibodies decreased adhesion to cluster IV up to 77% (*Fig 1C*). Adhesion was also reduced in the presence of the  $\beta$ 2-integrin ligand fibrinogen (*Fig 1C*). It should be noted that similar data were obtained when adhesion to cluster II was tested (data not shown). Thus, it appears that LRP provides a binding site for  $\beta$ 2-integrins and that  $\alpha_L\beta_2$ ,  $\alpha_M\beta_2$ , and  $\alpha_X\beta_2$  are able to associate with LRP.



**Figure 1. Adhesion of monocytes to LRP clusters II and IV.** Freshly isolated monocytes were stimulated with 100 nM PMA for 15 minutes and added to immobilized LRP cluster II or IV for 30 minutes at room temperature in the presence or absence of LRP-antagonist GST-RAP (50  $\mu$ g/mL). **A.** Typical experiment visualized by light microscopy, in which  $1.5 \times 10^5$  cells were added to each well. Original magnification, 400x. **Bi.** To quantify cell adhesion, different amounts of cells ( $0-1.5 \times 10^5$  cells/well) were added. After incubation and subsequent washing, bound cells were lysed using 1% Triton-X100/50 mM acetic acid (pH 5.0), and endogenous alkaline phosphatase activity was determined using PNP as substrate. Open circles indicates LRP cluster II; closed circles LRP cluster IV. **Bii.** Relative adhesion in the presence (black bars) or absence (white bars) of GST-RAP. Data are corrected for adhesion to uncoated wells (less than 20% of cluster IV coated wells) and represent mean  $\pm$  SD of 3 experiments performed in duplicate. **C.** PMA-stimulated U937 cells ( $1.5 \times 10^6$ ) were incubated with indicated antibodies (20  $\mu$ g/mL), fibrinogen (50  $\mu$ g/mL), or vitronectin (100  $\mu$ g/mL) for 15 minutes, or wells were preincubated with GST-RAP (50  $\mu$ g/mL) and added to immobilized IV for 60 minutes at 37°C. Adhered cells were detected as described in panel B. Presented is the percentage of adhesion relative to adhesion in the absence of antibodies or GST-RAP. Data represent the mean  $\pm$  SD of 3 to 10 experiments performed in duplicate. ns indicates not significant ( $p > .05$ ).



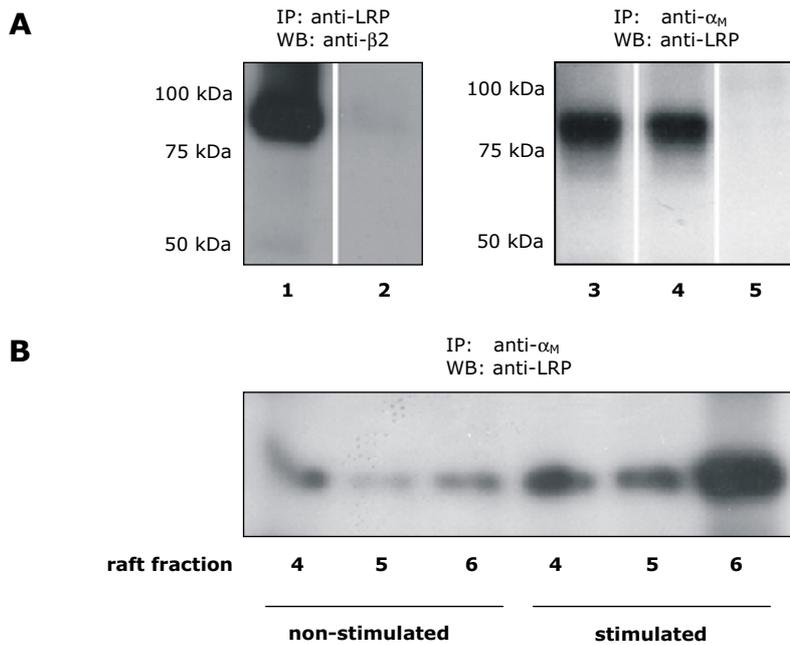
**Figure 2. Complex formation between LRP and  $\alpha_M\beta_2$ .** *Ai.* Purified recombinant cluster IV (0-2.0  $\mu\text{M}$ ) was incubated with immobilized  $\alpha_M\beta_2$  complex (2.5  $\mu\text{g}/\text{well}$ ) in Tris-buffered saline/3 mM  $\text{CaCl}_2$ /1 mM  $\text{MnCl}_2$  (pH 7.4) for 2 hours at 37°C. Bound cluster IV was subsequently determined using peroxidase-labeled polyclonal antibodies directed against cluster IV. Data represent mean  $\pm$  SEM of 4 experiments. *Aii.* Binding of 200 nM cluster IV to immobilized  $\alpha_M\beta_2$  in the presence or absence of a 10-fold excess of GST-RAP. *B.* 500 nM purified recombinant I-domain of the  $\alpha_M$ - (line I) or the  $\alpha_L$ -subunit (line II) were perfused over LRP immobilized onto a CM5-sensorchip (7.7 fmol/ $\text{mm}^2$ ) in 100 mM NaCl, 0.005% Tween-20, 2 mM  $\text{CaCl}_2$ , 2 mM  $\text{MnCl}_2$ , 25 mM HEPES (pH 7.4) at a flow rate of 5  $\mu\text{L}/\text{min}$  for 20 minutes at 25°C. Line III: Before perfusion, 500 nM  $\alpha_M$  was preincubated with a 5-fold molar excess of fibrinogen for 30 minutes. Line IV: 5  $\mu\text{M}$  fibrinogen. Line V: 500 nM GST. Ligand solution was replaced with buffer 10 minutes after injection to initiate dissociation. Depicted are sensorgrams corrected for specific binding, which was less than 5% of binding to LRP-coated channels.

### **LRP comprises a binding site for $\beta$ 2-integrins**

To assess the interaction between LRP and  $\beta$ 2-integrins at the level of purified proteins, binding of LRP cluster IV to the  $\alpha_M\beta$ 2 complex was assessed in a qualitative manner using an immunosorbent assay. Various concentrations of LRP cluster IV (0-2  $\mu$ M) were incubated with immobilized  $\alpha_M\beta$ 2 (2.5  $\mu$ g/well), and bound cluster IV was subsequently determined using polyclonal anti-cluster IV antibodies. Cluster IV bound to immobilized  $\alpha_M\beta$ 2 in a dose-dependent and saturable manner, and half-maximum binding was observed at a concentration of 0.5  $\mu$ M cluster IV (*Fig 2Ai*). Furthermore, binding of cluster IV to the immobilized complex could be blocked in the presence of GST-RAP (*Fig 2Aii*). In a second approach, we investigated whether the  $\alpha$ -subunits of the  $\alpha_L\beta$ 2 or  $\alpha_M\beta$ 2 complex contribute to the interaction with LRP. Therefore, SPR analysis was performed using LRP and the recombinant I-domains of both subunits fused to GST. SPR analysis demonstrated that both I-domain/GST fusion proteins associate with immobilized full-length LRP in a reversible manner (*Fig 2B, lines I and II*), whereas GST alone did not bind to LRP (*Fig 2B, line V*). In addition, binding of the  $\alpha_M$  I-domain to LRP was inhibited in the presence of the  $\alpha_M$  ligand fibrinogen (*Fig 2B, line III*). Steady state analysis further indicated that the  $\alpha_L$  and  $\alpha_M$  I-domains both interact with LRP with an apparent affinity constant of 0.5  $\mu$ M. These data demonstrate that  $\beta$ 2-integrins directly interact with LRP and that binding is at least in part mediated by the  $\alpha$ -subunits.

### **LRP and $\beta$ 2-integrins are targeted to detergent-insoluble membrane regions**

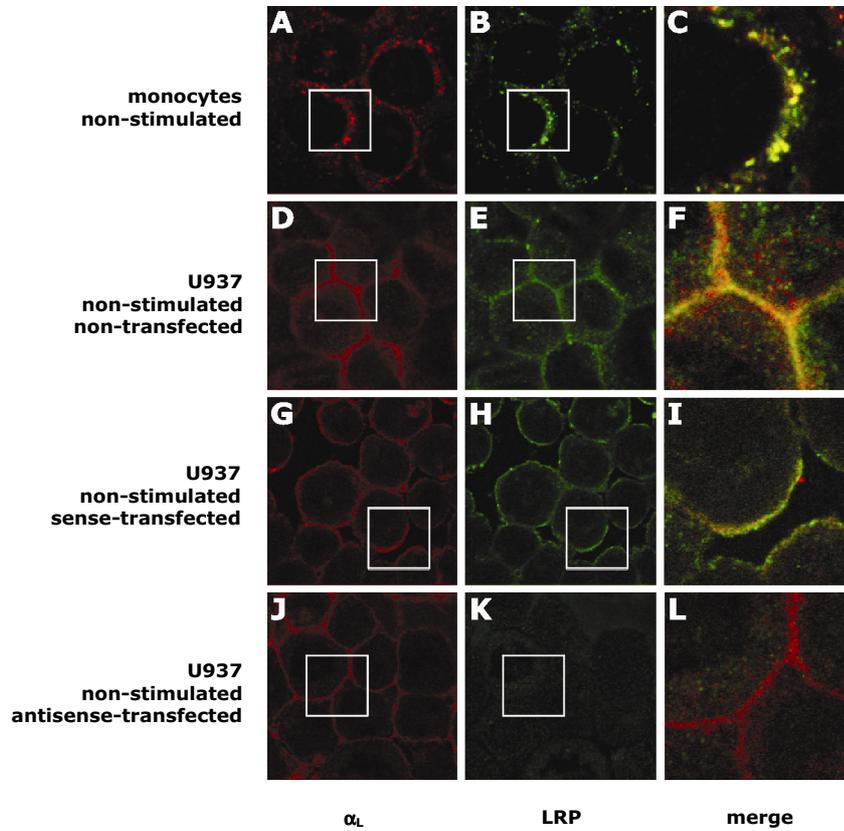
To address the possibility that both receptors associate in the cellular environment, complex formation was assessed in immunoprecipitation experiments using U937 cells. With non-stimulated U937 cells, the  $\beta$ 2-subunit was readily immunoprecipitated using an antibody against LRP but not an isotype-matched control antibody (*Fig 3A, lanes 1 and 2*). In turn, an anti- $\alpha_M$  antibody, but not a control antibody, was able to immunoprecipitate LRP (*Fig 3A, lanes 3 and 5*). LRP coprecipitated with an anti- $\alpha_M$ -integrin antibody to the same extent when cells were stimulated with PMA for 1 hour (*Fig 3A, lane 4*). Because leukocyte stimulation is associated with the redistribution of  $\beta$ 2-integrins at the cell surface to cholesterol-enriched regions<sup>44</sup>, we further examined the presence of LRP/ $\beta$ 2-integrin complexes in such membrane fractions of resting and stimulated U937 cells. In detergent-insoluble fractions of non-stimulated cells, some coprecipitation of LRP with  $\beta$ 2-integrins was observed (*Fig 3B, left 3 lanes*). An increase in coprecipitated LRP was found on stimulation of U937 cells with PMA (*Fig 3B, right 3 lanes*), suggesting that the LRP/ $\beta$ 2-integrin complex is targeted to cholesterol-enriched membrane regions.



**Figure 3. Coimmunoprecipitation of LRP and  $\beta$ 2-integrins.** *A.* U937 cells were lysed for 1 hour on ice and incubated with Protein A-Sepharose and anti-LRP clone  $\alpha$ 2-M-R-II2C7 (lane 1) or an isotype control (lane 2) at 4°C overnight. Beads were washed extensively and boiled to release bound proteins. Samples were analyzed by SDS-PAGE and Western blotting using anti- $\beta$ 2 (clone R2E7B) and peroxidase-conjugated rat anti-mouse. Non-stimulated (lane 3) or stimulated (lane 4) cells were lysed and incubated with protein G-Sepharose and an anti- $\alpha_M$  antibody (clone M1/70) (lanes 3 and 4) or an isotype control (lane 5). Precipitated proteins were analyzed by SDS-PAGE and Western blotting using anti-LRP antibody (clone  $\alpha$ 2-M-R-II4/8). *B.* Lipid rafts were isolated from non-stimulated or stimulated cells, as described elsewhere<sup>41</sup>. Immunoprecipitations were performed as described in the legend of Figure 3A.

### Surface expression of $\alpha_L\beta$ 2 is independent of LRP

The subcellular localization of endogenously expressed LRP and  $\beta$ 2-integrins was further studied by confocal immunofluorescence microscopy. First, the presence of  $\beta$ 2-integrins at the cell surface in blood-derived monocytes or the U937 cell line was confirmed using anti- $\alpha_L$  antibodies (*Fig 4A, D*). Similar surface staining was observed for LRP (*Fig 4B, E*), which overlapped with that of  $\alpha_L$  to a significant extent (*Fig 4C, F*). To investigate whether the surface location of  $\alpha_L\beta$ 2-integrin was linked to that of LRP, U937 cells were prepared in which LRP expression was downregulated using LRP-specific phosphorothioate antisense oligodeoxynucleotides. LRP expression appeared to be unaffected when resting

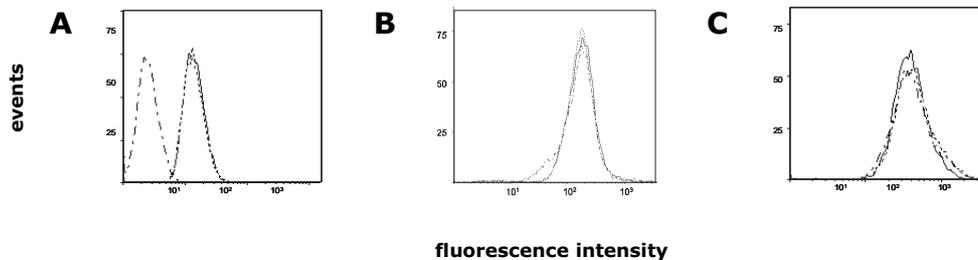


**Figure 4. Surface expression of LRP and  $\alpha_L$ -subunit in monocytes and U937 cells.** Freshly isolated monocytes (A-C), non-transfected U937 cells (D-F), and U937 cells transfected with sense- (G-I) or antisense LRP oligonucleotides (J-L) were collected on object glasses by cytospin centrifugation ( $10^5$  cells/spin). After fixation,  $\alpha_L$ -subunit (A, D, G, J) was visualized using monoclonal antibodies and a TexasRed-labeled secondary antibody. LRP (B, E, H, K) was detected using polyclonal antibodies and a FITC-labeled secondary antibody. Original magnification,  $\times 250$ . C, F, I, L. Enlarged merge images; original magnification,  $\times 750$ . White boxes indicate the positions of the enlarged merge images.

cells were treated with sense oligodeoxynucleotides (Fig 4H). In contrast, a strong downregulation of LRP expression was observed in cells that were transfected with antisense LRP oligodeoxynucleotides (Fig 4K). Downregulation of LRP-surface expression was examined in a quantitative manner using flow cytometric analysis, which revealed that the amount of surface-exposed LRP was similar in non- and sense-transfected U937 cells, but reduced to background levels in antisense-transfected cells (Fig 5A). The expression of the  $\alpha_L$  subunit at the cell surface remained unaffected in sense- and antisense-transfected cells (Fig 4G, J). Indeed, flow cytometric analysis showed similar mean fluorescence intensities for non-, sense-, and antisense-transfected U937 cells (Fig 5B). In addition, the surface expression of uPAR, which has the potential to associate with LRP and  $\beta_2$ -integrins, remained unchanged on transfection of the U937 cells (Fig 5C). These data support the view that the amount of  $\alpha_L$  expression at the cell surface is independent of the presence of LRP.

### Reduced PMA-mediated clustering of $\alpha_L$ subunit in LRP-deficient cells

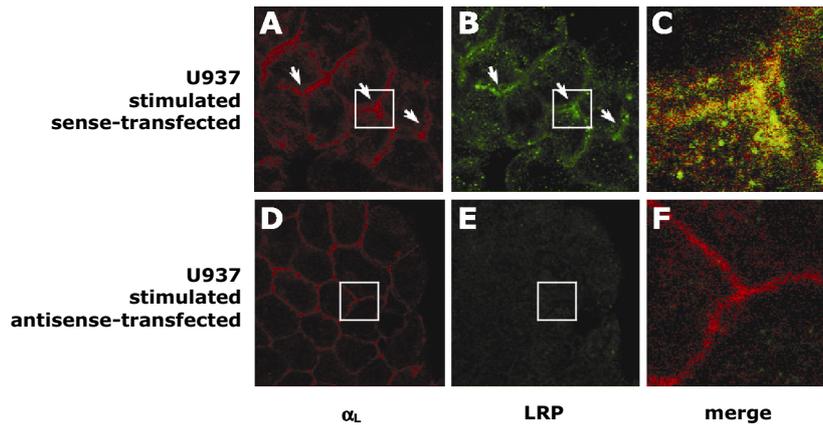
Because PMA-mediated stimulation results in the clustering of  $\alpha_L\beta_2$ -integrins, we further addressed the effect of PMA stimulation on sense and antisense oligodeoxynucleotide-treated U937 cells. PMA stimulation was indeed associated with an increase in the density of the  $\alpha_L$  subunit at the cell surface in sense-transfected cells (Fig 6A). A similar increase in density at indicated regions at the cell surface was observed for LRP (Fig 6B and C), suggesting that the translocation of  $\alpha_L$  and LRP is linked to some extent. With regard to the antisense-transfected cells, no obvious increase in density was observed for the  $\alpha_L$  subunit (Fig 6D), despite normal expression levels for this subunit. This suggests that clustering of  $\beta_2$ -integrins is dependent on the presence of LRP.



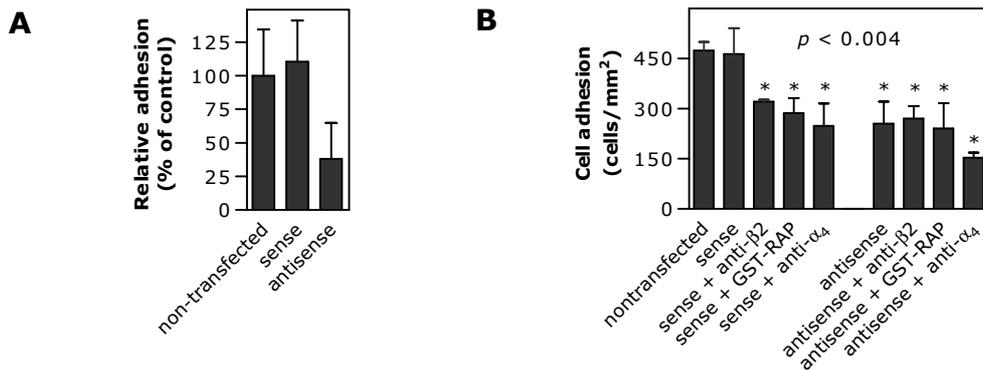
**Figure 5. Cell surface expression of LRP,  $\alpha_L$ , and uPAR.** Flow cytometric analysis of the cell surface expression of LRP (A),  $\alpha_L$  (B), and uPAR (C) in non- (—), sense- (....), and antisense- (-.-.-) transfected U937 cells.

**Reduced LRP levels result in impaired  $\beta$ 2-integrin-dependent adhesion to endothelial cells**

$\beta$ 2-Integrin-dependent adhesion of leukocytes to the endothelium requires clustering of these integrins. Given that our microscopic analysis suggested that clustering of  $\beta$ 2-integrins is LRP dependent, the effect of LRP deficiency on leukocyte adhesion to the endothelial surface was addressed. Therefore, the adhesion of normal U937 cells and U937 cells transfected with sense or antisense LRP oligos to HUVECs was compared. First, in a static adhesion assay using stimulated HUVECs as a surface, sense-transfected U937 cells were similar to non-transfected cells (*Fig 7A*). In contrast, a markedly reduced adhesion was observed for antisense-transfected cells (*Fig 7A*). This was further addressed in perfusion experiments. As shown in *Figure 7B*, U937 cells adhered efficiently to the HUVEC layer ( $474 \pm 25$  cells/mm<sup>2</sup>; n=3). A similar amount of adhesion was detected for sense-transfected cells ( $463 \pm 77$  cells/mm<sup>2</sup>; n=8; P > .05). Monoclonal antibodies directed against the  $\beta$ 2-subunit or the LRP antagonist GST-RAP effectively interfered with the adhesion of sense-transfected cells ( $321 \pm 6$  cells/mm<sup>2</sup>; n=3; P = .0005 and  $287 \pm 44$  cells/mm<sup>2</sup>; n=4; P = .0013, respectively). Thus, the adhesion of U937 cells to endothelial cells is a process that involves  $\beta$ 2-integrins and LRP. With regard to antisense-transfected cells, lesser cells were observed to adhere to the endothelial surface compared with non- and sense-transfected cells ( $255 \pm 66$  cells/mm<sup>2</sup>; n=8; P = .0004 compared with non-transfected cells). Moreover, no further decrease in adhesion was observed in the presence of either anti- $\beta$ 2 antibodies ( $270 \pm 38$  cells/mm<sup>2</sup>; n=3; P = .0015 compared with non-transfected cells and P > .05 relative to antisense-transfected cells) or the LRP antagonist GST-RAP ( $241 \pm 75$  cells/mm<sup>2</sup>; n=4; P = .0039 compared with non-transfected cells and P > .05 relative to antisense-transfected cells). In contrast, an additional decrease in adhesion was observed in the presence of an anti- $\alpha$ <sub>4</sub> antibody ( $153 \pm 15$  cells/mm<sup>2</sup>; n=3; P = .003 compared with antisense-transfected cells). In conclusion, these data indicate that the absence of functional LRP is associated with reduced  $\beta$ 2-integrin-dependent adhesion to the endothelial surface.



**Figure 6. Effect of sense and antisense nucleotides on expression of LRP and  $\alpha_L$ -subunit.** U937 cells were transfected with sense (A-C) or antisense oligonucleotides (D-F) and were stimulated with 100 nM PMA for 15 minutes. Cells were collected on object glasses and stained for  $\alpha_L$  and LRP, as described in the legend to Figure 4. Original magnification,  $\times 250$ . C and F. Enlarged merge images; original magnification,  $\times 1375$ . White boxes indicate the positions of the enlarged merge images.



**Figure 7. Adhesion of U937 cells to HUVECs under flow conditions.** A. Non-, sense-, and antisense-transfected U937 cells were added to wells coated with HUVECs, which were stimulated with TNF- $\alpha$  (100 U/mL) for 4 hours and fixed afterward. Bound U937 cells were analyzed as described in the legend to Figure 1B. B. Non-, sense-, and antisense-transfected U937 cells ( $2 \times 10^6$  cells/mL) were perfused for 10 minutes with a wall shear stress of 0.8 dyne/cm<sup>2</sup> over a glass coverslip confluent with HUVECs, which were stimulated with TNF- $\alpha$  (100 U/mL) for 4 hours before perfusion. Where indicated, cells were preincubated with anti- $\beta_2$ -integrin or anti- $\alpha_4$  antibodies (20  $\mu$ g/mL) or GST-RAP (50  $\mu$ g/mL) for 15 minutes. The number of firmly adhered cells per square millimeter was obtained from video-image analysis. Data represent the mean  $\pm$  SD of 3 to 8 perfusions.

## Discussion

Many receptors have evolved to fulfill one specific function, though a subset of receptors is known to display multispecificity and multifunctionality<sup>45</sup>. One such example is LRP, a member of the LDL-receptor family that, since its first description in 1988, has been classified as an endocytic receptor<sup>1</sup>. However, in the past few years, it has become evident that LRP function encompasses other processes as well. For instance, LRP expressed in brain vascular endothelial cells is involved in the regulation of the vascular tone and permeability of the blood-brain barrier<sup>46</sup>, whereas LRP expressed in primary neurons mediates calcium signaling through N-methyl-D-aspartate receptors<sup>12</sup>. This suggests that the functionality of LRP is dependent on the cell type in which this receptor is expressed.

LRP is abundantly present in leukocytes<sup>15;16</sup>, but its contribution to leukocyte function has remained poorly understood. In the present study, we obtained evidence that LRP is able to bind to leukocyte-specific  $\beta$ 2-integrin complexes: 1) adhesion of the monocytic cell line U937 to recombinant fragments of LRP, i.e. clusters II and IV, could be inhibited by antibodies directed against  $\beta$ 2-,  $\alpha$ <sub>L</sub>-,  $\alpha$ <sub>M</sub>-, or  $\alpha$ <sub>X</sub>-subunits (*Fig 1C*); 2) LRP and  $\beta$ 2-integrins coprecipitated in immunoprecipitation experiments (*Fig 3*); 3) recombinant fragments of the  $\alpha$ <sub>M</sub>- and  $\alpha$ <sub>L</sub>-subunits interacted with full-length LRP (*Fig 2B*); 4) LRP cluster IV displayed binding to purified  $\alpha$ <sub>M</sub> $\beta$ 2-integrin (*Fig 2A*).

The integrin superfamily has been reported to comprise 18 different  $\alpha$ -subunits and 8 different  $\beta$ -subunits, which can combine to make up to 24 different heterodimers. To the best of our knowledge, the  $\beta$ 2-integrin subfamily is the first to be reported to bind LRP. It should be mentioned that LRP has recently been implicated to promote the maturation and intracellular trafficking of  $\beta$ 1-integrins<sup>47</sup>. Furthermore, LRP has been reported to mediate endocytosis of complexes between plasminogen activator inhibitor-1 and the  $\alpha$ <sub>V</sub>-subunit<sup>48</sup>. However, no evidence of direct interaction between these integrin subunits and LRP could be obtained in these studies. Indeed, we could not detect any inhibition of cell adhesion to LRP fragments in the presence of inhibitors of  $\alpha$ <sub>4</sub> $\beta$ 1 or  $\alpha$ <sub>V</sub> $\beta$ 3 (*Fig 1C*).

$\beta$ 2-Integrins and LRP are transmembrane proteins containing cytoplasmic and extracellular domains. Our results obtained from cell adhesion and protein-interaction assays demonstrate that at least some of the interactive sites are located in the cluster II and IV regions of LRP and the I-domain of the  $\alpha$ -subunits (*Fig 1 and 2*), both of which are part of the respective extracellular regions. Cell adhesion was also inhibited in the presence of the anti- $\beta$ 2 antibody. It seems reasonable to assume that this subunit is involved in complex assembly as well. Alternatively, antibody binding to the  $\beta$ 2-subunit may prevent binding to

the complementary subunit by sterical hindrance. With regard to the cytoplasmic regions of the receptors, it is unclear whether they are involved in complex formation. This aspect is currently under investigation.

We considered the possibility that LRP is involved in the removal of  $\beta$ 2-integrins from the cell surface. As such, the amount of  $\beta$ 2-integrins expressed at the cell surface would be increased on the downregulation of LRP. However, examination of  $\alpha_L$  (Fig 5B) or  $\alpha_M$  (data not shown) surface expression using flow cytometric analysis revealed that sense- and antisense-transfected monocytic U937 cells display a fluorescence response similar to that of non-transfected cells. Apparently, the absence of LRP leaves the amount of  $\alpha_L$  or  $\alpha_M$  at the cell surface unaffected, suggesting another function for the interaction between LRP and  $\beta$ 2-integrins. Data obtained from immunoprecipitation experiments (Fig 3A) and confocal microscopy (Fig 4) point to the possibility that  $\beta$ 2-integrins and LRP form a complex at the cell surface. Indeed, we observed not only a cooperative clustering of both receptors (Fig 6), but also an increase of  $\beta$ 2-integrin/LRP complex in detergent-insoluble membrane fractions (Fig 3B). The notion that both receptors have the potential to be present in detergent-resistant membrane fractions is in line with previous reports showing the presence of  $\beta$ 2-integrins and LRP in lipid rafts of leukocytes and smooth muscle cells, respectively<sup>13;44</sup>.

Recently, one of us reported that  $\beta$ 2-integrins associate with matrix metalloproteases<sup>49</sup> and that this interaction is critical for  $\beta$ 2-integrin-dependent leukocyte migration. This indicates that  $\beta$ 2-integrins have the potential to form supramolecular complexes that are of functional importance. We have addressed this possibility for the LRP/ $\beta$ 2-integrin complex by testing  $\beta$ 2-dependent adhesion to stimulated HUVECs. Adhesion of U937 cells to HUVECs was reduced by almost 50% in the presence of anti- $\beta$ 2 subunit antibodies and to the same extent in the presence of GST-RAP (Fig 7A). Moreover, a similar reduction in adhesion was observed for LRP-deficient cells, and this reduced adhesion remained unchanged in the presence of anti- $\beta$ 2-integrin antibodies or GST-RAP. It should be noted that the residual adhesion observed could reflect adhesion mediated by selectins and  $\alpha_4\beta$ 1-integrins. Indeed, the adhesion of LRP-expressing and -deficient cells was significantly decreased in the presence of anti- $\alpha_4$ -antibodies (Fig 7B). Nevertheless, our data demonstrate that the absence or inhibition of LRP abrogates  $\beta$ 2-integrin-dependent cell adhesion. Thus, it seems conceivable that LRP and  $\beta$ 2-integrins form a functionally important complex at the leukocyte surface.

Of importance in this regard is the notion that LRP and  $\beta$ 2-integrins may form complexes with other receptors as well. For instance, LRP has been shown to be involved in the regulation of uPAR surface expression, whereas urokinase plas-

minogen activator receptor (uPAR) contributes to  $\beta$ 2-integrin function. As such, some of our observations could be explained by a model in which the absence of LRP modulates the surface expression of uPAR, which in turn could affect  $\beta$ 2-integrin-dependent adhesion. Thus, LRP and  $\beta$ 2-integrins could be part of a larger complex in which uPAR acts as an intermediate between both receptors. However, several observations argue against such a model. First, it has been established that binding of uPAR to LRP requires the presence of the uPA/PAI-1 complex<sup>50</sup>. In agreement with this notion, we were unable to detect coprecipitation between LRP and uPAR in immunoprecipitation experiments (data not shown). Second, the surface expression of uPAR was unaffected in sense- or antisense-transfected cells (*Fig 5C*). In view of our experimental data, it seems reasonable to assume that complex formation between LRP and  $\beta$ 2-integrins is independent of uPAR.

The  $\beta$ 2-integrins are known to recognize a variety of ligands to facilitate leukocyte adhesion to a broad range of surfaces. The main ligands are the ICAM molecules, but also fibrinogen and the platelet glycoprotein Ib/IX/V receptor may provide adhesive surfaces for these integrins. The interaction with these counter-receptors involves the I-domain of  $\alpha_M$ <sup>34</sup>, the same domain that contains a binding site for LRP. Because  $\beta$ 2-integrin-dependent leukocyte adhesion to these ligands is obviously not hampered by the presence of LRP, the possibility exists that the LRP-binding site is distinct from those of ICAMs, fibrinogen, and glycoprotein Ib. Alternatively, the relatively low affinity for LRP may allow easy dissociation by ICAMs, fibrinogen, or glycoprotein Ib, provided that they display higher affinity for  $\alpha_M$  than LRP. This alternative explanation seems favorable in view of our observation that fibrinogen was found to interfere with the binding of the  $\alpha_M$  I-domain to LRP (*Fig 2B*). Thus, it seems conceivable that LRP is required to position  $\beta$ 2-integrins appropriately to facilitate the interaction with counter-receptors such as fibrinogen. This "cis" interaction, however, does not exclude the possibility that  $\beta$ 2-integrins and LRP may interact in "trans" or, in other words, that  $\beta$ 2-integrins exposed on the surface of one cell interact with LRP on the surface of another cell. This possibility is supported by our experiments that show effective leukocyte adhesion to purified LRP fragments (*Fig 1*).

In conclusion, the present study provides evidence for a previously unrecognized link between LRP and the inflammatory system. Our data point to a model in which LRP is a regulator of  $\beta$ 2-integrin function, because LRP deficiency abrogates  $\beta$ 2-dependent cell adhesion. The possibility that LRP also contributes to other  $\beta$ 2-integrin-dependent functions, such as phagocytosis and migration, remains to be investigated.

## Acknowledgements

This study was supported by a grant from the Dutch Organization of Scientific Research ZonMW (no. 902-26-236) to P.J.L.

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chapter **3**

**Differential localization of LDL receptor-related protein / CD91 in lymphocytic subsets**

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## Abstract

LDL receptor-related protein (LRP), also known as CD91, is a multifunctional protein that is expressed in a broad range of cells, including leukocytes. For instance, we recently demonstrated the requirement of LRP for  $\beta$ 2-integrin function in leukocytes [Spijkers et al, 2005]. Whereas LRP expression and function in monocytes has been well established, little is known about LRP expression in lymphocytes. The present study was aimed to elucidate LRP expression patterns in various lymphocytic subsets. The presence of LRP mRNA and protein could be detected in B-lymphocytes, natural killer (NK) cells and T-lymphocytes as analyzed by RT-PCR, Western blotting, flow cytometry and confocal microscopy. LRP is predominantly present at the cell surface of B-lymphocytes and NK cells, but it was found selectively inside the cell in >95% of the CD4+ and CD8+ T-lymphocytes. This let us to hypothesize that LRP may become redistributed to the cell surface upon T-lymphocyte activation. This was first tested by stimulating T-lymphocytes using interleukin-2 and phytohemagglutinin. Interleukin-2/phytohemagglutinin stimulation resulted in a transient increase of cell-surface-located LRP protein. After 48 h, the number of T-lymphocytes containing LRP at the cellular surface was increased from 2% till 12%, which was followed by a gradual decline to baseline levels in the subsequent 5 days. In a second approach, changes in LRP-surface expression were monitored by mixing peripheral blood mononuclear cells derived from two different donors. This resulted in a near immediate translocation of LRP to the cell-surface in 60-80% of the T-lymphocytes. By using isolated cells, we found that this translocation occurred exclusively in the presence of CD14+ monocytes. The observation that LRP can be rapidly recruited to the cellular surface in T-lymphocytes upon interaction with non-autologous monocytes may point to a role of LRP in the pathophysiology of transfusion-related complications, such as graft-versus-host disease.

## Introduction

The LDL-receptor related protein (LRP), also known as  $\alpha_2$ -macroglobulin receptor or CD91, is a multiligand receptor of the LDL receptor family<sup>1</sup>. It is a heterodimeric protein consisting of an 85-kDa cytoplasmic and transmembrane domain, and a non-covalently linked 515-kDa extracellular domain<sup>2</sup>. The extracellular domain comprises four clusters of complement-type repeats<sup>3;4</sup> that mediate the binding of over 35 structurally and functionally unrelated ligands (for review see Herz and Strickland<sup>5</sup>). The cytoplasmic tail contains motifs for binding of numerous adaptor proteins implicated in endocytosis and signaling pathways<sup>6;7</sup>.

The physiological function of LRP has initially been reported to be related to the endocytic process in order to mediate the cellular uptake of its ligands from the circulation or from the cell membrane<sup>8;9</sup>. LRP is also implicated in various other transport pathways, such as transcytosis across the blood-brain barrier<sup>10;11</sup> and nuclear localization of the growth factor midkine<sup>12</sup>. Furthermore, LRP is involved in cellular signal transduction pathways leading to cell proliferation, migration, neurotransmission and others<sup>13</sup>. One intriguing aspect of LRP is that it exerts functions that are highly cell-type restricted. For example, LRP regulates platelet-derived growth factor-induced proliferation and migration specifically in smooth muscle cells through complex formation with the platelet-derived growth factor receptor<sup>14</sup>. In neurons, LRP associates with N-methyl-D-aspartate receptor, leading to calcium influx<sup>15</sup>. Thus, LRP appears to form heterologous receptor-complexes that control cellular processes.

LRP is expressed in a large spectrum of cell types, including hepatocytes, fibroblasts, adipocytes and astrocytes<sup>16</sup>. Concerning leukocytes, LRP is known to be present in polymorphonuclear cells (PMNs), monocytes and a subset of dendritic cells<sup>16-19</sup>. Several studies point to essential physiological functions of LRP in these leukocytes: 1) the cytoplasmic tail of LRP is an essential element in the phagocytosis in macrophages<sup>20;21</sup>; 2) LRP acts as a cofactor in 12/15 lipoxygenase-dependent LDL oxidation in macrophages<sup>22</sup>; 3) LRP mediates uptake of heat shock protein / peptide complexes in dendritic cells<sup>19</sup>; 4) LRP is required for  $\beta$ 2-integrin-dependent leukocyte adhesion<sup>23</sup>.

In view of its versatility, it is conceivable that LRP also contributes to functions in lymphocytes. However, conflicting information exists with regard to LRP expression in these cells. Moestrup et al were unable to detect LRP in T-lymphocytes when assessed by immunohistochemistry<sup>16</sup>. In contrast, Gläser and coworkers reported the presence of LRP in these cells based on *in situ* hybridization and immunohistochemistry<sup>24</sup>. Apparently, whether or not LRP is present in T-lymphocytes is still inconclusive. With regard to other lymphocytic cells, like B-

lymphocytes and natural killer (NK) cells, no information is available to the best of our knowledge.

Therefore, the aim of the present study was to elucidate the LRP expression pattern in lymphocytes. Our data show that LRP mRNA and protein is present in B-lymphocytes, NK cells and T-lymphocytes. LRP protein is predominantly located at the cell surface of B-lymphocytes and NK cells, whereas it is stored intracellularly in T-lymphocytes. In these cells, near immediate translocation of LRP to the cell surface occurs in a mixed lymphocyte reaction. Our data show that LRP is differentially expressed at the cell membrane of lymphocytes, suggesting distinct functions for LRP in these leukocytes.

## **Experimental Procedures**

### **Materials**

Ficoll-Paque was from Amersham Biosciences (Uppsala, Sweden). MicroBeads-conjugated antibodies and MiniMACS were from Miltenyi Biotec (Bergisch Gladbach, Germany). All antibodies against CD-markers were obtained from BD Biosciences (San Jose, CA), anti-LRP 85-kDa from RDI (Flanders, NJ), fluorescein isothiocyanate (FITC)-conjugated mouse anti-LRP clone  $\alpha 2$ -M-R-I4C2 from Biomac (Leipzig, Germany), peroxidase-conjugated rabbit anti-mouse IgG from DAKO (Glostrup, Denmark). RPMI 1640 and penicillin/streptomycin were from Gibco Life Technologies (Paisley, UK), fetal bovine serum from Cambrex (Verviers, Belgium), interleukin-2 (IL-2) from Chiron (Emeryville, CA) and phytohaemagglutinin (PHA) from Remel (Lenexa, KS).

### **Lymphocyte isolation**

Anticoagulated blood was collected from healthy volunteers. Peripheral blood mononuclear cells (PBMCs) were isolated using Ficoll-Paque density centrifugation as instructed by the manufacturer. The following cells were isolated: total T-lymphocytes (CD3+), T-helper cells (CD3+CD4+), cytotoxic T-lymphocytes (CD3+CD8+), NK cells (CD16+CD56+) and B-lymphocytes (CD19+). CD3+ T-lymphocytes were isolated by magnetic cell sorting using the Pan T-cell isolation kit. Other lymphocyte subsets were incubated with an antibody against a specific CD-marker and subsequently with MicroBeads-conjugated anti-mouse IgG, and isolated using magnetic cell sorting. Peripheral blood monocytes were isolated using CD14-MicroBeads. For all cell isolations, a MiniMACS cell separator was used. Cell purity appeared to be greater than 95% for all isolations, as examined by flow cytometry.

### **Culturing of T-lymphocytes**

PBMCs were isolated using Ficoll-Paque and resuspended at  $1 \times 10^6$  cells/ml in RPMI, 10% FCS, 1% penicillin/streptomycin and 50  $\mu$ M  $\beta$ -mercapto-ethanol. 300 U/ml IL-2 and 1  $\mu$ g/ml PHA were added to the cell suspension and cells were cultured in a humidified CO<sub>2</sub> (5%) incubator at 37°C for 7 days. Cells were kept in culture at a density of  $0.5 \times 10^6$  cells/ml. Activated T-lymphocytes were characterized by their CD25 expression. LRP levels were measured at day 0, 1, 2, 4 and 7.

### **RT-PCR analysis**

Total RNA was purified from isolated leukocytes by means of the RNeasy Mini kit (Qiagen, Hilden, Germany) according to the manufacturer's protocol. After a cDNA synthesis step, a PCR was performed using Amplitaq DNA polymerase (Roche, Basel, Switzerland) and the following primers: forward primer 5' GAACGAGCCAGTAGACCG 3' and reverse primer 5' CAATCTTGCTGTCGACGA 3'. The forward primer spans exon 5 nucleotide 125 to exon 6 nucleotide 13; the reverse primers spans exon 8 nucleotide 78-85. 30 cycles of cDNA amplification were performed with annealing temperature of 62°C. This reaction resulted in a specific LRP product of 527 bp. To confirm the identity of the product, the cDNA product was cloned into a TOPO vector (Invitrogen, Carlsbad, CA) and subsequently sequenced, or digested using BglI. This resulted in cleavage of the 527 bp band into two bands of 305 and 222 bp.

### **Confocal scanning laser microscopy**

Purified lymphocytes were collected on glass coverslips by cytospin centrifugation ( $10^5$  cells/spin). Cells were fixed by 4% paraformaldehyde (PFA) in PBS. Cells were washed extensively and free aldehyde groups were quenched using 50 mM NH<sub>4</sub>Cl in PBS. Cells were optionally permeabilized with 0.1% saponin. Subsequently, cells were stained with FITC-conjugated mouse anti-LRP (Biomac) and mounted in mowiol containing 2.5% 1,4-diazabicyclo[2.2.2]octane. Cells were visualized using a Leica confocal scanning laser microscope and a TCS 4D system.

### **Flow cytometric analysis**

PBMCs were stained with FITC-conjugated anti-LRP and antibodies against different cell markers. For total cell staining, cells were first fixed in 4% PFA and permeabilized using 0.1% saponin. Lymphocytes were identified in the scatter plot by their CD-marker staining and analyzed for their LRP expression at the cell surface and/or intracellular.

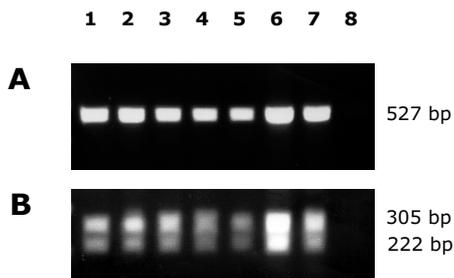
### Mixed lymphocyte reaction

Either PBMCs or isolated T-lymphocytes and monocytes from 2 different donors were mixed during 30 minutes ( $0.5 \times 10^6$  cells from donor 1 +  $0.5 \times 10^6$  cells from donor 2) and fixed with 4% PFA. Cells were stained for their specific CD-marker and LRP and subsequently analyzed by flow cytometry. Using isolated cells, cells were selected by their CD-marker and passed twice over MS columns to ensure ultrapure populations.

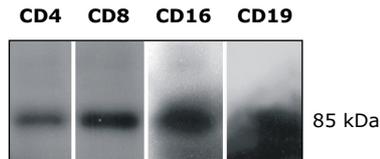
## Results

### LRP mRNA is present in lymphocytic subsets

In order to investigate LRP expression by lymphocytes, various lymphocytic subsets were analyzed for the presence of LRP mRNA and protein. First, RNA was purified from isolated CD4+ T-helper cells, CD8+ cytotoxic T-lymphocytes, CD16+CD56+ NK cells and CD19+ B-lymphocytes. Primers specifically detecting LRP mRNA were subsequently used in an RT-PCR reaction to generate a 527 bp product. This LRP-specific product could indeed be obtained from RNA isolated from the monocytic cell line THP-1 and freshly isolated monocytes (*Fig 1A*). Analysis of mRNA content in CD3+ lymphocytes showed an identical 527 bp band. Further purification of the CD3+ T-lymphocytes obtaining CD4+ and CD8+ cells revealed that both T-lymphocytic subsets express LRP mRNA. Two other lymphocytic cell types, B-lymphocytes (CD19+) and NK cells (CD16+CD56+) also appeared to contain LRP mRNA. To confirm the identity of the 527 bp band, restriction site analysis using BglI results in specific digestion of the 527 bp into 2 fragments of 305 and 222 bp (*Fig 1B*). Also, sequence analysis showed that the obtained PCR products encoded for LRP cDNA (data not shown). Thus, LRP mRNA seems to be produced in each of these lymphocyte subsets.



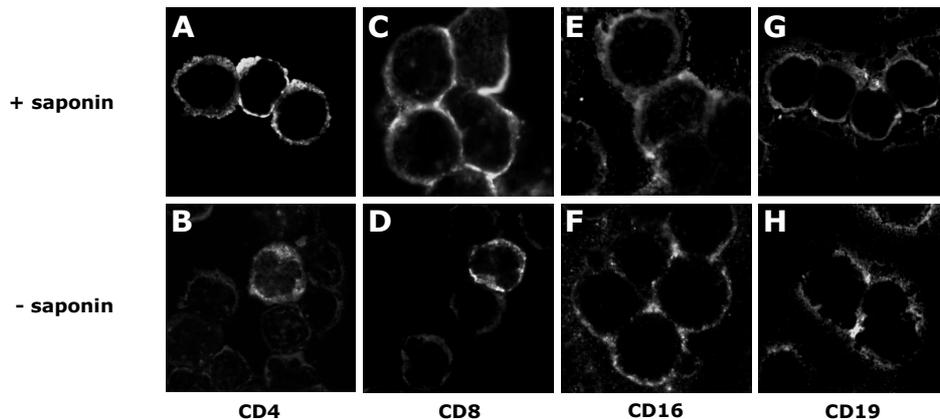
**Figure 1. LRP mRNA expression in lymphocytes.** A. Lymphocytes were isolated from the PBMCs fraction after Ficoll-Paque density centrifugation. After total RNA isolation, an RT-PCR was performed using equal amounts of RNA and specific primers for LRP. PCR products were visualized on an ethidium-bromide stained agarose gel. The length of the expected LRP fragment was 527 bp. B. The obtained PCR product was digested using BglI, resulting in cleavage of the 527 bp product in a 305 bp and a 222 bp fragment. 1: CD3, 2: CD4, 3: CD8, 4: CD16, 5: CD19, 6: CD14, 7: THP-1, 8: no template.



**Figure 2. Western blot analysis of LRP expression.** Lymphocytes were isolated and lysed in 1% NP-40 and 0.5% DOC in PBS. Samples were resolved on SDS-PAGE and subsequently transferred onto a PVDF membrane. Blots were incubated with anti-LRP 85-kDa (RDI, 1:50) and peroxidase-conjugated rabbit anti-mouse IgG (DAKO, 1:1000).

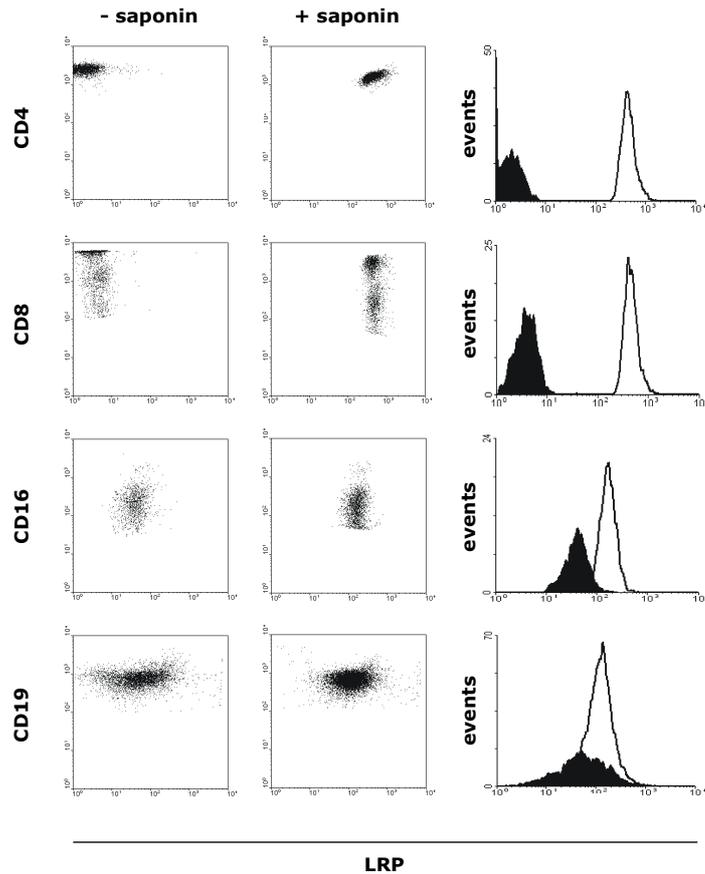
### LRP protein is differentially expressed at the cell surface lymphocytes

To test whether these cells also contain LRP protein, two experimental approaches were employed to detect LRP protein in lymphocytes: 1) Western blotting using an antibody directed against the 85-kDa intracellular tail of LRP, and 2) confocal microscopy using an antibody against the extracellular domain. First, total cell lysates of isolated lymphocytes were subjected to Western blot analysis. The presence of the 85-kDa band could be detected in both T-lymphocytic subsets (*Fig 2*). In addition, LRP protein was also detected in lysates of B-lymphocytes and NK cells. In a second approach, purified lymphocytes were collected on glass coverslips for confocal microscopy studies, permeabilized with saponin, and subsequently stained using a FITC-conjugated antibody against the extracellular domain of LRP (*Fig 3*). T-lymphocytes (CD4+ (*Fig 3A*) and CD8+ (*Fig 3C*)), NK cells (*Fig 3E*) and B-lymphocytes (*Fig 3G*) were all positive for LRP. Thus, both experimental approaches point to the presence of LRP protein in these lymphocytes.



**Figure 3. Localization of LRP in lymphocytes.** PBMCs were isolated using Ficoll-Paque density centrifugation and collected on glass coverslips ( $10^5$  cells/spin). After fixation, cells were treated with (A, C, E, G) or without saponin (B, D, F, H). LRP was stained with a FITC-conjugated antibody and cells were visualized by confocal microscopy. A, B. CD4+ T-helper cells. C, D. CD8+ cytotoxic T-lymphocytes. E, F. CD16+ NK cells. G, H. CD19+ B-lymphocytes.

To distinguish between LRP located at the cell surface or intracellularly, LRP staining was also examined in non-permeabilized lymphocytes. LRP staining was detected in NK cells (*Fig 3F*) and B-lymphocytes (*Fig 3H*). Interestingly, most CD4+ (*Fig 3B*) and CD8+ cells (*Fig 3D*) were negative for LRP cell surface staining. Apparently, CD3+ T-lymphocytes differ from NK and B-lymphocytes in that LRP seems to be maintained within the cell.



**Figure 4. Quantification of intracellular and cell surface located LRP in lymphocytes.** PMBCs were isolated using Ficoll-Paque density centrifugation. Cells were fixed and treated with 0.1% saponin. Cells were stained for LRP and CD4, CD8, CD16 or CD19, respectively and analyzed by FACS measurements. Right panels: histogram of LRP levels in lymphocytes. Black: without saponin, white: with saponin.

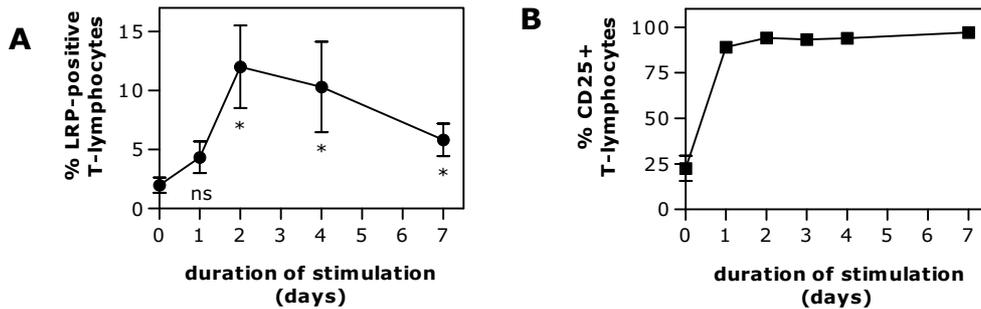
**LRP is present in intracellular compartments in T-lymphocytes**

The apparent dissimilar expression of LRP in NK cells and B-lymphocytes versus T-lymphocytes was further addressed in flow cytometric experiments. To quantify the ratio of intracellularly and cell-surface distributed LRP, permeabilized and non-permeabilized lymphocytes were compared for LRP staining in FACS experiments (Fig 4). In the non-permeabilized condition, high numbers of NK cells and B-lymphocytes expressed LRP at the cell surface ( $91.4 \pm 4.4\%$ , and  $93.0 \pm 3.9\%$ , respectively). Saponin treatment increased the percentage of LRP positive cells up to 100% and the MFI was 2-3-fold higher than in non-permeabilized cells. Thus, these results confirm that LRP is located at the cell surface of most NK cells and B-lymphocytes. In contrast, LRP cell-surface expression was as low as background levels in both CD4+ and CD8+ T-lymphocytes, and only a small number of T-lymphocytes were LRP positive (< 5% of the CD4+ and CD8+ population). To exclude the possibility that the epitope for the antibody was not available at the cell surface of T-lymphocytes, also 3 other monoclonal and polyclonal antibodies against the extracellular domain of LRP were tested. However, none of these antibodies were able to detect LRP on non-permeabilized T-lymphocytes (data not shown), supporting the view that LRP is absent from the T-lymphocytic cell surface.

Under permeabilized conditions the number of LRP-positive T-lymphocytes increased from 2% to 100% for both CD4+ and CD8+ T-lymphocytes. The intensity of staining increased over 50-fold upon permeabilization for both cell types. CD4 and CD8 staining remained unaffected upon permeabilization, indicating that saponin treatment is not associated with non-specific binding of antibodies (data not shown). In conclusion, our data indicate that LRP is located at the cell surface of virtually all B-lymphocytes and NK cells, whereas LRP is mainly distributed intracellularly in T-lymphocytes.

**Activation of T-lymphocytes in a mixed lymphocyte reaction results in redistribution of LRP**

We hypothesized that LRP may become exposed at the cell surface during activation of circulating T-lymphocytes upon an immune response. Therefore, two activation methods were applied to investigate their effect on LRP cell surface expression. First, activation of the immune system may lead to proliferation of naïve T-lymphocytes, which can be mimicked *in vitro* by IL-2 and PHA stimulation. Indeed, a marker for activation (*ie* CD25) increased after 1 day and sustained during 7 day follow-up (Fig 5B). LRP was present at the cell surface of  $2.0 \pm 0.7\%$  of the T-lymphocytes prior to activation (Fig 5A). Stimulation caused a 6-fold increase in the number of LRP positive T-lymphocytes, reaching its maxi-



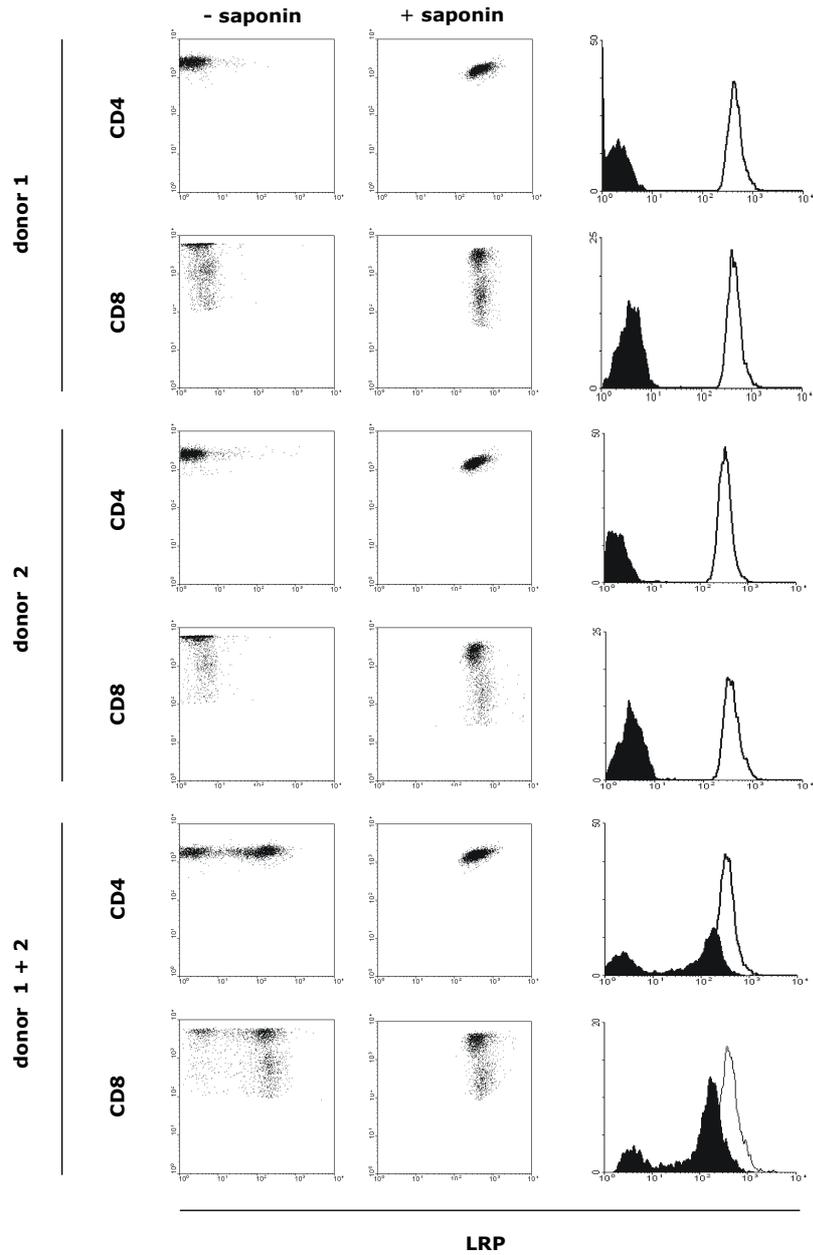
**Figure 5. Stimulation of T-lymphocytes with IL-2 and PHA.** A. T-lymphocytes were cultured in medium containing IL-2 and PHA during 7 days. At day 0, 1, 2, 4 and 7, LRP expression was measured in activated T-lymphocytes by FACS analysis. B. Activated cells were determined by their CD3 and CD25 cell surface expression. Data represent mean  $\pm$  SEM. ns = not significant, \* =  $P < 0.02$ , as compared to  $t=0$ .

num after 2 days ( $12.0 \pm 3.5\%$  of the total vital T-lymphocytes). After 7 days, the number of LRP expressing cells gradually declined to  $5.8 \pm 1.4\%$ .

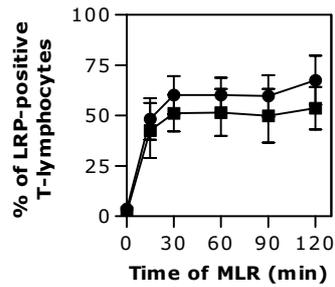
In a second approach, PBMCs from one donor were mixed with PBMCs from a second donor. Only a few non-mixed T-lymphocytes from various donors were positive for LRP at the cell surface before mixing ( $3.2 \pm 1.1\%$  and  $5.5 \pm 2.9\%$  of CD4+ and CD8+ T-lymphocytes, respectively) (Fig 6). Remarkably, combining PBMCs from 2 donors caused cell surface LRP expression in  $68.5 \pm 6.6\%$  and  $79.9 \pm 1.9\%$  of the CD4+ and CD8+ T-lymphocytes, respectively. Compared to non-mixed T-lymphocytes, cell surface LRP levels were increased up to  $40.0 \pm 18.6$  and  $26.6 \pm 14.1$  fold in the combined CD4+ and CD8+ T-lymphocytes, respectively (Fig 6). The number of T-lymphocytes positive for LRP at their cellular surface increased in time, reaching maximum and steady-state levels after 30 minutes (Fig 7). Under permeabilizing conditions, similar total LRP levels were observed for the non-mixed and mixed T-lymphocytes, indicating that appearance of LRP is independent of *de novo* synthesis. Cell surface and intracellular staining using a non-relevant antibody was not increased upon permeabilization or activation (data not shown). Thus, in T-lymphocytes intracellularly positioned LRP may be translocated to the cell surface in an MLR.

### Monocytes are required for the induction of LRP cell surface expression in T-lymphocytes

Next, we investigated which cells were responsible for the upregulation of cell surface LRP in an MLR. To this end, monocytes, CD4+ and CD8+ T-lymphocytes were purified from various donors. Cells were mixed in different combinations for 30 minutes and subsequently fixed (Fig 8). In the resting condition, only a

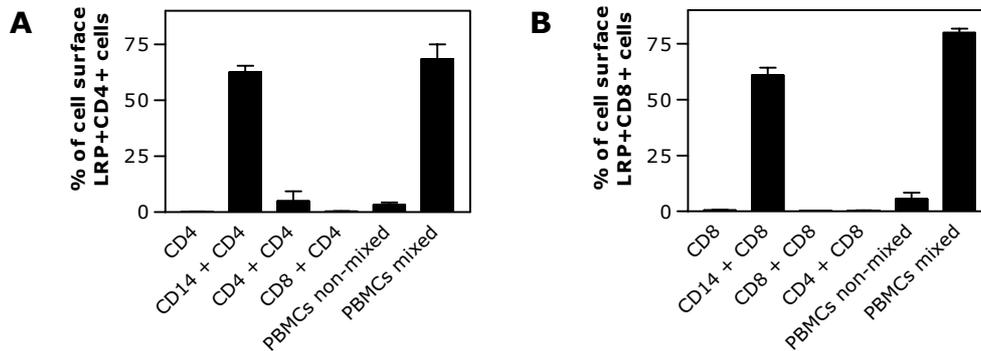


**Figure 6. LRP expression in T-lymphocytes in a mixed lymphocyte reaction.**  $0.5 \times 10^6$  cells from donor 1 were mixed with  $0.5 \times 10^6$  cells from donor 2 for 30 min. Cells were subsequently fixed and optionally permeabilized with 0.1% saponin. Cells were stained for LRP and CD4 or CD8 and analyzed by flow cytometry. Right panels: histogram of LRP levels in lymphocytes. Black: without saponin, white: with saponin.



**Figure 7. Time-dependent upregulation of LRP cell surface levels in an MLR.**  $0.5 \times 10^6$  cells from donor 1 were mixed with  $0.5 \times 10^6$  cells from donor 2 for indicated time. Cells were subsequently fixed, stained for LRP and CD4 (squares) or CD8 (circles) and analyzed by flow cytometry.

few T-lymphocytes showed LRP at the cellular surface, and mixing T-lymphocytes from one donor with T-lymphocytes from a second donor did not increase these cell surface levels (*Fig 8*). Interestingly, monocytes from one donor with CD4+ or CD8+ T-lymphocytes from a second donor resulted in LRP cell surface expression of  $62.6 \pm 2.8\%$  and  $60.9 \pm 3.4\%$  of the CD4+ and CD8+ T-lymphocytes, respectively. Thus, these results indicate that the interaction between monocytes as antigen presenting cells from one donor and T-lymphocytes from a second donor is responsible for the observed upregulation of cell-surface-exposed LRP in T-lymphocytes.



**Figure 8. LRP expression in T-lymphocytes in a mixed lymphocyte reaction using purified cells.** Monocytes (CD14+), CD4+ and CD8+ T-lymphocytes were purified from 2 donors.  $0.5 \times 10^6$  cells from donor 1 were mixed with  $0.5 \times 10^6$  cells from donor 2 for 30 min and subsequently fixed. Cells were stained for LRP and CD4 or CD8 and analyzed by flow cytometry. The x-axis represents the investigated combinations of cells from the 2 donors. The y-axis represents the percentage of CD4+ (A) or CD8+ T-lymphocytes (B) that express LRP at their cell surface. Data are represented as mean  $\pm$  SEM.

## Discussion

The multifunctional receptor LRP is expressed in a wide spectrum of cell types, including adipocytes, neurons, hepatocytes and smooth muscle cells<sup>16</sup>. Whereas initial studies have identified LRP as an endocytic receptor<sup>8;9</sup>, more recent studies have revealed that LRP is involved in other cellular processes as well, like transcytosis and cellular signaling<sup>10;11;13</sup>. However, these latter functions seem to be dependent on the cell-type in which LRP is expressed, suggesting that cell-specific components regulate the functionality of LRP. This is for instance exemplified in monocytes and granulocytes, for which we recently found that LRP forms a functional active complex with  $\beta$ 2-integrins<sup>23</sup> that are selectively expressed in leukocytes. Thus, LRP may contribute to leukocyte function by interacting with leukocyte-specific components.

Although the expression pattern of LRP in myeloid cells has been well established<sup>16;21;23;25;26</sup>, data regarding LRP expression in lymphoid cells have been inconclusive. In the present study we have clearly demonstrated the presence of LRP mRNA and protein in various lymphocytic cells: B-lymphocytes, NK cells and CD4+ and CD8+ T-lymphocytes (*Fig 1-4*). Surprisingly, B-lymphocytes and NK cells differ from both types of T-lymphocytes in that the former two cell types expressed LRP abundantly at the cellular surface, whereas T-lymphocytes were characterized by a virtual absence of LRP at their surface (*Fig 3 and 4*). Confocal and flow cytometric analysis demonstrated that only a small number of individual cells were positive for LRP surface expression (*Fig 3 and 4*). Dual staining using anti-CD4 or anti-CD8 antibodies confirmed the identity of these cells, excluding the possibility that this small number of cells represent contamination of other LRP positive leukocytes.

Despite the lack of cell surface LRP in these cells, various experimental approaches to detect total LRP protein (confocal and flow cytometric analysis in the presence of saponin, and western blotting using total cell lysates) demonstrated that LRP is expressed by T-lymphocytes. In addition, expression levels roughly equal the levels in B-lymphocytes and NK cells. Thus, LRP seems to be maintained inside T-lymphocytes, possibly within storage pools. It is of interest to consider in this respect that B-lymphocytes, NK cells and T-lymphocytes derive from the same lymphoid precursor cell. The B-lymphocyte pathway branches from the T-lymphocyte/NK cell pathway, the latter two cell types sharing a bipotential precursor. During further maturation this precursor cell diverges in either T-lymphocytes or NK cells<sup>27-30</sup>. At some point during this differentiation process, regulatory mechanisms should occur that determine the cellular location of LRP, either by forcing surface expression in B-lymphocytes and NK cells, or by keeping LRP intracellularly in T-lymphocytes. So far, no other LRP-expressing cells

have been identified that are able to maintain the complete LRP population intracellularly. Therefore, we feel the second possibility to be favorable.

It should be noted that based on the MFI values, part of LRP (~ 50-60%) is present intracellularly in B-lymphocytes and NK cells. Such a distribution is also reported for other cells. For instance, LRP is present at the cellular surface of adipocytes. However, insulin treatment transiently increased uptake of LRP ligands  $\alpha_2$ -macroglobulin and cholesteryl esters 2-3 fold within minutes, suggesting translocation of intracellularly stored LRP to the cell surface<sup>31</sup>. A similar upregulation of LRP surface expression was noticed in nerve growth factor-stimulated neurons<sup>32</sup>. Of importance, LRP is already present at the surface in both resting adipocytes and neurons, whereas this is untrue for LRP in T-lymphocytes. Such a dramatic increase in LRP cell surface expression as observed in T-lymphocytes, has not been detected in other stimulated cells.

The notion that LRP is maintained within T-lymphocytes prompted us to explore conditions that are associated with the translocation of LRP from the inside to the surface of these cells. In response to IL-2 and PHA, a transient raise in LRP surface expression was observed in 12% of the T-lymphocytes after 2 days (*Fig 5*). The observation that increased levels of LRP develop during T-lymphocyte activation is in line with a previous report by Banerjee et al<sup>33</sup>, who described that differentiation of naïve T-lymphocytes into specific T-helper subsets parallels increased binding (approximately 15 fold) of the LRP ligand heat shock protein 96.

A more pronounced effect on the presence of LRP at the T-lymphocyte cell surface was observed upon the incubation of these cells with allogenic antigen-presenting cells (APCs) (*Fig 6-8*). In these MLRs, LRP cell surface expression is rapidly upregulated, suggesting the existence of a readily available storage pool. Indeed, our confocal microscopy studies using permeabilizing conditions demonstrated that LRP is located intracellularly just beneath the cell surface, ready for redistribution upon a trigger such as allogenic antigen recognition.

Activation of T-lymphocytes by allogenic APCs is a serious problem in graft-versus-host disease (GVHD)<sup>34</sup>. Many receptors on T-lymphocytes and their counterreceptors on APCs are involved in this process. A major group are the B7/CD28 family of receptors and their counterreceptors<sup>35-37</sup>. Recognition of foreign MHC/peptide complexes by T-lymphocytes induces several stimulatory and inhibitory pathways. For example, the CD28/B7 interactions lead to activation of the T-lymphocyte, whereas CTLA-4 on the T-lymphocyte turns activation down. This latter protein may be of special relevance, since it is also stored intracellularly in naïve T-lymphocytes and becomes exposed at the cell surface upon activation<sup>38</sup>.

Although from our data it is tempting to speculate that LRP is involved in the complex T-lymphocyte response, further studies should be performed concerning the role of LRP in GVHD. Recently, we showed that LRP is required for proper function of  $\beta$ 2-integrins at the cell surface of leukocytes<sup>23</sup>.  $\beta$ 2-Integrins are another family of receptors that support interaction between a T-lymphocyte and an APC and evoke stimulatory signals towards the T-lymphocyte<sup>39</sup>. Therefore, it would be of interest to examine a possible role for LRP in modifying this interaction by  $\beta$ 2-integrins.

In conclusion, our study focused on the elucidation of LRP expression patterns in lymphocyte subsets. Whereas NK cells, B-lymphocytes and T-lymphocytes all express LRP mRNA and protein, they show a distinct LRP protein localization. LRP is mainly present at the cell surface of NK cells and B-lymphocytes and intracellularly in T-lymphocytes. In T-lymphocytes, LRP cell surface expression is rapidly upregulated in an experimental set-up in which T-lymphocytes were incubated with allogenic monocytes. This suggests that LRP may play a role in GVHD. Future experiments must obtain more evidence on the precise function of LRP in this disease.

## Acknowledgements

We gratefully acknowledge T. Mutis, A. Martens, S. Ebeling and F. Miedema (University Medical Center Utrecht, Utrecht, The Netherlands) for helpful discussions. This study was supported by a grant from the Dutch Organization of Scientific Research ZonMW (no. 902-26-236) to P.J.L.

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# chapter 4

## LRP / amyloid $\beta$ -peptide interaction mediates differential brain efflux of A $\beta$ isoforms

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*Neuron*, 2004, 43(3):333-344

## Abstract

LRP (low-density lipoprotein receptor-related protein) is linked to Alzheimer's disease (AD). Here, we report that amyloid  $\beta$ -peptide, A $\beta$ 40, binds to immobilized LRP clusters II and IV with high affinity ( $K_d = 0.6-1.2$  nM) compared to A $\beta$ 42 and mutant A $\beta$ , and LRP-mediated A $\beta$  brain capillary binding, endocytosis and transcytosis across the mouse blood-brain barrier are substantially reduced by the high  $\beta$ -sheet content in A $\beta$  and deletion of the receptor-associated protein gene. Despite low A $\beta$  production in the brain, transgenic mice expressing low LRP-clearance mutant A $\beta$  develop robust A $\beta$  cerebral accumulations much earlier than Tg-2576 A $\beta$ -overproducing mice. While A $\beta$  does not affect LRP internalization and synthesis, it promotes proteasome-dependent LRP degradation in endothelium at concentrations  $> 1$   $\mu$ M consistent with reduced brain capillary LRP levels in A $\beta$ -accumulating transgenic mice, AD and patients with cerebrovascular  $\beta$ -amyloidosis. Thus, low affinity LRP/A $\beta$  interaction and/or A $\beta$ -induced LRP loss at the BBB mediate brain accumulation of neurotoxic A $\beta$ .

## Introduction

The low density lipoprotein receptor-related protein (LRP), a member of the low-density lipoprotein (LDL) receptor family, is a multiligand receptor whose physiological functions are carried out by endocytosis of ligands and activation of multiple signal transduction pathways<sup>1;2</sup>. The precursor form of LRP, a 600-kDa transmembrane glycoprotein, is cleaved in *trans*-Golgi network by furin to generate a 515-kDa  $\alpha$ -subunit and an 85-kDa  $\beta$ -subunit that remain non-covalently associated during LRP transport to the cell membrane<sup>3</sup>. A 39-kDa receptor-associated protein (RAP) is a specialized chaperone molecule which binds to LRP and regulates its proper folding<sup>4</sup>. LRP interacts with a broad range of secreted proteins and resident cell surface molecules in the brain (> 30 structurally diverse ligands) mediating their endocytosis and/or activating signaling pathways through multiple cytosolic adaptor and scaffold proteins<sup>5</sup>. Phosphorylation of LRP's tail regulates ligand internalization and signal transduction<sup>6;7</sup>.

LRP is linked to Alzheimer's disease (AD) by genetic and biochemical evidence. Genetic polymorphism in *LRP* gene is associated with late-onset AD<sup>8-11</sup>. In addition, LRP binds amyloid  $\beta$ -peptide (A $\beta$ ) precursor protein (APP), Apolipoprotein E (ApoE) and  $\alpha_2$ -macroglobulin ( $\alpha_2$ M), which have been genetically linked to AD<sup>1</sup>. The exact pathogenic mechanism(s) by which LRP contributes to neurotoxic A $\beta$  accumulations is unclear. LRP binds secreted APP and influences its degradation<sup>12</sup> and processing<sup>13</sup> leading to increased A $\beta$  production<sup>14</sup>. It may be involved in  $\alpha_2$ M-mediated slow reuptake of secreted A $\beta$  by neurons derived from transgenic APP mice and in mouse fibroblasts with no degradation of A $\beta$ <sup>15;16</sup>. Slow degradation of A $\beta$  via  $\alpha_2$ M in mouse fibroblasts and U97 human glioma cells has also been suggested<sup>17</sup>. ApoE3 may prevent A $\beta$  neurotoxicity by forming complexes with A $\beta$ , and its protective effect possibly requires LRP<sup>18</sup>, although the role of other lipoprotein receptors, e.g., LDLR, megalin, cannot be ruled out. However, overexpression of functional LRP mini-receptors in neurons of Alzheimer's PDAPP mice results in age-dependent increases of soluble A $\beta$  in the brain<sup>19</sup>, suggesting LRP on neurons *in vivo* does not mediate A $\beta$  clearance.

LRP is expressed in brain capillary endothelium<sup>20;21</sup>. The idea that LRP along the brain capillary membranes clears A $\beta$  *in vivo* has been supported by brain efflux studies with exogenous soluble A $\beta$ 40 suggesting the blood-brain barrier (BBB) is a primary clearance site for A $\beta$ <sup>20</sup>. However, neither this nor other studies have clarified at the molecular and cellular level how LRP at the BBB works as a cargo/clearance receptor for A $\beta$ , and whether LRP on brain capillaries clears neurotoxic A $\beta$ 42<sup>22</sup> and vasculotropic A $\beta$  mutants<sup>23</sup>. Reduced levels of LRP in AD have been associated with A $\beta$  brain accumulations<sup>8;16;20</sup>, but whether A $\beta$  directly

interferes with LRP's internalization, degradation and/or synthesis at the clearance site(s) in brain, is not known.

Here, we provide evidence for a direct LRP/A $\beta$  protein-protein interaction and demonstrate this interaction on brain capillaries regulates in an isoform-specific manner binding of A $\beta$  to LRP followed by A $\beta$  endocytosis and transcytosis across the BBB. We report LRP at the BBB favors clearance of A $\beta$ 40 relative to a high  $\beta$ -sheet content A $\beta$ 42 and vasculotropic mutant A $\beta$ , while deletion of the RAP gene/depletion of LRP<sup>24</sup>, but not of the LDLR and VLDLR genes, precludes rapid A $\beta$  brain capillary clearance and efflux at the BBB. Transgenic mice producing low LRP-clearance mutant A $\beta$  in the brain develop robust A $\beta$  accumulations compared to Tg-2576 wild-type A $\beta$ -overproducing mice<sup>25</sup>, despite > 20-fold lower brain APP expression and > 30-fold reduced A $\beta$  neuronal production. Finally, we show A $\beta$  does not affect LRP internalization or synthesis, but promotes proteasome-dependent LRP degradation at pathological levels consistent with low LRP activity in brain microvessels in A $\beta$ -accumulating mice and in patients with AD and familial cerebrovascular  $\beta$ -amyloidosis.

## Experimental Procedures

### Reagents

Wild-type and mutant A $\beta$  (Dutch40, Dutch42, Dutch/Iowa40) peptides were synthesized by solid-phase F-moc (9-fluorenylmethoxycarbonyl) amino acid chemistry, purified by reverse phase-HPLC and structurally characterized, as described<sup>26,27</sup>. Recombinant LRP fragments encompassing clusters II and IV were produced using stable transfected baby hamster kidney cell lines<sup>28</sup>. We used human recombinant RAP (EMD Biosciences, Inc. San Diego, CA), polyclonal goat anti-human LRP N20 antibody which cross reacts with mouse LRP (1:200, Santa Cruz Biotech. Inc., Santa Cruz, CA), monoclonal mouse antibody against C-terminal domain of human LRP  $\beta$ -chain which cross reacts with mouse LRP (5A6, 1:350, 5  $\mu$ g/ml; EMD Biosciences, Inc. San Diego, CA), monoclonal mouse antibody against human LRP  $\alpha$ -chain (8G1, 1:240, 5  $\mu$ g/ml; EMD Biosciences, Inc. San Diego, CA), monoclonal mouse antibody P2-1 specific for human APP (1:1000, 1 mg/ml), 22C11 which recognizes mouse and human APP (1:100, 0.5 mg/ml; Chemicon International, Temecula, CA), 66.1 to residues 1-8 of human A $\beta$  (1:1000, 1 mg/ml)<sup>29</sup>, rat anti-mouse CD31 antibody (1:200, BD Pharmigen, Lexington, KY), polyclonal rabbit antibody to human Von Willebrand Factor (VWF) (1:200, DAKO, Glostrup, Denmark), RAGE-specific IgG (1:500, 1 mg/ml)<sup>29</sup>, goat anti-human VLDLR (1:50, 200  $\mu$ g/ml, Santa Cruz Biotech Inc.,

Santa Cruz, CA) and monoclonal mouse human LDLR-specific IgG C7 (1:50, 200  $\mu$ g/ml, Research Diagnostics, Flanders, NJ).

### Surface plasmon resonance analysis

LRP clusters II and IV were immobilized at CM5 chips at a density of 10-20 fmol/mm<sup>2</sup> and incubated with A $\beta$ 40, A $\beta$ 42, and mutant A $\beta$  (Dutch/Iowa40) (0-50 nM) in 150 mM NaCl, 0.005% (v/v) Tween-20 and 25 mM Hepes (pH 7.4) at a flow rate of 5  $\mu$ l/min for 2 min at 25°C, as described<sup>28</sup>. RAP was used at 500 nM. The data were analyzed to calculate apparent association rate constants  $K_{on,app}$  and apparent dissociation rate constants  $K_{off,app}$  using a single-site binding model<sup>28</sup>. Apparent affinity constants  $K_{d,app}$  were inferred from the ratio  $K_{off,app}/K_{on,app}$ . Data are based on 3-5 measurements using 6-9 different concentrations for each measurement and presented as the average ( $\pm$  SEM). Analysis was performed using Biacore X biosensor system (Uppsala, Sweden) and BIA evaluation 3.0 software (Biacore, Sweden).

### Secondary structure analysis

Secondary structure of peptides was analyzed by circular dichroism<sup>30;31</sup>. Briefly, 20-25  $\mu$ g of hexafluoroisopropanol treated seedless peptide was dissolved in 980  $\mu$ L of 10 mM phosphate buffer, pH 7.4, and centrifuged to remove precipitated/undissolved material. The CD spectrum was recorded within 24 h using a 1 mm path length cell, on an Aviv 202 CD spectropolarimeter (Proterion, Piscataway, NJ). Results are expressed as molar ellipticity and the percentage of  $\alpha$ -helix,  $\beta$ -sheet,  $\beta$ -turn and random coil determined.

### Radio-iodination of A $\beta$

Radioiodination of A $\beta$  was carried out by mild "lactoperoxidase" method (Thorell and Johansson, 1971). Typically, 10  $\mu$ g of A $\beta$  was labeled for 18 min at room temperature with 2 mCi of Na[<sup>125</sup>I]. After radiolabeling, the preparations were subjected to reverse-phase HPLC separation using a Vydac C4 column and a 30 minute linear gradient of 25% to 40% acetonitrile in 0.059% trifluoroacetic acid to separate the monoiodinated non-oxidized form of A $\beta$  (which is the tracer we are using) from diiodinated A $\beta$ , non-labeled non-oxidized A $\beta$ , and oxidized A $\beta$  species, as we reported<sup>29</sup>. The content of material in the peaks eluted from HPLC was determined by MALDI-TOF mass-spectrometry to ensure the purity of the radiolabeled species. For MALDI-TOF mass spectrometry A $\beta$  peptides were labeled under identical conditions using Na[<sup>127</sup>I] instead of the radioactive nuclide. The specific activities were in the range of 45 to 65  $\mu$ Ci/ $\mu$ g of peptide. For brain capillary uptake studies and animal clearance studies, we used in most experiments the preparations within 24 h of labeling that were  $\geq$  99% TCA-

precipitable. If used within 72 h of labeling, the radiolabeled peptides were stabilized in ethanol as a quenching agent. The HPLC/SDS-PAGE analysis was used to confirm the monomeric state of infused radiolabeled A $\beta$ .

#### **Brain capillary binding, endocytosis and uptake of $^{125}\text{I}$ -A $\beta$**

Brain capillaries from wild type, RAP null, LDLR null and VLDLR null mice were isolated using a modified procedure, as we described<sup>32</sup>.

First, we determined  $^{125}\text{I}$ -labeled A $\beta$ 40, A $\beta$ 42, Dutch/Iowa A $\beta$ 40 and reverse 40-1 A $\beta$  binding to endogenous LRP at the abluminal side of brain capillaries. Capillaries were incubated in 0.6 ml Eppendorf tubes in the assay buffer (mock CSF containing 1 mM sodium perchlorate to prevent free iodide binding/uptake) with 2 nM  $^{125}\text{I}$ -A $\beta$  ligand at 4°C for 30 min in the absence and presence of RAP (500 nM), LRP-specific IgG (Fab<sub>2</sub>) and unlabeled A $\beta$  (500 nM). After 30 min, incubation buffer containing unbound ligand was removed and capillaries washed, lysed in low SDS lysis buffer and counted.

Second, we determined internalization of  $^{125}\text{I}$ -labeled A $\beta$ 40, A $\beta$ 42 and Dutch/Iowa A $\beta$ 40 at the abluminal surface of brain capillaries and the rate of LRP endocytosis in the presence of each isoform. Capillaries were incubated in 0.6 ml Eppendorf tubes in cold assay buffer with 2 nM  $^{125}\text{I}$ -A $\beta$  ligand at 4°C for 30 min in the absence and presence of RAP (500 nM), LRP-specific IgG (Fab<sub>2</sub>) and unlabeled A $\beta$  (500 nM), as above. After 30 min, unbound ligand was removed by washing capillaries with cold assay buffer. Ice-cold stop/strip solution (0.2 M acetic acid, pH 2.6, 0.1 M NaCl) was added to one Eppendorf tube that was kept on ice. The remaining Eppendorf tubes were placed in a 37°C water bath, and the assay buffer at 37°C added quickly to capillaries to initiate ligand internalization. At predetermined times at 30 s, 1, 2 and 5 min, Eppendorf tubes were quickly placed on ice, and incubated for 12 min with the ice-cold stop/strip solution (0.2 M acetic acid, pH 2.6, 0.1 M NaCl) to remove ligand from the capillary abluminal cell surface. Capillaries were solubilized with low SDS lysis buffer and counted. The sum of internalized ligand plus the ligand associated with the abluminal capillary cell surface represented the amount of ligand available for internalization. The fraction of ligand internalized at each time point was plotted, and the half-time for internalization ( $t_{1/2}$ ) calculated, as described<sup>33;34</sup>.

$^{125}\text{I}$ -A $\beta$  ligand degradation was determined in brain capillary endothelial cell lysates and in the incubation medium at the end of internalization studies using the TCA precipitation assay, as reported<sup>29</sup>.

To determine kinetics of  $^{125}\text{I}$ -A $\beta$  brain capillary unidirectional uptake, capillaries were incubated with  $^{125}\text{I}$ -labeled A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40) at 37°C for 1 min. Self- and cross-inhibition studies were performed with unlabeled A $\beta$ 40 from 1 to 120 nM, and unlabeled A $\beta$ 42 and mutant A $\beta$  (Dutch40, Dutch42,

Dutch/Iowa40) at 40 nM, RAP (500 nM) and LRP-specific polyclonal N20 antibody (60  $\mu$ g/ml).

### Brain clearance studies

Studies were performed according to the National Institutes of Health guidelines using an approved institutional protocol. CNS clearance of  $^{125}$ I-labeled A $\beta$  ligands was determined simultaneously with  $^{14}$ C-inulin (reference marker) in male C57Bl/6, RAP null, LDLR null and VLDLR null mice 8-10 weeks old, using a procedure as described<sup>20</sup>. Briefly, a stainless steel guide cannula was implanted stereotaxically into the right caudate-putamen of anesthetized mice (0.5 mg/kg ketamine and 5 mg/kg xylazine I.P.). Coordinates for tip of the cannula were 0.9 mm anterior and 1.9 mm lateral to the bregma and 2.9 mm below the surface of the brain. Animals were allowed to recover after surgery prior to radiotracer studies. The experiments were performed before substantial chronic processes have occurred, as assessed by histological analysis of tissue, *ie*, negative staining for astrocytes (glial fibrillar acidic protein) and activated microglia (anti-phosphotyrosine), but allowing time for BBB repair to large molecules<sup>35</sup>, typically 4–6 h after the cannula insertion, as reported<sup>36</sup>. Tracer fluid (0.5  $\mu$ L) containing [ $^{125}$ I]-A $\beta$  and  $^{14}$ C-inulin was injected into brain ISF over 5 min via an ultra micro-pump with a micro4-controller (World Precision Instruments, Sarasota, FL). When the effects of different unlabeled molecular reagents are tested, these were injected simultaneously with radiolabeled ligands. For self-inhibition studies, the uptake of  $^{125}$ I-A $\beta$ 40 and  $^{125}$ I-mutant A $\beta$  (Dutch/Iowa40) was studied over a range of carrier concentrations from 0.5-120 nM. For cross-inhibition studies, efflux of  $^{125}$ I-test-peptides was studied at a carrier concentration of 40 nM and the inhibitory concentration of unlabeled A $\beta$  peptides at 120 nM. Brain and blood were sampled 30 min after tracers injection and prepared for radioactivity analysis by TCA, HPLC and SDS-PAGE/immunoprecipitation analysis to determine the molecular forms of test-tracers, as described<sup>20;29;30</sup>. Gamma counting was performed using Wallac Vizard Gamma Counter (Perkin Elmer, Meriden, CT) and beta-counting using Tri-carb 2100 Liquid Scintillation Counter (Perkin Elmer, CT). Previous studies with  $^{125}$ I-labeled A $\beta$  demonstrated an excellent correlation between TCA and HPLC methods. The intactness of  $^{125}$ I-labeled A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40) injected into the brain ISF was > 99% by TCA/HPLC analysis. The A $\beta$  standards eluted between 29.1 and 31.2 min for different A $\beta$  peptides. For SDS-PAGE analysis, TCA precipitated samples were resuspended in 1% SDS, vortexed and incubated at 55°C for 5 min, then neutralized, boiled for 3 min, homogenized and analyzed by electrophoresis in 10% Tris-tricine gels followed by fluorography. Methodological details were as we reported<sup>20;29;30</sup>.

### Calculations

$^{125}\text{I}$ -A $\beta$  brain capillary uptake was corrected for the distribution of  $^{14}\text{C}$ -inulin (extracellular space marker) and determined as the tissue to medium ratio as: c.p.m. for TCA-precipitable  $^{125}\text{I}$ -radioactivity (mg capillary protein)/c.p.m. for TCA-precipitable  $^{125}\text{I}$ -radioactivity (ml medium) (1). All calculations of clearance parameters were as reported<sup>20</sup>. Briefly, the percentage of radioactivity remaining in the brain after microinjection was determined as % recovery in brain =  $100 \times (N_b/N_i)$  (2), where,  $N_b$  is the radioactivity remaining in the brain at the end of the experiment and  $N_i$  is the radioactivity injected into the brain ISF, ie, the d.p.m. for  $^{14}\text{C}$ -inulin and the c.p.m. for TCA-precipitable  $^{125}\text{I}$ -radioactivity (intact A $\beta$ ). The percentage of A $\beta$  cleared through the BBB was calculated as  $[(1 - N_{b(\text{A}\beta)}/N_{i(\text{A}\beta)}) - (1 - N_{b(\text{inulin})}/N_{i(\text{inulin})})] \times 100$ , using a standard time of 30 min (3). Efflux of A $\beta$  from brain ISF via transport across the BBB at different concentrations of peptides,  $J_{\text{out}}$ , was calculated as  $[(1 - N_{b(\text{A}\beta)}/N_{i(\text{A}\beta)}) - (1 - N_{b(\text{inulin})}/N_{i(\text{inulin})})]/T \times C_{\text{A}\beta}$  (4) where  $C_{\text{A}\beta}$  is A $\beta$  concentration in the infusate. The half-saturation concentration for A $\beta$  elimination via BBB transport,  $K_m$ , was calculated from  $J_{\text{out}} = \text{Cl}_{\text{max}}/(K_m + C_{\text{A}\beta})$  (4), where  $\text{Cl}_{\text{max}}$  (pmol/s/L ISF) represents the maximal efflux capacity for the saturable A $\beta$  efflux across the BBB corrected for the rate of ISF flow, as we reported<sup>20</sup>. The  $K_m$  value for A $\beta$ 40 uptake by isolated brain capillaries was calculated using Michaelis-Menten analysis. The inhibitory constants,  $K_i$ , were calculated from the velocity ratios<sup>30</sup> as  $K_i = (J_i \times K_m \times C_i)/(J_{\text{out}} - J_i)(K_m + C_{\text{A}\beta 40})$ , where  $C_i$  and  $C_{\text{A}\beta 40}$  were the inhibitory concentrations of test-A $\beta$  peptide and A $\beta$ 40 in the infusate *in vivo* or incubation medium *in vitro*. Kinetic constants were obtained by a non-linear regression curve fitting (Prism 3.0).

### Transgenic mice

We used Tg-2576 mice in a C57Bl6/SJL background<sup>25</sup> and Tg-DI (Dutch/Iowa) mice in C57Bl6 background<sup>37</sup>. Human APP (770 isoform) cDNA harboring the Swedish (KM670/671NL), Dutch (E693Q), and Iowa (D694N) mutations was subcloned between exons II and IV of a Thy-1.2 expression cassette (gift from Dr. F. LaFerla, University of California, Irvine). The 9 kb transgene was liberated by NotI/PvuI digestion, purified, and microinjected into pronuclei of C57BL/6 single-cell embryos at the Stony Brook Transgenic Mouse Facility. Founder transgenic mice were identified by Southern blot analysis of tail DNA. Transgenic offspring were determined by PCR analysis of tail DNA using the following primers for human APP: 5'-CCTGATTGATACCAAGGAAGGCATCCTG-3': 5'-GTCATCATCGGCTTCTTCTTCTCCACC-3' (generating 500 base pair product).

RAP null, VLDLR null and LDLR null mice were purchased from Jackson Laboratories, Bar Harbor, ME.

**Quantification of A $\beta$** 

Soluble and insoluble pools of A $\beta$  were determined by ELISA of carbonate extracted forebrain tissue and of guanidine lysates of the insoluble pellets resulting from the carbonate extracted brain tissue, respectively<sup>38</sup>. Levels of total A $\beta$  were compared between Tg-2576 and Tg-DI mice.

**Histological analysis**

For neuropathological analysis on mouse brain tissue in Tg-2576 and Tg-DI mice, tissue sections were cut from mouse brain hemispheres in the sagittal plane either at 5  $\mu$ m (paraffin embedded fixed tissue) or 14  $\mu$ m (fresh frozen tissue). A $\beta$  immunoreactive deposits were identified with human specific monoclonal mouse antibody 66.1 to A $\beta$ <sup>29</sup>. For LRP staining on brain microvessels in RAP null, Tg-2576, Tg-DI and wild-type mice, we used 14  $\mu$ m frozen acetone fixed tissue sections and double immunostaining for LRP and CD31 (endothelial marker). LRP-specific IgG (5A6) was used as a primary antibody. Biotinylated anti-mouse IgG was used as a secondary antibody and was detected with fluorescein streptavidin (1:1000, Vector Laboratories, Inc., Burlingame, CA). M.O.M. kit (Vector Laboratories, Inc., Burlingame, CA) was used to block endogenous IgG, as described<sup>39</sup>. For CD31 staining, mouse CD31-specific IgG was used as a primary antibody, and Alexa Fluor 594 donkey anti-rat IgG (1:500, Molecular Probes, Inc. Eugene, OR) as a secondary antibody. For immunocytochemical analysis of LRP on brain microvessels in tissue from AD patients, patients with cerebrovascular  $\beta$ -amyloidosis Dutch-type and age-matched controls, we used cryostat sections (10  $\mu$ m) of frontal cortex (Brodmann area 9/10) that were air dried. LRP was detected with 8G1 human LRP-specific IgG as a primary antibody and rhodamine red goat anti-mouse IgG (1:150, Molecular Probes, Eugene, OR) as a secondary antibody. The anti-vWF (endothelial marker) primary antibody was detected with fluorescein goat anti-rabbit IgG (1:150, Molecular Probes, Eugene, OR). Image analysis was performed using Olympus AX70 microscope equipped with the SPOT digital camera. Ten randomly selected fields in each region from ten sections spanning the entire hemisphere from 4 mice/group or from Brodmann A9/10 areas in humans was performed.

**Human brain endothelial cells**

Human BEC were isolated from rapid autopsies of neurologically normal young individuals after trauma. BEC were characterized and cultured as we described<sup>40</sup> and incubated with different A $\beta$  isoforms at concentrations ranging from 1 nM to 20  $\mu$ M within 48 h. Cells were lysed and equal amounts of proteins electrophoresed (10  $\mu$ g/ml) on 10% SDS-polyacrylamide gel, transferred onto nitrocellulose membrane and probed with 5A6 ( $\beta$ -chain) or 8G1 ( $\alpha$ -chain) human LRP-specific

IgGs, RAGE-specific IgG<sup>29</sup> or transferrin-specific IgG. The relative density of each protein was determined by scanning densitometry using  $\beta$ -actin as an internal control.

LRP endocytosis rates were studied with <sup>125</sup>I- $\alpha_2$ M\*, an LRP-specific ligand. BEC were incubated with 20  $\mu$ M A $\beta$ 40, A $\beta$ 42 and Dutch/Iowa A $\beta$ 40 at 37°C for 48 h in 12-well plates, and then switched to A $\beta$ -free medium containing 5 mM CaCl<sub>2</sub> and 5 nM <sup>125</sup>I- $\alpha_2$ M\* at 4°C for 30 min followed by incubations with  $\alpha_2$ M\*-free medium at 37°C to initiate internalization, as reported<sup>33</sup>.

#### **Metabolic labeling**

Human BEC ( $4 \times 10^5$ ) were pulsed for 1 h at 37°C with 400  $\mu$ Ci of [<sup>35</sup>S]-methionine (> 1000 Ci/mmol; Perkin Elmer Life Sci. Inc., Boston, MA) in methionine-free Dulbecco modified Eagle medium (GIBCO BRL, New York, NY), as described<sup>41</sup>. Cells were chased at indicated times within 48 h. Cell lysates were immunoprecipitated with LRP-515-kDa  $\alpha$ -chain specific IgG (8G1) on SDS-PAGE. The intensity of signal was quantified in pixels using Storm 860 Phosphoimager (Amersham Biosciences Corp., Piscataway, NJ).

#### **Neuronal cultures and A $\beta$ production**

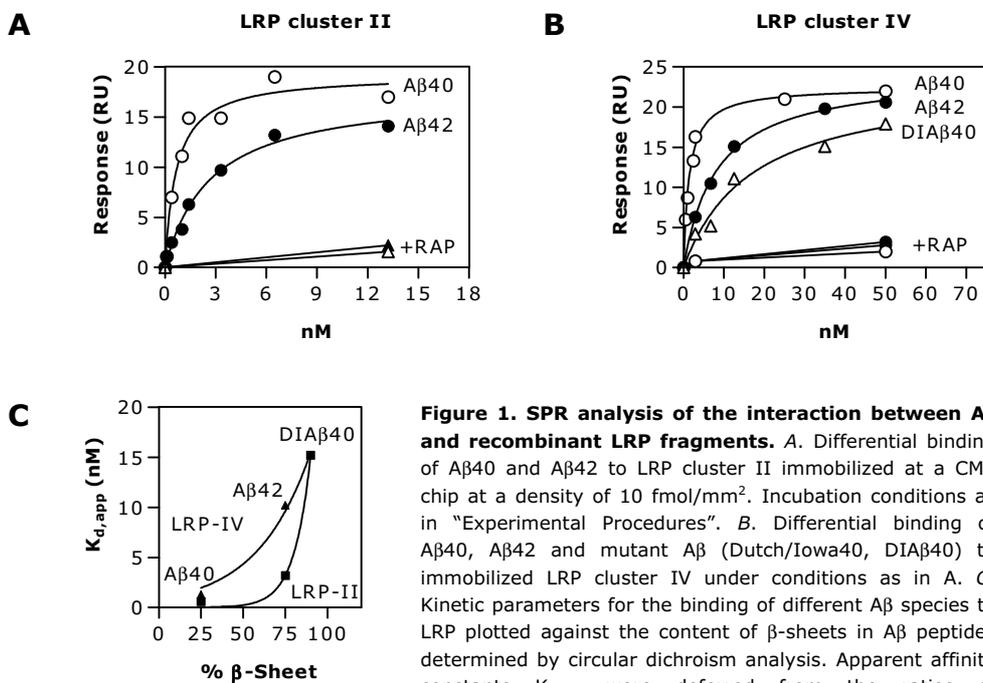
Primary mouse neuronal cultures were prepared as we described<sup>42</sup>. Briefly, cerebral cortices were dissected from fetal heterozygous Tg-DI mice<sup>37</sup> and Tg-2576 mice<sup>25</sup> at 16 days of gestation, treated with trypsin for 10 min at 37°C, and dissociated with trituration. Cell suspensions were plated at  $5 \times 10^5$  cells per well on 12-well plates in serum-free Neurobasal medium plus B27 supplement (GIBCO-BRL, Rockville, MA). The absence of astrocytes was confirmed by the lack of glial fibrillar acidic protein. Cultures were maintained in a humidified 5% CO<sub>2</sub> incubator at 37°C for 5 days and the levels of A $\beta$  in the culture medium determined by a quantitative immunoblot analysis using 4G8 antibody (1:1,000, 1  $\mu$ g/ml, Signet, Dedham, MA), similar as described<sup>15</sup>.

#### **Statistical Analysis**

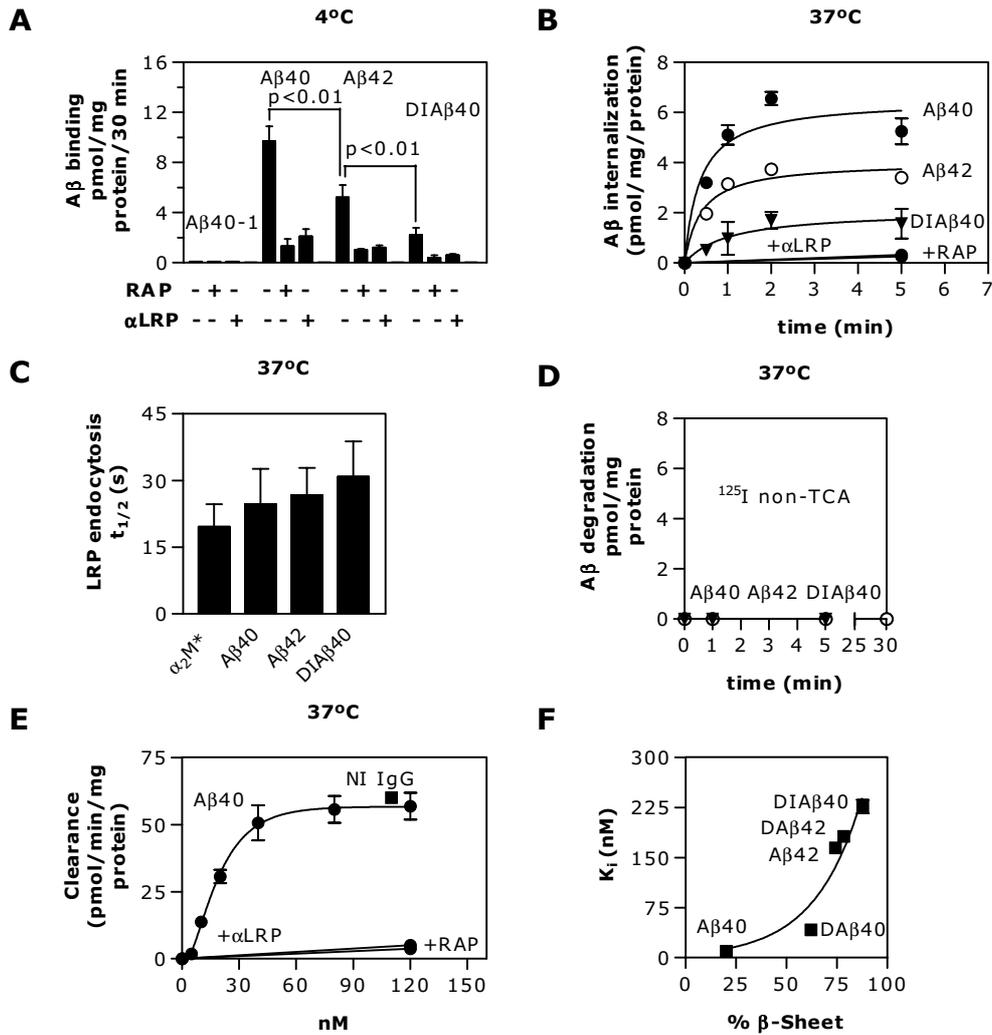
Data were analyzed by multifactorial analysis of variance, Student's t test and Dunnett's t test.

## Results

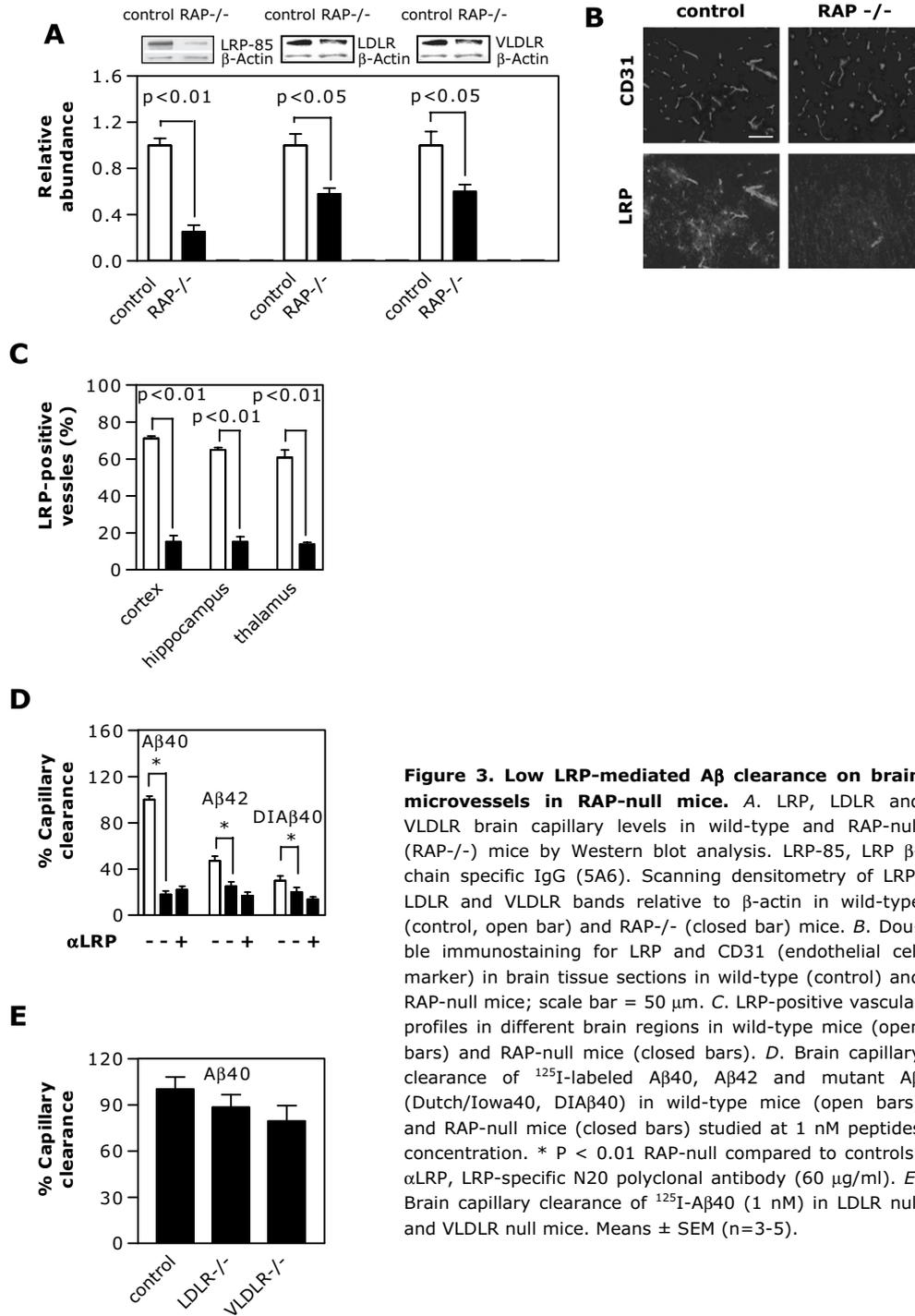
Fig. 1A-B show high affinity binding of monomeric A $\beta$ 40 to immobilized LRP clusters II and IV with  $K_d$  values of  $0.57 \pm 0.12$  and  $1.24 \pm 0.01$  nM, respectively, determined by the surface plasmon resonance (SPR) analysis. In contrast, A $\beta$ 42 and vasculotropic mutant A $\beta$  (Dutch/Iowa A $\beta$ 40 double mutant model peptide<sup>27</sup>) exhibit greatly reduced binding affinity for LRP clusters II and IV by 6 and 9-fold and 28 and 12-fold, respectively, compared to A $\beta$ 40. The  $K_d$  values for A $\beta$ 42 binding to LRP II and IV clusters were  $3.00 \pm 0.11$  and  $10.10 \pm 0.03$  nM, respectively, and for mutant A $\beta$   $15.10 \pm 0.10$  and  $15.30 \pm 0.07$  nM, respectively. Binding of A $\beta$  to LRP was abolished by RAP, an LRP antagonist<sup>1</sup> (Fig 1A-B). Its affinity to immobilized LRP fragments was greatly reduced by the high content of  $\beta$ -sheets (Fig 1C).



**Figure 1. SPR analysis of the interaction between A $\beta$  and recombinant LRP fragments.** A. Differential binding of A $\beta$ 40 and A $\beta$ 42 to LRP cluster II immobilized at a CM5 chip at a density of 10 fmol/mm<sup>2</sup>. Incubation conditions as in "Experimental Procedures". B. Differential binding of A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40, DIA $\beta$ 40) to immobilized LRP cluster IV under conditions as in A. C. Kinetic parameters for the binding of different A $\beta$  species to LRP plotted against the content of  $\beta$ -sheets in A $\beta$  peptides determined by circular dichroism analysis. Apparent affinity constants  $K_{d,app}$  were derived from the ratios of  $K_{off,app}/K_{on,app}$  as described in "Experimental Procedures". Means  $\pm$  SEM ( $n=3-5$ ); SEM  $\leq$  5% of the mean in A and B.  $K_{d,app}$  (mean  $\pm$  SD) values were determined from 6-9 different concentrations of A $\beta$  and 3-5 independent measurements at each concentration. RAP, receptor-associated protein (500 nM); RU, resonance units.



**Figure 2. LRP-mediated Aβ binding, internalization and clearance on mouse brain capillaries.** A. Binding of <sup>125</sup>I-Aβ ligands (2 nM) to the abluminal brain capillary surface was determined at 4°C for 30 min. RAP (500 nM); αLRP, LRP-specific IgG (Fab<sub>2</sub>, 60 μg/ml); Aβ40-1, reverse Aβ40 peptide. B. Internalization of Aβ peptides at the abluminal brain capillary surface. Capillaries were incubated with <sup>125</sup>I-Aβ ligands (2 nM) at 4°C for 30 min and then at 37°C for the indicated times. The fraction of internalized ligand was calculated as in the "Experimental Procedures" and plotted against the internalization time. RAP (500 nM); αLRP (60 μg/ml). C. The half-time ( $t_{1/2}$ ) of LRP endocytosis determined with α<sub>2</sub>M\* and different Aβ ligands from Fig. 2B. D. Undetectable <sup>125</sup>I non-TCA precipitable counts (<sup>125</sup>I non-TCA) in lysed BEC during internalization of <sup>125</sup>I-Aβ ligands (2 nM). E. Rapid saturable Aβ40 uptake by brain capillaries determined at 37°C within 1 min using <sup>125</sup>I-Aβ40 (1 nM) and increasing concentrations of unlabeled Aβ (1-120 nM). RAP (1 μM); αLRP and a non-immune IgG, NI IgG (60 μg/ml). F.  $K_i$  (inhibitory constant) for LRP-mediated brain capillary clearance of Aβ isoforms plotted against Aβ β-sheet content.  $K_i$  values were determined with <sup>125</sup>I-Aβ40 (2 nM) and 40 nM unlabeled Aβ40, Aβ42 and mutant Aβ (Dutch40, DAβ40; Dutch42, DAβ42; Dutch/Iowa40, DIAβ40). Means ± SEM (n=3-5).

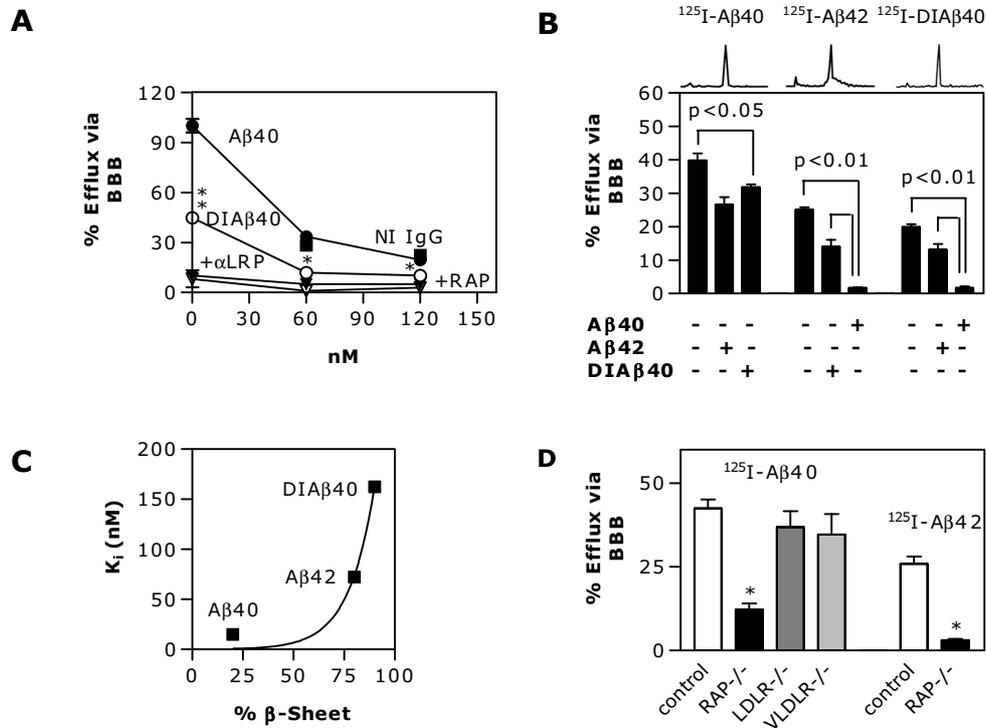


**Figure 3. Low LRP-mediated A $\beta$  clearance on brain microvessels in RAP-null mice.** *A.* LRP, LDLR and VLDLR brain capillary levels in wild-type and RAP-null (RAP<sup>-/-</sup>) mice by Western blot analysis. LRP-85, LRP  $\beta$ -chain specific IgG (5A6). Scanning densitometry of LRP, LDLR and VLDLR bands relative to  $\beta$ -actin in wild-type (control, open bar) and RAP<sup>-/-</sup> (closed bar) mice. *B.* Double immunostaining for LRP and CD31 (endothelial cell marker) in brain tissue sections in wild-type (control) and RAP-null mice; scale bar = 50  $\mu$ m. *C.* LRP-positive vascular profiles in different brain regions in wild-type mice (open bars) and RAP-null mice (closed bars). *D.* Brain capillary clearance of <sup>125</sup>I-labeled A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40, DIA $\beta$ 40) in wild-type mice (open bars) and RAP-null mice (closed bars) studied at 1 nM peptides concentration. \* P < 0.01 RAP-null compared to controls;  $\alpha$ LRP, LRP-specific N20 polyclonal antibody (60  $\mu$ g/ml). *E.* Brain capillary clearance of <sup>125</sup>I-A $\beta$ 40 (1 nM) in LDLR null and VLDLR null mice. Means  $\pm$  SEM (n=3-5).

Figure 2A shows differential LRP-dependent binding of A $\beta$  isoforms at the abluminal side of brain capillaries, *ie*, A $\beta$ 40 > A $\beta$ 42 > Dutch/Iowa A $\beta$ 40, and no binding of the reverse 40-1 peptide, consistent with the SPR findings (Fig 1). Binding of A $\beta$  was displaced by RAP and LRP-specific IgG (Fig 2A), and excess of unlabeled ligand (not shown). Fig 2B illustrates differential LRP-mediated internalization of <sup>125</sup>I-labeled A $\beta$  isoforms into brain endothelium, *ie*, A $\beta$ 40 > A $\beta$ 42 > DIA $\beta$ 40, and inhibition (> 85%) with RAP and LRP-specific IgG. The half-time ( $t_{1/2}$ ) of LRP internalization in the presence of different A $\beta$  isoforms was similar between 26 and 30 s (Fig 1C). During internalization, there was no significant degradation of A $\beta$  by brain endothelium as indicated by negligible non-trichloroacetic acid (TCA)-precipitable counts in brain capillary cell lysates (Fig 1D) and in the medium (not shown). The short-term kinetic uptake studies at 37°C confirmed a saturable LRP-dependent clearance of A $\beta$ 40 by brain capillaries with the Michaelis constant  $K_m$  of  $10 \pm 2$  nM, and > 85% inhibition of uptake by RAP and LRP-specific IgG (Fig 2D). The kinetic inhibitory constants,  $K_i$ , confirmed that Dutch A $\beta$ 40, wild-type A $\beta$ 42, Dutch A $\beta$ 42 and Dutch/Iowa A $\beta$ 40 exhibit 6, 14, 18 and 22-fold lower affinities for LRP-mediated brain capillary clearance than A $\beta$ 40, respectively (Fig 2E).

Next, we compared <sup>125</sup>I-A $\beta$  clearance by brain microvessels derived from RAP null, LDLR null, VLDLR null and control mice. The Western blot analysis indicated that LRP in brain capillaries of RAP null mice was decreased by > 75% compared to controls (Fig 3A), while VLDLR and LDLR levels were reduced approximately by 40%. LRP-positive vascular profiles in different brain regions in RAP null mice were reduced to 14%-16% vs. 60%-70% in controls, as indicated by double staining for LRP and endothelial cell marker CD31 (Fig 3B and C). Deletion of the RAP gene resulted in about 80% reduction in brain capillary clearance of A $\beta$ 40, A $\beta$ 42 and DIA $\beta$ 40 (Fig 3D), while deletion of the VLDLR and LDLR genes did not affect significantly A $\beta$  clearance (Fig 3E).

Fig 4A compares efflux across the BBB of [<sup>125</sup>I]-labeled A $\beta$ 40 and mutant A $\beta$  (Dutch/Iowa A $\beta$ 40) microinfused simultaneously with <sup>14</sup>C-inulin (reference marker) into brain ISF, as described<sup>20</sup>. *In vivo* transcytosis rates of <sup>125</sup>I-labeled A $\beta$  at the BBB were obtained after correcting for the tracer's passive diffusion via the ISF bulk flow by subtracting the elimination rate of <sup>14</sup>C-inulin, as explained in the "Experimental Procedures". At concentrations comparable to physiological levels of soluble A $\beta$  in brain ISF (*ie*,  $\leq 1$  nM)<sup>36</sup>, A $\beta$ 40 was cleared rapidly across the BBB within few seconds. In contrast, only 40% of mutant A $\beta$  was cleared across the BBB over a much longer time period of 30 min. At higher concentrations, mutant A $\beta$  was almost devoid of clearance at the BBB, while A $\beta$ 40 still exhibited a substantial clearance. RAP and an anti-LRP antibody, but not a non-



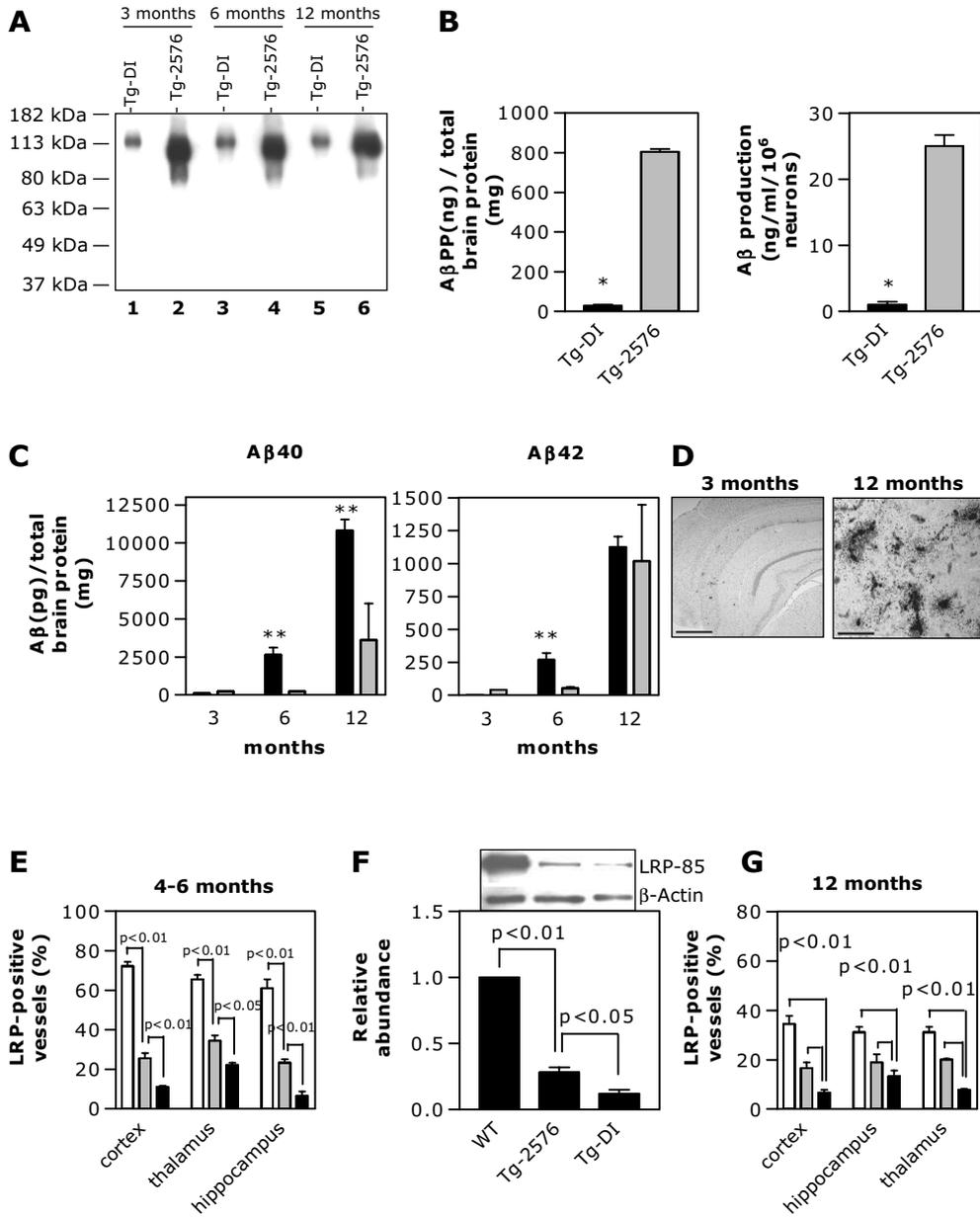
**Figure 4. LRP-mediated transport of A $\beta$  across the mouse BBB *in vivo*.** A. Efflux of mutant A $\beta$  (Dutch/Iowa40, DIA $\beta$ 40; open circles) vs. wild-type A $\beta$ 40 (closed circles) across the BBB within 30 min of brain ISF  $^{125}$ I-A $\beta$  microinjections at 1-120 nM carrier concentration. RAP (1  $\mu$ M);  $\alpha$ LRP, LRP-specific IgG (60  $\mu$ g/ml); NI IgG, non-immune IgG (60  $\mu$ g/ml; closed square). \*\*  $P < 0.01$  and \*  $P < 0.05$  for DIA $\beta$ 40 vs. A $\beta$ 40. B. Efflux across the BBB of  $^{125}$ I-labeled A $\beta$ 40, A $\beta$ 42, and mutant A $\beta$  (Dutch/Iowa40, DIA $\beta$ 40) at 40 nM in the absence and presence of unlabeled A $\beta$  (120 nM).  $^{125}$ I-A $\beta$  monomers (HPLC analysis) in brain homogenates 30 min after microinfusion of  $^{125}$ I/unlabeled A $\beta$  mixture into brain ISF (insets above bars). C.  $K_i$  values for LRP-mediated efflux at the BBB of A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40; DIA $\beta$ 40) plotted against A $\beta$   $\beta$ -sheets content.  $K_i$  values were determined with  $^{125}$ I-A $\beta$ 40 at 40 nM and unlabeled A $\beta$  at 120 nM. D. A $\beta$ 40 and A $\beta$ 42 efflux across the BBB in RAP null mice (RAP $^{-/-}$ ; closed bars) vs. wild-type mice (control, open bars), and A $\beta$ 40 efflux in LDLR null and VLDLR null mice. Means  $\pm$  SEM ( $n=3-8$ ).

immune immunoglobulin G (NI IgG) (Fig 4A), almost abolished A $\beta$  elimination across the BBB.

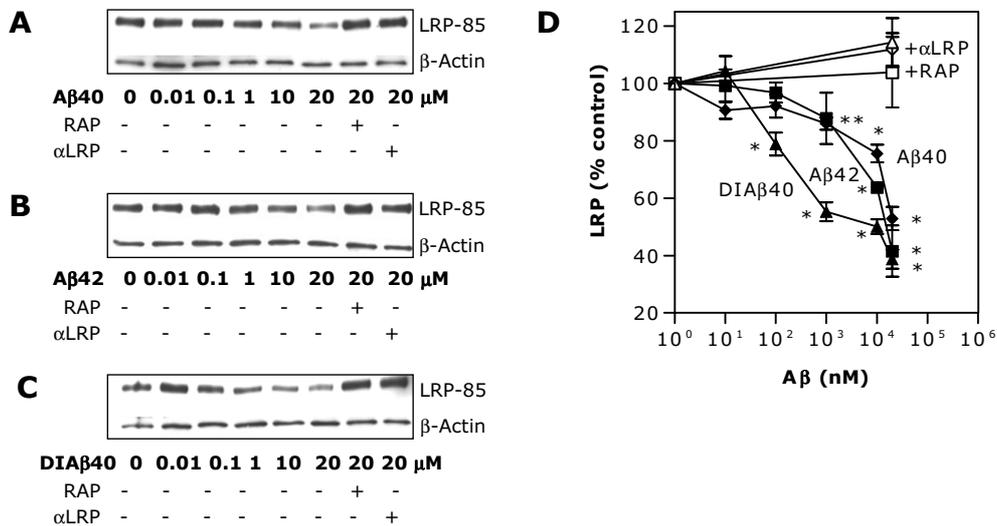
A significant ( $p < 0.05$ ) cross-inhibition of [ $^{125}\text{I}$ ]-A $\beta$ 40 efflux at the BBB by unlabeled A $\beta$ 42 and mutant A $\beta$ , and a pronounced  $> 95\%$  inhibition of [ $^{125}\text{I}$ ]-labeled A $\beta$ 42 and mutant A $\beta$  clearance by unlabeled wild-type A $\beta$ 40 (Fig 4B), indicated all A $\beta$  peptides compete for the same LRP-mediated efflux system to exit the brain, and A $\beta$ 40 exerts a significant retention effect on A $\beta$ 42 and mutant A $\beta$  *in vivo*. All A $\beta$  test-ligands microinfused into brain ISF in the presence of unlabeled A $\beta$  remained  $> 97\%$  in their monomeric forms<sup>20;35</sup>, as confirmed by the HPLC (Fig 4B, insets) and SDS-PAGE analysis (not shown) of brain homogenates. During relatively short incubation time and low nM levels of  $^{125}\text{I}$ -A $\beta$  ligand and the cold peptide,  $^{125}\text{I}$ -A $\beta$  did not bind to unlabeled A $\beta$ , consistent with reports suggesting significantly higher concentrations of A $\beta$  and longer times are required for A $\beta$  oligomerization or aggregation<sup>43;44</sup>.

The  $K_i$  values indicated A $\beta$ 42 and mutant A $\beta$  exhibit 8 and 15-fold lower affinity for LRP-mediated efflux at the BBB *in vivo* than A $\beta$ 40 consistent with their high  $\beta$ -sheet content (Fig 4C). There was 75% to 85% inhibition of A $\beta$ 40 and A $\beta$ 42 rapid efflux across the BBB in RAP null/severely depleted LRP mice (Fig 4D), while deletion of the LDLR and VLDLR genes did not affect significantly A $\beta$  clearance at the BBB (Fig 4D). Since LDLR and VLDLR do not contribute to rapid A $\beta$  clearance in RAP null mice (Fig 3E, 4D), and LRP on neurons *in vivo* does not mediate A $\beta$  clearance<sup>19</sup>, one can speculate that increased amount of amyloid in double crossed RAP null/APP overexpressing mice<sup>45</sup> could be related to inefficient LRP-mediated A $\beta$  clearance across the BBB.

(Page 95) **Figure 5. A $\beta$  accumulation in transgenic mice expressing low LRP-clearance mutant A $\beta$  vs. wild-type A $\beta$ .** A. Immunoblot analysis of human APP in the brain of transgenic Tg-DI mice expressing mutant APP harboring Dutch and Iowa mutations vs. Tg-2576 mice. B. APP levels in the brain of Tg-DI mice vs. Tg-2576 mice at 6 months of age (left) and A $\beta$  production from isolated mouse neurons into culture medium (right) determined by quantitative immunoblot analyses. C. Brain accumulation of low LRP-clearance mutant A $\beta$ 40 and A $\beta$ 42 (Dutch/Iowa, black bars) in Tg-DI mice vs. wild-type A $\beta$ 40 and A $\beta$ 42 (gray bars) in Tg-2576 mice. D. Early deposits of mutant A $\beta$  (Dutch/Iowa) in the brain of Tg-DI mice at 3 months of age (scale bar = 200  $\mu\text{m}$ ), and abundant diffuse deposits with significant intracerebral microvascular A $\beta$  deposits in Tg-DI mice at 12 months of age (scale bar = 50  $\mu\text{m}$ ) detected by immunostaining for A $\beta$ . E. LRP-positive brain microvessels in Tg-DI mice (black bars), Tg-2576 mice (gray bars) and controls (open bars) at 4-6 months of age. F. Western blot analysis of LRP levels in mice in E. G. LRP-positive microvessels in Tg-DI mice (black bars), Tg-2576 mice (gray bars) and controls (open bars) at 12 months of age. Mean  $\pm$  SEM ( $n = 4$  mice). \*  $P < 0.001$ ; \*\*  $P < 0.01$  in B and C for APP and A $\beta$  levels in Tg-DI mice compared to Tg-2576 mice.



To validate our clearance hypothesis for endogenous A $\beta$ , we compared A $\beta$  accumulation in transgenic Dutch/Iowa (Tg-DI) mice expressing low levels of human APP under control of Thy 1.2 neuronal promoter harboring the Dutch<sup>46</sup> and Iowa<sup>47</sup> vasculotropic mutations<sup>37</sup> vs. Tg-2576 APP overexpressing mice<sup>25</sup>. Tg-DI mice produce mainly mutant A $\beta$ 40(Dutch/Iowa) (Fig 5C) that compared to the wild-type A $\beta$ 40 binds to LRP with significantly lower affinity (Fig 2B and C) and exhibits low LRP-clearance on brain capillaries (Fig 2B, 3D) and across the BBB (Fig 4A and C). At 3, 6 and 12 months of age APP levels in the brain of Tg-DI mice vs. Tg-2576 mice were considerably lower by 24-fold (Fig 5A and B, left). In fact, the Tg-DI mice express human APP at < 50% the levels of endogenous mouse APP, as reported in three independent lines<sup>37</sup>. Despite substantially lower levels of human APP (Fig 5B, left) and > 30-fold lower neuronal production of A $\beta$  compared to Tg-2576 mice (Fig 5B, right), the Tg-DI mice exhibited earlier onset and more robust brain accumulations of A $\beta$  than Tg-2576 mice (Fig 5C), *ie*, by 15- and 5-fold higher for the 40 and 42 isoforms, respectively, at 6 months of age.

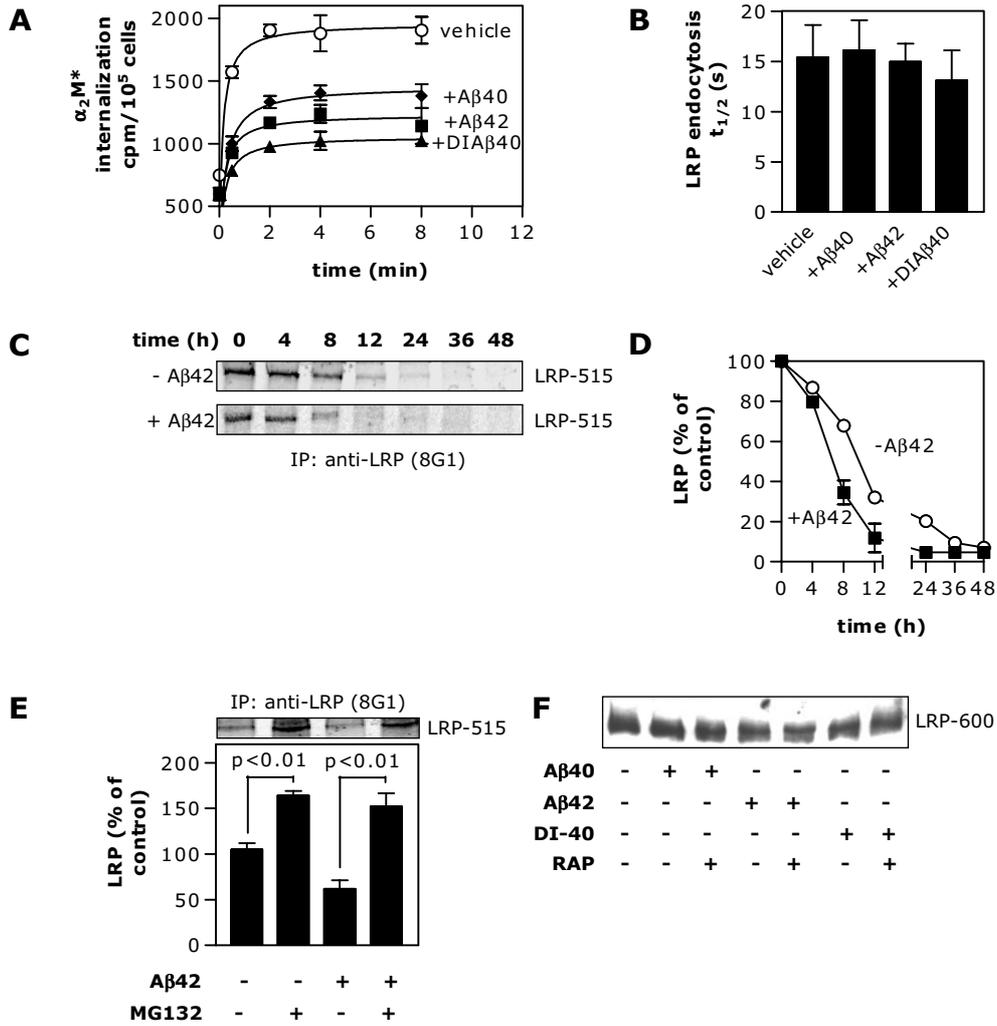


**Figure 6. LRP downregulation in human brain endothelium by excess A $\beta$ .** A-C. Western blot analysis of LRP  $\beta$ -subunit (LRP-85) in brain endothelium exposed for 48 h to different concentrations of A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (Dutch/Iowa40; DIA $\beta$ 40) from 1 nM to 20  $\mu$ M. RAP (1.25  $\mu$ M) and  $\alpha$ LRP, LRP  $\beta$ -chain-specific IgG F(ab)<sub>2</sub> (25  $\mu$ g/ml) were incubated 1 h prior to the addition of A $\beta$  peptides and than for 48 h simultaneously with different A $\beta$  peptides. D. Relative intensity of LRP bands in brain endothelium exposed to different A $\beta$  peptides was determined by scanning densitometry and normalized for  $\beta$ -actin. Mean  $\pm$  SEM (n=3-5 independent experiments); \* P < 0.01; \*\* P < 0.05 for the relative expression of LRP levels in the presence of a given A $\beta$  concentration compared to the absence A $\beta$ .

Tg-DI mice developed early A $\beta$  plaque-like deposits in the cortex and hippocampus at 3 months of age (Fig 5D, left), while the Tg-2576 mice<sup>25;48</sup> initially present A $\beta$  deposits at  $\approx$  9 months of age (not shown). Similarly, the APP/PS-1 knock in mice homozygous for both the humanized APP and the PS-1 knock in mutation, which causes an increase in the amount of A $\beta$ 42, only develop A $\beta$  deposits at  $\approx$  6 months of age<sup>49</sup>. Abundant A $\beta$  plaque-like diffuse deposits with significant microvascular intracerebral association were found in Tg-DI mice at 12 months (Fig 5D, right). Plasma levels of mutant A $\beta$  in Tg-DI mice were extremely low, *ie*, < 25 pM, consistent with the low clearance across the BBB (Fig 4A). Double immunostaining for LRP and endothelial cell marker CD31 indicated substantial reduction of LRP-positive vascular profiles in several brain regions in Tg-DI and Tg-2576 mice at 4-6 months of age, *ie*, only 5-20% and 25-30% of microvessels were positive for LRP, respectively, compared to 65-75% of LRP-positive vessels in age-matched littermate controls (Fig 5E). Significant LRP brain capillary reduction in transgenic mice was confirmed by Western blot analysis (Fig 5F). The decrease in LRP vascular profiles at 12 months of age was also more pronounced in A $\beta$ -accumulating transgenic mice than in controls (Fig 5G).

Fig 6A-D indicate that A $\beta$ 40, A $\beta$ 42 and/or mutant A $\beta$  (Dutch/Iowa A $\beta$ 40) at lower concentrations ( $\leq$  10 nM) do not alter LRP expression in primary human brain endothelial cells (BEC). However, at levels  $\geq$  1  $\mu$ M over longer periods of time (*ie*, 48 h), all A $\beta$  species down regulated LRP in BEC in a concentration-dependent manner, as demonstrated for LRP 85-kDa  $\beta$ -subunit (Fig 6A-D) and 515-kDa  $\alpha$ -subunit (not shown). RAP and LRP-specific IgG, but not a control non-immune IgG (not shown), blocked A $\beta$ -induced LRP down regulation. A $\beta$  treatment did not reduce the expression of other cell surface receptors in BEC, as for example, the levels of the receptor for advanced glycation end products (RAGE) were increased (not shown) as reported<sup>29</sup> and the levels transferrin receptor remain unchanged (not shown). TUNEL and Hoechst staining in BEC were negative, and the levels of lactic acid dehydrogenase in the medium did not increase from control values (not shown).

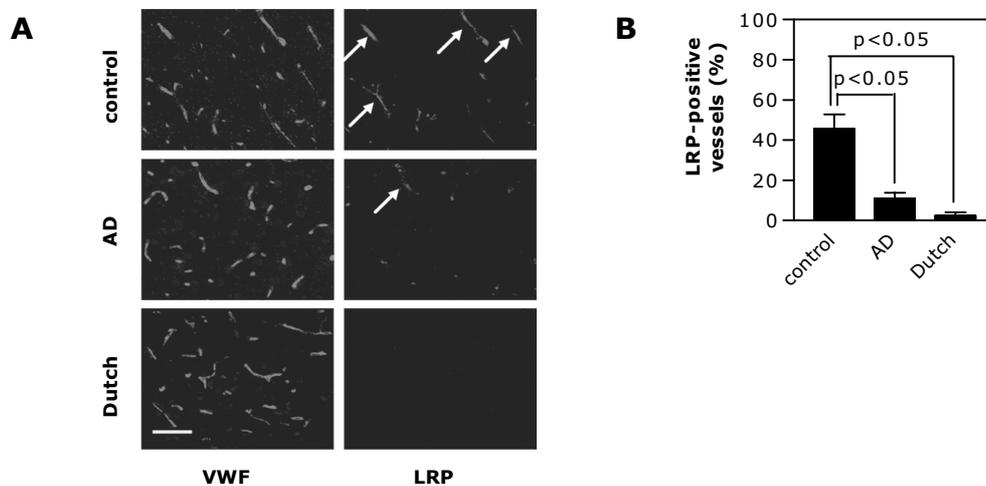
A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  at 20  $\mu$ M over 48 h significantly reduced (> 60%) the amount of internalized <sup>125</sup>I- $\alpha_2$ M\* into human BEC (Fig 7A) consistent with significantly reduced LRP levels (Fig 6D). However, this treatment did not alter significantly the LRP endocytosis rate, and the  $t_{1/2}$  for LRP endocytosis in the presence of different A $\beta$  isoforms remained unchanged compared to controls, *ie*, about 15 s (Fig 7B). On the other hand, the pulse-chase studies revealed that A $\beta$  (20  $\mu$ M) decreases the half-life ( $t_{1/2}$ ) for LRP from 9.6 h to 6.4 h (Fig 7C and D). MG132, an inhibitor of the proteasome-dependent LRP degradation<sup>50</sup>, increased



**Figure 7. High A $\beta$  levels reduce LRP's half-life in human brain endothelium.** *A.* Internalization of <sup>125</sup>I- $\alpha_2M^*$  by human BEC treated with 20  $\mu$ M of A $\beta$ 40, A $\beta$ 42 and mutant A $\beta$  (DIA $\beta$ 40) for 48 h. *B.* The half-life ( $t_{1/2}$ ) for rapid LRP endocytosis in human BEC in *A.* determined with <sup>125</sup>I- $\alpha_2M^*$ . *C.* Cells were labeled by [<sup>35</sup>S]-methionine for 1 h and chased for the indicated times. Autoradiograph of [<sup>35</sup>S]-methionine labeled LRP immunoprecipitated with LRP 515 kDa  $\alpha$ -subunit-specific IgG in the presence and absence of A $\beta$ 42 (20  $\mu$ M). *D.* Graph of LRP levels from three independent experiments as in *A.* \*  $P < 0.05$ ; \*\*  $P < 0.01$  in the presence of A $\beta$  compared to the corresponding control value. *E.* Effects of the proteasomal inhibitor MG132 (20  $\mu$ M) on LRP levels in the presence and absence of A $\beta$ 42 (20  $\mu$ M) determined at 8 h. Autoradiograph of [<sup>35</sup>S]-methionine labeled LRP immunoprecipitated with LRP 515 kDa  $\alpha$ -subunit specific IgG. *F.* Graph of LRP levels from three independent experiments as in *C.* *G.* Western blot analysis for immature 600 kDa LRP (LRP-600) in the presence of A $\beta$ 40, A $\beta$ 42 or mutant A $\beta$  (Dutch/Iowa40, DIA $\beta$ 40) at 20  $\mu$ M within 48 h, as demonstrated with LRP-specific IgG directed to the C-terminal region of LRP (5A6). RAP was used at 1.2  $\mu$ M. Mean  $\pm$  SEM (n=3) in *B* and *D*.

the  $t_{1/2}$  for LRP in control and A $\beta$ -treated cells (Fig 7E and F). Immunoblot analysis with an anti-LRP C-terminal antibody that recognizes immature 600-kDa form of LRP confirmed the levels of immature LRP are not affected by A $\beta$  treatment (Fig 7G), suggesting no effect on LRP synthesis.

LRP-positive vascular profiles in human brain tissue revealed greatly reduced LRP expression in patients with AD and cerebrovascular  $\beta$ -amyloidosis Dutch-type compared to age-matched controls (Fig 8A and B). The number of LRP-positive cerebral vessels dropped from 45% in age-matched controls to  $\approx$  12% in AD, and was barely detectable in patients with Dutch-type cerebrovascular  $\beta$ -amyloidosis.



**Figure 8. LRP expression in brain microvessels in Alzheimer's and Dutch patients.** A. Immunostaining for Von Willebrand Factor (VWF, endothelial cell marker) and LRP in cortical sections (Brodmann area 10) in age-matched controls and AD and Dutch patients with cerebrovascular  $\beta$ -amyloidosis. Scale bar = 50  $\mu$ m. B. LRP-positive vessels in controls, AD and Dutch patients. Mean  $\pm$  SEM (n=4 cases/group).

## Discussion

A $\beta$  accumulation in the brain is a chief pathogenic event<sup>22</sup>. Increased A $\beta$ 42 levels lead to formation of neurotoxic A $\beta$  oligomers resulting in progressive synaptic, neuritic and neuronal dysfunction<sup>43;44;51;52</sup>. Missense mutations inside the A $\beta$  sequence associate with vascular deposits in patients with the Dutch mutation (G to C at codon 693, Glu to Gln at position 22<sup>46</sup>) and Iowa mutation (G to A at codon 694, Asp to Asn at position 23<sup>47</sup>). How A $\beta$  accumulates in the brain is unclear. Increased A $\beta$  production can explain a small fraction of early onset AD familial cases bearing inherited mutations in the APP gene flanking the A $\beta$  coding region (*ie*, Swedish mutation) or PS-1 or 2 genes. This leaves inefficient A $\beta$  clearance as a major mechanism mediating A $\beta$  cerebral accumulation in late-onset AD<sup>53;54</sup> and familial forms of cerebrovascular  $\beta$ -amyloidoses<sup>55</sup>.

Here, we report LRP/A $\beta$  direct interaction at the BBB critically influences A $\beta$  brain accumulation by promoting retention of high  $\beta$ -sheet neurotoxic A $\beta$ 42 and vasculotropic mutant A $\beta$ , while clearing soluble A $\beta$ 40 from brain ISF to blood. The SPR analysis (*Fig 1*) used as a first step to characterize A $\beta$  interaction with the LRP ligand-binding domains<sup>1</sup> revealed A $\beta$ 40 binds to LRP with high affinity compared to A $\beta$ 42 and mutant A $\beta$ . Consistent with the SPR findings, we next showed that LRP-mediated A $\beta$  brain capillary binding and endocytosis *in vitro* and transcytosis across the BBB *in vivo* in mice are substantially reduced by the high  $\beta$ -sheet content in A $\beta$  (*Fig 2-4*), and deletion of the RAP gene, but not the LDLR and VLDLR genes (*Fig 3 and 4*).

In contrast to remarkable LRP-mediated A $\beta$ 40 binding and endocytosis at the abluminal side of brain microvessels (*Fig 2 and 3*) and abluminal to luminal transcytosis across the BBB *in vivo* (*Fig 4*), our earlier study failed to detect significant receptor-mediated A $\beta$ 40 basolateral to apical transport in cultured brain endothelial monolayers<sup>56</sup>. It is possible that the density of LRP receptors at the basolateral membrane in the monolayer cultures<sup>56</sup> was much lower than at the abluminal membrane of freshly isolated brain capillaries and at the BBB *in vivo* (*Fig 2-4*), which could account for the loss of LRP-dependent A $\beta$ 40 basolateral to apical flux in monolayers<sup>56</sup> and may explain the discrepancies between results obtained in the monolayer transport model vs. present and previous study<sup>20</sup>.

The present data show that rapid interaction of A $\beta$  with LRP does not require chaperone molecules. However, this does not rule out the possibility that A $\beta$  may also interact with LRP via chaperone molecules. Thus, both direct and indirect interaction of A $\beta$  with LRP may exist in parallel. For example,  $\alpha_2$ M is implicated in slow and/or delayed reuptake of secreted A $\beta$  into neurons isolated from transgenic APP overexpressing mice which might require LRP<sup>15</sup>, and may also influ-

ence slow uptake of A $\beta$  into mouse fibroblasts via LRP<sup>16</sup> and/or degradation of A $\beta$  in U97 human glioblastoma cells and mouse fibroblasts via  $\alpha_2$ M requires LRP<sup>17</sup>.

Despite extremely low human APP expression/A $\beta$  production, *ie*, 25-30-fold lower than Tg-2576 mice, Tg-DI mice<sup>37</sup> expressing low LRP-clearance mutant A $\beta$  develop robust A $\beta$  brain accumulations much earlier than Tg-2576 A $\beta$ -overproducing mice<sup>25;48</sup> (Fig 5). Significant intra-parenchymal microvascular A $\beta$  association points to brain's blood vessels as a site of deficient clearance in Tg-DI mice consistent with prominent cerebrovascular pathology in Dutch and Iowa patients<sup>23</sup>. Increased fibrillogenicity of mutant Dutch/Iowa A $\beta$ <sup>27</sup> may contribute to its decreased clearance from brain, but the presence of mainly non-fibrillar A $\beta$  parenchymal deposits in Tg-DI mice would argue against this possibility. In addition to reduced LRP-mediated A $\beta$  efflux, an increased formation of non-fibrillar A $\beta$  species might favor A $\beta$  deposition in this model.

The present data show that A $\beta$  at pathological concentrations reduces LRP levels in brain endothelium by accelerating proteasome-dependent LRP degradation<sup>50</sup>, while the receptor's internalization and synthesis are not affected (Fig 6 and 7). In contrast to LRP downregulation by simultaneous overexpression of secreted ligands such as ApoE, resulting in retention and degradation of LRP in the endoplasmic reticulum (ER)<sup>2;57</sup>, binding of internalized A $\beta$  to immature LRP in the ER compartment is not implicated in LRP degradation by extracellular A $\beta$ . One can speculate that A $\beta$  accumulations adjacent to brain vasculature may down regulate LRP at the BBB *in vivo* consistent with reduced brain capillary LRP levels in transgenic A $\beta$ -accumulating mice (Fig 5E-G), AD and patients with the Dutch-type cerebrovascular  $\beta$ -amyloidosis (Fig 8), and reduced total brain LRP in AD<sup>16;20</sup>.

In contrast to LRP, RAGE mediates a continuous influx of circulating A $\beta$  into the brain and is overexpressed in brain vasculature in transgenic APP models and in AD<sup>29</sup>. Thus, an imbalance between LRP-mediated and RAGE-mediated A $\beta$  transport at the BBB would create a positive feedback amplification mechanism for A $\beta$  accumulation in AD and related familial cerebrovascular disorders. The present findings raise a possibility that efficacy of A $\beta$  lowering therapies with peripheral A $\beta$  sequestering agents<sup>29;58;59</sup> may depend at least in part on preserved LRP activity at the BBB. Whether therapeutic modifications of A $\beta$ /LRP interaction at the BBB will have a major impact on controlling dementia remains to be seen in future.

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chapter **5**

**Cellular uptake of C4b-binding protein is coordinated by heparan sulfate proteoglycans and LDL receptor-related protein**

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*In preparation*

## Abstract

C4b-binding protein (C4BP) is a plasma protein involved in both complement and coagulation pathways. C4BP interacts with anticoagulant protein S via its  $\beta$ -chain, while the  $\alpha$ -chain comprises binding sites for C4b, heparin and bacterial components. Recently, low-density lipoprotein receptor-related protein (LRP) was found to mediate the cellular uptake of C4BP *in vitro* and *in vivo* [Westein et al, 2002]. Moreover, this uptake involved the binding of LRP to the C4BP  $\alpha$ -chains. In the present study, the interaction between C4BP and LRP is further characterized. By using an array of C4BP mutants with replacements of positively charged residues in the  $\alpha$ -chain, we demonstrated that several cationic amino acids contribute to the interaction with LRP when assessed by surface plasmon resonance analysis. Furthermore, the triple mutant rC4BP $\alpha$ /R39Q-R64Q-R66Q is also less efficiently degraded by CHO cells expressing LRP. Since C4BP efficiently binds heparin, we explored the potential cooperative action of LRP with cell-surface exposed heparan sulfate proteoglycans (HSPGs) in the endocytosis of C4BP. Indeed, CHO cells lacking HSPGs bound and internalized less C4BP than wild-type CHO cells, which endogenously express LRP. Finally, we assessed the effect of protein S on the interaction between C4BP and LRP. Using two distinct approaches, we observed that protein S is unable to interfere with complex formation between LRP and C4BP. In conclusion, binding and uptake of C4BP is coordinated by both HSPGs and LRP. Furthermore, the formation of the trimolecular LRP/C4BP/protein S complex suggests that LRP indirectly contributes to the clearance of C4BP-bound protein S.

## Introduction

C4b-binding protein (C4BP) is a component of both complement and coagulation pathways<sup>1;2</sup>. In the complement pathway, it enhances factor I-mediated degradation of C4b and C3b<sup>3;4</sup>, and prevents the assembly of the C3-convertase complex by binding C4b<sup>5</sup>. C4BP is an approximately 570-kDa plasma protein with an average plasma concentration of 150 mg/L<sup>6-8</sup>. It is synthesized in the liver as a heterologous, multimeric protein: it contains 6 or 7 identical  $\alpha$ -chains of 70 kDa (C4BP $\alpha$ ) and 80% of the molecules also contains a unique  $\beta$ -chain of 45 kDa (C4BP $\alpha\beta$ )<sup>9;10</sup>. All chains are interconnected via their C-terminal domains through disulfide bridges<sup>11</sup>. The  $\alpha$ - and  $\beta$ -chains are composed of 8 or 3 so-called complement control protein (CCP)- or Sushi domains, respectively.

C4BP $\alpha\beta$  plays a role in the coagulation pathway via its  $\beta$ -chain, which has been found to bind protein S<sup>2;12</sup>. Protein S acts as a cofactor for activated protein C in the degradation of coagulation factors Va and VIIIa<sup>13;14</sup>. 60% of plasma protein S is associated with C4BP $\alpha\beta$ . This interaction inhibits the cofactor activity of protein S<sup>15</sup>. The  $\alpha$ -chains harbor binding sites for C4b<sup>11;16</sup>. C4BP is also able to bind other ligands, such as serum amyloid P<sup>17</sup>, factor VIII<sup>18</sup>, components of pathogenic bacteria<sup>19</sup> and heparin<sup>3;16</sup>. The plasma concentration of C4BP is determined by the balance between synthesis and clearance. C4BP $\alpha$  is increased dramatically during an inflammatory process, whereas C4BP $\alpha\beta$  displays stable levels<sup>20</sup>. Recently, we identified low-density lipoprotein (LDL) receptor-related protein (LRP) as a receptor that mediates the cellular clearance of C4BP *in vitro* and *in vivo*<sup>21</sup>.

LRP is a member of the LDL receptor family that is expressed in a wide spectrum of cells, including hepatocytes, fibroblasts, neurons and monocytes<sup>22</sup>. LRP consists of a non-covalently linked heavy and light chain. The 85-kDa light chain comprises the intracellular and transmembrane domain. The intracellular domain harbors 2 NPXY and 1 YXXL motifs that are implicated in binding of adaptor proteins involved in endocytosis and cellular signaling<sup>23;24</sup>. The 515-kDa heavy chain contains four clusters of complement-type repeats. Clusters II and IV play a dominant role in ligand binding<sup>25;26</sup>. Ligands that are recognized by LRP are structurally and functionally unrelated<sup>21</sup>. Among them are apolipoproteins, lipases, coagulation factors and  $\beta$ 2-integrins<sup>27-30</sup>.

The molecular mechanism of cellular C4BP binding has not been completely investigated yet. For other heparin-binding LRP ligands, such as tissue factor pathway inhibitor, factor VIII, activated factor IX and thrombospondin<sup>31-33</sup>, it has been described that their cellular binding is facilitated by heparan sulfate proteoglycans (HSPGs). HSPGs are abundantly expressed at the cell surface and consist of a core protein and heterogeneous heparan sulfate side chains<sup>34</sup>. After

sequestration from the plasma by HSPGs, ligands can be efficiently transferred to their specific receptor, such as LRP.

Heparin has been demonstrated to prevent the binding of C4BP to LRP<sup>21</sup>. The present study focused on the role of the heparin-binding sites of C4BP in its binding to the cell surface and LRP. Our data show that rC4BP $\alpha$  mutants with decreased heparin-affinity<sup>35</sup> were able to bind to immobilized LRP less efficiently. The R39Q-R64Q-R66Q triple mutant showed reduced clearance by mouse fibroblasts. The initial binding of rC4BP to CHO cells was predominantly dependent on the presence of HSPGs, resulting in a delayed LRP-mediated cellular clearance. Finally, the protein S/C4BP $\alpha\beta$  complex was able to bind to LRP, suggesting that LRP indirectly contributes to the binding and probably clearance of protein S.

## Experimental Procedures

### Materials

The Biacore2000 biosensor system and reagents, including an amine-coupling kit and CM5 biosensor chips (research grade), were from Biacore AB (Uppsala, Sweden). All cell culture disposables were from Nunc (Roskilde, Denmark). Bovine Serum Albumin (BSA) fraction V was from Sigma (Zwijndrecht, The Netherlands).

### Proteins

Recombinant C4BP, consisting of the  $\alpha$ -chains but lacking the  $\beta$ -chain (rC4BP $\alpha$ ), was produced using stably transfected baby hamster kidney cell lines, purified by immunoaffinity chromatography as reported previously<sup>36</sup>, and stored in 125 mM NaCl, 0.005% (v/v) Tween 20, 25 mM Hepes (pH 7.4) at -20°C until use. rC4BP $\alpha$  isoforms mutated in positively charged regions were previously described in ref 35. Plasma-derived C4BP containing both  $\alpha$ - and  $\beta$ -chains (C4BP $\alpha\beta$ ) was purified as described<sup>37</sup>. rC4BP $\alpha$  was radio-labeled with Na<sup>125</sup>I (Amersham Biosciences) using IodoBeads (Pierce, Rockford, IL) according to the manufacturer's protocol. Free <sup>125</sup>I was removed by extensive dialysis against 125 mM NaCl, 25 mM Hepes pH 7.4. Purified full-length LRP was purchased from Biomac (Leipzig, Germany). Recombinant LRP cluster IV was purified from the cell culture supernatant using a GST-RAP sepharose column, as reported previously<sup>26</sup>. Protein S was isolated from plasma as described by Hackeng et al<sup>38</sup>.

### Cell culture

Mouse embryonic fibroblasts (MEF-1, ATCC CRL-2241) and their LRP-deficient counterparts PEA-13 cells<sup>39</sup> (ATCC CRL-2216), CHO-K1 (wild-type, ATCC CRL-

61), CHO LRP-/-<sup>40</sup> and xylosyltransferase-deficient CHO cells (resulting in the inability to produce cell surface heparan sulfate proteoglycans (HSPG), ATCC CRL-2242) were from the American Tissue Culture Collection. MEFs and PEAs were cultured in DMEM:F12 medium (Invitrogen, Breda, The Netherlands) and CHO cells in HAM's F12 medium (Cambrex, Verviers, Belgium). Both cell culture media were supplemented with 10% fetal bovine serum (Cambrex, Verviers, Belgium) and 100 U/mL penicillin and 100 µg/mL streptomycin (Invitrogen, Breda, The Netherlands). Cells were grown in a humidified incubator containing 5% CO<sub>2</sub> at 37°C.

#### **Surface plasmon resonance (SPR)**

Binding studies were performed employing a Biacore2000 biosensor system, and SPR analysis was done essentially as described<sup>21;32;41</sup>. LRP was immobilized on a CM5 sensor chip at 7 fmol/mm<sup>2</sup> using the amine-coupling kit as instructed by the supplier. Routinely, a control channel was activated and blocked using the amine-coupling reagents in the absence of protein. rC4BP $\alpha$  was injected until saturation was achieved. Directly after binding, rC4BP $\alpha$  at a higher concentration was injected. Alternatively, C4BP $\alpha\beta$  was perfused over the LRP chip until saturation, directly followed by an injection of protein S. For other experiments, C4BP $\alpha\beta$  or C4BP $\alpha$  were immobilized on a CM5 sensor chip at 7.2 and 6.9 fmol/mm<sup>2</sup>, respectively, as described for LRP. After saturation of immobilized C4BP $\alpha\beta$  with protein S, cluster IV was allowed to bind to the complex. Binding to coated channels was corrected for binding to non-coated channels (< 5% of binding to coated channels). SPR analysis was performed in 125 mM NaCl, 1 mM CaCl<sub>2</sub>, 0.005% (v/v) Tween 20, 25 mM Hepes (pH 7.4) at 25°C at 10 µL/min. Regeneration of the surface of the LRP sensor chip was performed using 100 mM H<sub>3</sub>PO<sub>4</sub>.

#### **Analysis of SPR data**

The maximally obtained response was determined from each injection. Using GraphPad software, response units were plotted against the protein concentration. K<sub>d</sub> values were determined employing one-site binding nonlinear regression.

#### **Cellular binding and degradation**

For cellular binding experiments, CHO cells seeded at least 48h before the start of the experiment on glass coverslips in a 24-wells plate and grown until 90-95% confluence in normal culture medium. Cells were extensively washed with HAM's F12 medium, supplemented with 1% BSA and 2 mM CaCl<sub>2</sub> (assay medium), followed by an incubation for 30 min at 37°C to deplete serum compo-

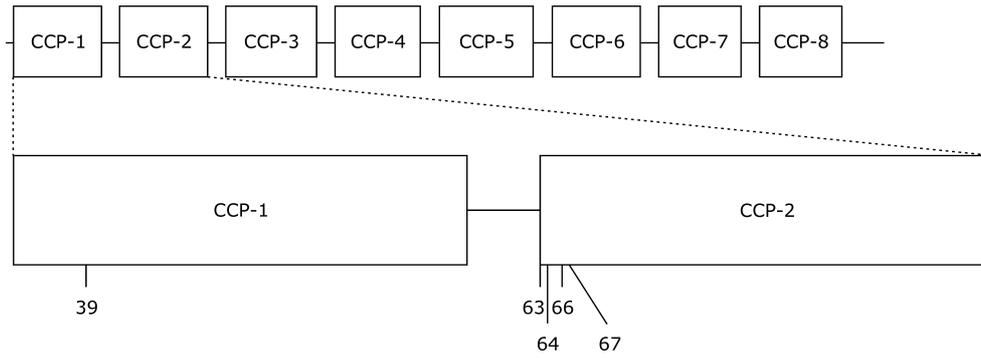
nents. Cells were transferred to ice and then incubated with 250  $\mu$ l assay medium containing 10-100 nM  $^{125}$ I-labeled rC4BP $\alpha$  or  $^{125}$ I-labeled C4BP $\alpha\beta$  mixed with non-labeled rC4BP $\alpha$  or C4BP $\alpha\beta$  to a 1:1 molar ratio for 2h. Subsequently, cells were washed 3 times to remove non-bound material and coverslips were transferred to a new plate. Finally, cells were lysed in 1 M NaOH and supernatant was counted in a  $\gamma$ -counter. In each experiment, also a control coverslip without cells was included to determine non-specific binding. Non-specific binding appeared to be less than 20% compared to the wells that contained cells.

For cellular degradation experiments, MEF and PEA, or CHO cells were directly seeded in 24-wells plates at least 48h before the experiment and grown until 90-95% confluence in normal culture medium. Cells were extensively washed and incubated in assay medium at 37°C for 30 min. 50 nM  $^{125}$ I-rC4BP $\alpha$  mixed with 50 nM non-labeled rC4BP $\alpha$  was added for 1h at 4°C. Cells were washed three times with 500  $\mu$ l of assay medium to remove non-bound material. Subsequently, incubation was allowed to proceed at 37°C in a volume of 500  $\mu$ l. At indicated time points, 30  $\mu$ l samples were taken to determine the amount of degraded material and mixed with 10% trichloroacetic acid (TCA) for at least 30 min. Degraded material is defined as the radioactivity that is soluble in 10% TCA. In all experiments a control was included in which the amount of degradation was examined in the absence of cells.

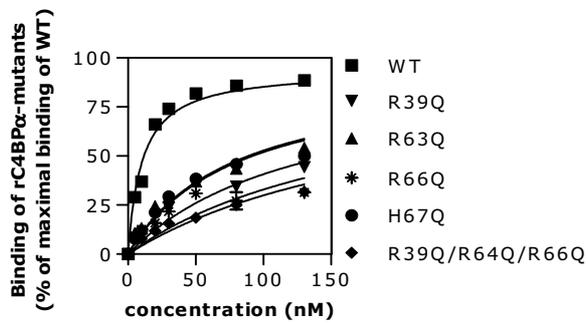
## Results

### **rC4BP $\alpha$ mutants display decreased binding to LRP**

To investigate whether the heparin-binding region of the C4BP $\alpha$  chains contribute to the interaction with LRP, we used a number of previously described mutants (*Fig 1*). In these mutants, one or three positively charged residues have been replaced by glutamine, and each mutation is associated with suboptimal heparin-binding<sup>35</sup>. The interaction of the purified mutants with LRP was tested in equilibrium-binding assays using SPR analysis. Various concentrations of wt-rC4BP $\alpha$  or its mutants were applied to immobilized LRP until equilibrium was reached. For wt-rC4BP $\alpha$ , this approach resulted in a binding-isotherm that allowed the calculation of an apparent  $K_d$ , the value of which being 10.4 nM (*Fig 2*). With regard to the mutants, all of them were less efficient than wt-rC4BP $\alpha$ , albeit to a different extent (*Table 1*). Most severely affected was the triple mutant rC4BP $\alpha$ /R39Q-R64Q-R66Q, which displayed 20.6-fold less affinity for LRP than wt-rC4BP $\alpha$ . Apparently, the LRP-binding site includes residues that are also involved in heparin binding.



**Figure 1. Schematic representation of the C4BP  $\alpha$ -chain.** The  $\alpha$ -chain of C4BP contains 8 complement control protein domains (CCP). CCP-1 and -2 are enlarged. Indicated are positively charged amino acids that were mutated to glutamine residues. In the present study, the following mutants were investigated: R39Q, R63Q, R66Q, H67Q, R39Q-R64Q-R66Q.



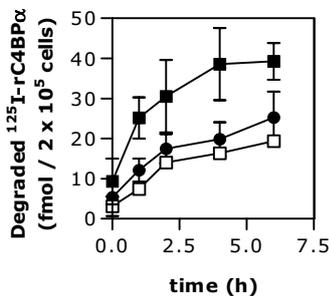
**Figure 2. Binding of rC4BP $\alpha$  mutants to immobilized LRP.** LRP was immobilized on a CM5 sensor chip at 7 fmol/mm<sup>2</sup>. rC4BP $\alpha$  was injected until saturation was achieved. Directly after binding, rC4BP $\alpha$  at a higher concentration was injected. Data were fitted using a one-site regression model.

**Table 1. Kinetic parameters for the binding of rC4BP $\alpha$  mutants to LRP.**

rC4BP $\alpha$ mutant	$K_{d,app}$ (nM)	standard error
WT	10.4	0.7
R39Q	130.9	4.7
R63Q	78.2	3.5
R66Q	185.5	16.7
H67Q	80.9	3.5
R39Q-R64Q-R66Q	214.0	8.4

### Reduced cellular uptake of rC4BP $\alpha$ /R39Q-R64Q-R66Q by LRP-expressing fibroblasts

In a second approach, we tested whether impaired binding of C4BP to LRP is indeed associated with a reduced cellular uptake of this protein. Therefore, experiments were performed using mouse fibroblasts that are genetically deficient for LRP (*ie* PEA-13 cells) or their counterparts that express LRP endogenously (*ie* MEF-1 cells). As expected,  $^{125}\text{I}$ -labeled wt-rC4BP $\alpha$  was efficiently degraded by MEF-1 cells, whereas the amount of degraded wt-rC4BP $\alpha$  was reduced in the LRP-deficient PEA-13 cells (*Fig 3*). With regard to the triple mutant rC4BP $\alpha$ /R39Q-R64Q-R66Q, the amount of degraded protein in LRP-expressing MEF-1 cells was reduced to levels that paralleled that of wt-rC4BP $\alpha$  in LRP-deficient cells. Thus, abrogation of the combined heparin/LRP-binding site in C4BP results in impaired cellular uptake of the protein.

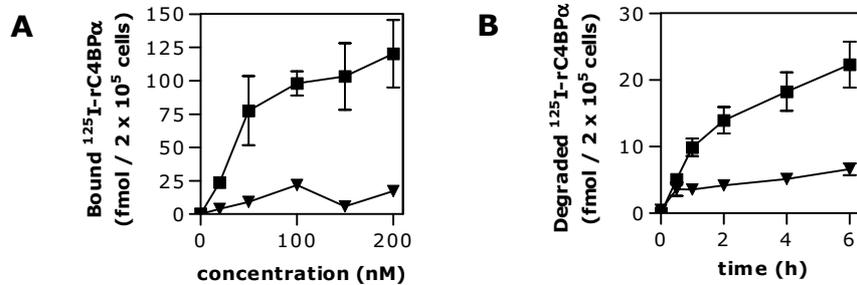


**Figure 3. Cellular degradation of wt and mutant rC4BP $\alpha$ .** LRP+ MEF-1 cells (squares) and LRP- PEA-13 cells (circles) were incubated with a 1:1 mixture of unlabeled and  $^{125}\text{I}$ -labeled wt-rC4BP $\alpha$  (closed symbols) or rC4BP $\alpha$  R39Q-R64Q-R66Q (open symbols) (final concentration 100 nM) at 4°C for 60min. After washing, cells were subsequently incubated at 37°C for indicated time points, and degradation of rC4BP $\alpha$  was determined as described under "Experimental Procedures". Data represent mean  $\pm$  SEM (n=2).

### HSPGs are required for optimal endocytosis of C4BP

The observation that the triple mutant rC4BP $\alpha$ /R39Q-R64Q-R66Q is taken up by the cell in a suboptimal manner, does not automatically allow the conclusion that the reduced uptake is selectively due to an aberrant interaction with LRP. For a number of LRP ligands such as factor VIII and thrombospondin, it has been described that LRP-dependent endocytosis is preceded by the sequestration of these ligands on HSPGs that are present at the cellular surface. It was of interest therefore to examine the contribution of HSPGs to the cellular uptake of C4BP. To this end, binding and degradation of wt-rC4BP $\alpha$  by normal CHO cells (which express LRP endogenously) and HSPG-deficient CHO cells were compared. Binding of  $^{125}\text{I}$ -labeled wt-rC4BP $\alpha$  to normal CHO cells was saturable and dose-dependent, with half-maximal binding occurring at approximately 70 nM (*Fig 4A*). In addition, radioactive TCA-soluble products appeared in time in the medium, indicating that the radiolabeled wt-rC4BP $\alpha$  was degraded by these nor-

mal CHO cells (Fig 4B). In contrast, binding and subsequent degradation were severely reduced in CHO cells that lack HSPG at their cellular surface (Fig 4A and B). Taken together, these data indicate that HSPGs and LRP act in concert in the binding and internalization of C4BP.

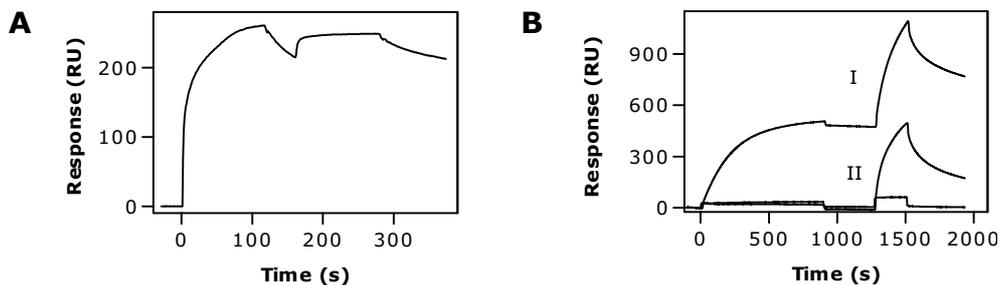


**Figure 4. Cellular binding and degradation of  $^{125}\text{I}$ -rC4BP $\alpha$  to CHO cells.** A. CHO cells (WT (squares) and HSPG (triangles)) were grown on coverslips. After incubation in HAM's F12, 1% BSA and 2 mM  $\text{CaCl}_2$  at 37°C for 30 min, indicated concentrations of  $^{125}\text{I}$ -rC4BP $\alpha$  and nonlabeled rC4BP $\alpha$  were added simultaneously in a 1:1 molar ratio on ice for 2h. Cells were washed extensively and coverslips were transferred to a new plate. Then, cells were lysed and supernatant was counted in a  $\gamma$ -counter. B. CHO cells were incubated with a 1:1 mixture of unlabeled and  $^{125}\text{I}$ -labeled wt-rC4BP $\alpha$  (final concentration 100 nM) at 4°C for 60min. After washing, cells were subsequently incubated at 37°C for indicated time points, and degradation of rC4BP $\alpha$  was determined as described under "Experimental Procedures". Counts were corrected for non-specific binding by subtracting values from wells without cells. Data represent mean  $\pm$  SEM (n=3).

### Simultaneous binding of C4BP $\alpha\beta$ to LRP and protein S

In plasma, the  $\beta$ -chain containing C4BP molecules (spanning approximately 80% of the C4BP population) circulate in complex with anticoagulant protein S. We examined therefore the possibility that not only C4BP $\alpha\beta$  alone but also the C4BP $\alpha\beta$ /protein S complex may interact with LRP. Two distinct experimental approaches employing SPR-analysis were used in this respect. First, immobilized LRP was saturated with C4BP $\alpha\beta$  followed by application of protein S. As shown in figure 5A, C4BP $\alpha\beta$  efficiently bound to immobilized LRP, resulting in a response of 261 RU that gradually declined to 214 RU just before application of protein S. These 214 RU correspond to approximately 0.4 fmol C4BP $\alpha$ /mm $^2$ . Shortly after finalizing LRP injection, protein S was applied. Perfusion of protein S resulted in an increased response, which was calculated to be 34 RU, corresponding to 0.45 fmol protein S/mm $^2$ . Control experiments demonstrated that protein S alone

does not associate with LRP<sup>21</sup>, indicating that the observed response originates from binding of protein S to LRP-bound C4BP $\alpha\beta$ . In an alternative approach, C4BP consisting of both the  $\alpha$ - and  $\beta$ -chains or only of its  $\alpha$ -chains was immobilized onto a CM5-sensorchip (7.2 fmol/mm<sup>2</sup> and 6.9 fmol/mm<sup>2</sup> for C4BP $\alpha\beta$  and C4BP $\alpha$ , respectively). Protein S was subsequently perfused over both channels, resulting in the absence of protein S on the C4BP $\alpha$ -channel and binding of 500 RU (*ie* 6.6 fmol protein S/mm<sup>2</sup>) onto the C4BP $\alpha\beta$ -channel (*Fig 5B*). Subsequently, both channels were perfused with recombinant LRP cluster IV. As depicted in figure 5B, binding of near equal amounts of LRP cluster IV to the C4BP $\alpha$ - and C4BP $\alpha\beta$ -channel was observed. These results are consistent with the view that C4BP $\alpha\beta$  is able to interact simultaneously with LRP and protein S.



**Figure 5. Binding of the C4BP $\alpha\beta$  / protein S complex to LRP.** A. Immobilized LRP (7.0 fmol/mm<sup>2</sup>) was saturated with C4BP $\alpha\beta$  followed by application of protein S. B. C4BP consisting of both the  $\alpha$ - and  $\beta$ -chains (*line I*) or only of its  $\alpha$ -chains (*line II*) were immobilized onto a CM5-sensorchip (7.2 fmol/mm<sup>2</sup> and 6.9 fmol/mm<sup>2</sup> for C4BP $\alpha\beta$  and C4BP $\alpha$ , respectively). Protein S was subsequently perfused over both channels. Then, both channels were perfused with recombinant LRP cluster IV.

## Discussion

C4BP is a multifunctional protein associated with various physiological processes, like complement regulation, coagulation and removal of apoptotic cells<sup>14;42</sup>. Recently, we described that C4BP is a ligand for the endocytic receptor LRP<sup>21</sup>, and in the present study the interaction between both proteins was investigated in more detail. By using a number of previously described mutants (*Fig 1* and ref 35), we have now identified a number of positively charged amino acids located in the C4BP  $\alpha$ -chains that are critical for optimal interaction between C4BP and LRP. Conversion of these residues (*ie* R39, R63, R64, R66, H67) into glutamine residues all resulted in reduced binding to LRP as assessed by SPR

analysis. In addition, the mutant with the triple mutation (rC4BP $\alpha$ /R39Q-R64Q-R66Q) did not only display reduced binding to LRP, but also was taken up by LRP-expressing cells at levels that parallel those in the absence of LRP (Fig 2). The same mutations also affect the interaction with other components, like factor I, C4b, heparin and bacterial components<sup>19;35</sup>. The triple mutant showed a strongly reduced cofactor function in the factor I-mediated C4b binding. In addition, all mutants demonstrated decreased affinity for heparin and streptococcal M proteins. Whether the interaction between C4BP and LRP is completely dependent on charged residues interactions seems unlikely, because even in the triple mutant significant residual binding was observed. Furthermore, binding could not be completely inhibited in the presence of high salt concentrations (> 300 mM, data not shown). This suggests that also hydrophobic residues within the C4BP  $\alpha$ -chains may contribute to the interaction with LRP. In this respect, C4BP is similar to activated factor IXa, which interaction with LRP also involves a number of charged and hydrophobic residues<sup>43</sup>.

The presence of the heparin-binding site within the C4BP  $\alpha$ -chains opens the possibility that C4BP may also associate to HSPGs that were present at the cellular surface. Indeed, a strongly reduced binding and uptake of C4BP was observed in cells that lack the expression of HSPGs at their cell surface. The notion that an LRP ligand binds also to cell surface HSPGs is not unique for C4BP, as several other LRP ligands such as thrombospondin, factor VIII and activated factor IX also have been described to interact with HSPGs<sup>32;33;44</sup>. The mechanism that underlies the cooperation between LRP and HSPGs is unclear. It is assumed that HSPGs first bind the bulk of ligands in order to increase their local concentration. In a second step, these ligands are then transported to LRP that facilitates internalization. Since LRP is known to participate in heterologous receptor complexes<sup>30;45;46</sup>, it is tempting to speculate that part of the LRP and HSPG molecules combine into the same receptor complex. Alternatively, the possibility exists that both receptors meet during dynamic lateral movements at the cell surface. One other issue that is worth mentioning relates to the observations that HSPGs have the capacity to internalize proteins themselves<sup>34</sup>. Since there is considerable residual uptake of C4BP in LRP-deficient cells, it is conceivable that HSPGs are also directly involved in the internalization of C4BP.

We have previously shown that LRP contributes to the initial phase of C4BP clearance<sup>21</sup>. Since in the human situation (but not in mice) the C4BP $\alpha\beta$  isoform circulates in complex with anticoagulant protein S<sup>2;12</sup>, it seems reasonable to assume that C4BP may facilitate clearance of protein S via LRP. Such a facilitation requires formation of trimolecular complexes. Indeed, in two experimental approaches we demonstrated the possibility of the formation of the LRP/C4BP/protein S complex. In this regard it is important to note that protein S itself is unable to interact with LRP<sup>21</sup>. Supporting evidence for the view that

C4BP bridges protein S and LRP is obtained by *in vivo* survival experiments in mice, which showed that the half-life time of C4BP is roughly identical to that of C4BP-bound protein S (Denis and Lenting, unpublished observation).

C4BP interacts with several other proteins as well, such as amyloid P component<sup>17</sup>, factor VIII<sup>18</sup> and bacterial proteins<sup>19</sup>. Therefore, C4BP may function as a chaperone for these proteins to mediate their interaction with LRP. Interestingly, LRP may be involved in the binding of bacteria via the interaction of bacterial surface proteins with C4BP, thereby serving as an entry receptor for pathogenic bacteria.

Taken together, C4BP binds to LRP via positively charged residues that are also involved in binding of C4b and heparin. We demonstrated that HSPGs are involved in the primary binding of C4BP to the cellular surface. Docking of C4BP is followed by an LRP-dependent internalization process. LRP is also able to indirectly bind protein S via C4BP that serves as a physical bridge, suggesting that LRP can also mediate the clearance of protein S, and probably, other C4BP-binding proteins.

## Acknowledgements

This study was supported by a grant from the Dutch Organization of Scientific Research ZonMW (no. 902-26-236) to P.J.L.

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# chapter 6

## General Discussion

Eureka!  
(*Archimedes*)

Receptors have long been thought to interact with a single ligand, but current insights have recognized the existence of receptors that are able to bind multiple ligands<sup>1;2</sup>. One well-known group of these so-called multiligand receptors is the low-density lipoprotein (LDL)-receptor family. At present, 12 members are known, which show structural homology<sup>3-5</sup>. In addition to their multiligand nature, they also display multifunctionality and are expressed in a broad spectrum of cell types. The intriguing aspect in this regard is that despite their versatile nature, these receptors distinguish themselves from each other by distinct cell-expression patterns and cell-specific functions.

### **Novel Aspects of LDL Receptor-Related Protein (LRP) Biology**

Based on its structural homology to the LDL receptor and its capacity to bind lipoproteins, LRP has initially been considered as a second receptor involved in lipoprotein metabolism<sup>6;7</sup>. Indeed, the LDL receptor and LRP act in concert in the clearance of LDL particles<sup>8</sup>. However, it soon became clear that LRP functions as a multiligand receptor, because more and more ligands were identified in time<sup>9;10</sup>. Most of these ligands are internalized upon binding to LRP and subsequently degraded in lysosomal compartments<sup>3;11;12</sup>. LRP not only targets ligands to lysosomes, but also participates in other transport pathways, like transcytosis<sup>13</sup>, nuclear targeting<sup>14</sup> and phagocytosis<sup>15;16</sup>. A number of ligands have been found to modulate cellular properties through their interaction with LRP. In addition, LRP can initiate cellular signaling pathways<sup>16-18</sup>, or can influence signaling by other receptors, thereby functioning as a co-receptor<sup>19-21</sup>.

In the present thesis, various aspects of LRP function in relation to its cellular environment are described.

#### **LRP and its regulation of $\beta$ 2-integrin function**

Another example of multiligand receptors is the  $\beta$ 2-integrin family. These integrins are exclusively expressed by leukocytes and have a role in mediating cell adhesion to different surfaces<sup>22;23</sup>. Their main counter receptors are the intercellular adhesion molecules (ICAMs) present at the endothelium<sup>24</sup>, but they also interact with several other ligands, like fibrinogen<sup>25</sup>, glycoprotein Ib $\alpha$ <sup>26</sup>, factor X<sup>27</sup> and neutrophil inhibitory factor (NIF)<sup>28</sup>.

We demonstrated that LRP is able to directly bind to  $\beta$ 2-integrins at the cell surface of blood-derived polymorphonuclear cells (PMNs) and monocytes or the monocytic cell line U937 (chapter 2). This interaction is at least mediated by the extracellular domains of LRP and the  $\alpha$ -subunits of  $\beta$ 2-integrins. Evidence for this

conclusion includes 1) granulocytes, monocytes and U937 cells bind to immobilized cluster II and IV, which are part of the extracellular chain of LRP, 2) the LRP-antagonist receptor-associated protein (RAP) inhibits this interaction, and 3) adhesion of U937 cells to immobilized cluster II and IV is inhibited by antibodies against the I-domains of the  $\alpha$ -subunits, which are located at the extracellular domain. Whether the intracellular domains also contribute to complex formation remains to be investigated.

Binding of U937 cells to cluster II and IV can be inhibited by antibodies directed against the I-domains of the  $\alpha_L$ ,  $\alpha_M$  and  $\alpha_X$  subunits. Thus, LRP is able to interact with each of these  $\alpha$ -subunits at the cell surface. The importance of the I-domains in LRP-binding is in line with previous findings that show that other ligands interact with this region as well<sup>29-31</sup>. However, adhesion is also decreased when an antibody against the  $\beta$ -subunit is used. This may suggest that LRP is also able to bind to the  $\beta$ -subunit. Indeed, some ligands are known to directly interact with the  $\beta$ -subunit<sup>32</sup>. Alternatively, the anti- $\beta$ 2-subunit antibody may prevent binding of LRP to the  $\alpha$ -subunit via sterical hindrance, or by disrupting the interaction between the  $\alpha$ - and  $\beta$ -subunits. The  $\beta$ -subunit has been shown to be involved in the proper conformation and ligand-binding capacity of the  $\alpha$ -subunit<sup>33-37</sup>. By making use of the purified  $\beta$ -subunit, one may be able to discriminate between these proposed mechanisms.

Proper functioning of  $\beta$ 2-integrins requires their clustering upon immunological activation<sup>38</sup>. The redistribution of  $\beta$ 2-integrins is a process that depends on the liberation of this receptor from the actin cytoskeleton<sup>39</sup>. This process is influenced by a number of intracellular components, such as calpain<sup>40</sup>, activated phospholipase A2<sup>41</sup>, the small GTPase RAP1<sup>42</sup> and its regulator RAPL<sup>43</sup>. Whether or not LRP is involved in this intracellular process is unclear. However, the cytoplasmic domains of both LRP and  $\beta$ 2-integrins contain a number of phosphorylation sites that may affect interactions with adaptor proteins<sup>17;44</sup>.

Redistribution leads to the translocation of  $\beta$ 2-integrins to lipid rafts<sup>45</sup>. Our confocal microscopy and immunoprecipitation experiments clearly demonstrate that LRP is colocalized with  $\beta$ 2-integrins in these clusters. Interestingly, clustering is completely abolished in U937 cells lacking LRP. This suggests that the presence of LRP is necessary for the clustering and, thus, functioning of the  $\beta$ 2-integrins. Indeed, adhesion of LRP-negative U937 cells to endothelial cells is significantly reduced compared to LRP-positive U937 cells. This functional liaison between LRP and  $\beta$ 2-integrins opens the possibility that LRP may also mediate (pathological) processes in which leukocytes are involved. For example, LRP may contribute to  $\beta$ 2-integrin-dependent migration of leukocytes to the subendothelial space, and as such facilitate the development of atherosclerotic lesions. LRP may also control  $\beta$ 2-integrin-dependent phagocytosis in macrophages. Future

studies should be designed that address the issue in which processes LRP plays a regulatory role.

### **LRP and its expression in lymphocytes**

Since LRP plays an important role in monocyte and PMN adhesion, it was of interest to investigate the contribution of LRP to the function in other leukocytes as well. Whereas the expression and function of LRP in monocytes and dendritic cells have been well characterized<sup>15;16;46-48</sup>, LRP expression patterns and functions in other leukocytes have not been fully understood yet.

In chapter 3, we described the presence of both LRP mRNA and protein in different lymphocyte subsets: T-lymphocytes, B-lymphocytes and natural killer (NK) cells. Interestingly, a different localization of LRP in T-lymphocytes compared to B-lymphocytes and NK cells is observed. Whereas LRP is present at the cell surface of NK cells and B-lymphocytes, only a small amount of T-lymphocytes have LRP at their surface. Thus, LRP appears to be located in intracellular storage compartments in T-lymphocytes.

We speculate that LRP may be recruited to the cell membrane upon activation in T-lymphocytes. Stimulation of T-lymphocytes by interleukin-2 (IL-2) and phytohaemagglutinin (PHA) indeed increases the number of cells positive for cell surface LRP from 2% to 12% of the T-lymphocytes after 48h. However, this increase is transient and not immediate, making it unlikely that LRP is redistributed from intracellular storage pools to the cell surface. In a second approach, we showed that fast upregulation of LRP takes place in a mixed-lymphocyte reaction (MLR) in 60-70% of the T-lymphocytes. This suggests that LRP is promptly translocated from storage pools to the cell surface in these cells. A fast translocation of LRP to the cell surface is also observed in insulin-treated adipocytes<sup>49</sup> and NGF-treated neurons<sup>50</sup>. It is important to note that in resting neurons and adipocytes LRP is also present at the cell surface, whereas in T-lymphocytes the total LRP pool is maintained within the cell.

It appeared that allogenic monocytes were able to induce this observed increase in LRP cell surface expression, whereas mixing T-lymphocytes from 2 donors did not increase cell surface expression. This may point to a role for LRP in conditions in which antigen presenting cells (APCs) trigger T-lymphocyte activation, like in graft-versus-host disease (GVHD)<sup>51</sup>. Many receptors that play a role in the interaction with APCs are present at the cell surface of T-lymphocytes<sup>52-57</sup>. Interestingly, one of these cell surface receptors, CTLA-4, is also recruited to the cell membrane in a mixed lymphocyte reaction (MLR)<sup>55</sup>. It is possible that CTLA-4 and LRP are stored in similar storage pools. Another receptor that is important for T-lymphocyte activation is  $\alpha_L\beta_2$ <sup>57</sup>. Since LRP and  $\alpha_L\beta_2$  are functionally interlinked in monocytes, it deserves further study to de-

termine if such connection also exists in T-lymphocytes. If so, then LRP may serve as a novel target to interfere with immune disorders such as GVHD. It should be noted that such studies also should focus on the elucidation of the intracellular signaling pathways that force the translocation of LRP to the cell surface.

### **LRP and its function as brain endothelial clearance receptor for amyloid $\beta$ (A $\beta$ ) peptides**

LRP is a key component in the pathogenesis of Alzheimer's disease (AD). A genetic polymorphism in the *LRP* gene itself is associated with late-onset AD<sup>58-61</sup>. In addition, the LRP ligands amyloid precursor protein (APP), Apolipoprotein E (ApoE) and  $\alpha_2$ -macroglobulin ( $\alpha_2$ M) are also genetically linked to AD<sup>10</sup>. LRP mediates the endocytosis of APP in neurons, leading to increased A $\beta$  production<sup>62-64</sup>. Although LRP can mediate the re-uptake of these A $\beta$  peptides in neurons<sup>65-67</sup>, this mechanism is unlikely to contribute to A $\beta$  clearance *in vivo*, but rather leads to the formation of amyloid plaques<sup>68</sup>. Thus, an alternative pathway should exist that results in the removal of A $\beta$  peptides from the brain to the circulation. In chapter 4, the contribution of brain endothelial cell LRP in the clearance of A $\beta$  peptides is described.

A $\beta$  peptides are known to bind to LRP indirectly via complex formation with ApoE or  $\alpha_2$ M<sup>65-67</sup>. However, our study clearly shows that A $\beta$  peptides also directly interact with LRP clusters. The affinity of A $\beta$ 40 for LRP is higher than of A $\beta$ 42, which contains 2 extra hydrophobic residues, and of the A $\beta$ 40 isoform harboring the Dutch/Iowa mutation (DIA $\beta$ 40). This increased affinity correlates with the  $\beta$ -sheet content of A $\beta$ -peptides: A $\beta$ 40 contains only 25%  $\beta$ -sheet, and A $\beta$ 42 and DIA $\beta$ 40 75% and 80%, respectively. LRP mediates binding and internalization of A $\beta$  peptides by brain endothelial cells. However, this uptake is not followed by degradation. This suggests that LRP may mediate the transcytosis of A $\beta$  over the blood-brain barrier (BBB). *In vivo* experiments showed that A $\beta$ 40 is indeed more rapidly cleared from brain tissue than the low-affinity binders A $\beta$ 42 or DIA $\beta$ 40. Remarkably, the presence of high levels of A $\beta$  peptides decreases their efflux across the BBB, caused by an induction of LRP degradation in the proteasome pathway. The relevance of this observation *in vivo* has been demonstrated by studying the phenotype of RAP-null mice (which show severe LRP deficiency) crossed with APP-overexpressing mice<sup>69</sup>. In these mice A $\beta$  peptides accumulate in the interstitial fluid of the brain. This implicates that A $\beta$  peptides are inefficiently cleared across the BBB. Indeed, A $\beta$  peptide clearance is decreased significantly compared to wild-type mice in mice expressing either the low-clearance mutant DIA $\beta$ 40 or overexpressing APP. Interestingly, these latter mice show reduced LRP levels in brain, suggesting that an excess of A $\beta$  availability

also downregulates LRP in brain endothelial cells *in vivo*. In AD patients carrying the DIA $\beta$  mutant also an elevated A $\beta$  deposition is observed<sup>70</sup>. Studying LRP levels of brain endothelial cells from these patients reveals a reduction in LRP expression.

Receptor for advanced glycation end products (RAGE) is another A $\beta$  receptor present at brain endothelium. In contrast to LRP, it mediates continuous influx of A $\beta$  from blood to the brain<sup>71</sup>. An imbalance between influx and efflux may increase A $\beta$  accumulation in brain<sup>72</sup>. It has been demonstrated that A $\beta$ -peptide-binding agents present in blood shift the transport balance towards the blood, thereby lowering the A $\beta$  concentration in brain tissue. Therefore, sequestering of A $\beta$  peptides via LRP may be a novel therapeutic target in the development of AD.

### **LRP and its role in C4b-binding protein (C4BP) clearance**

Mediating internalization of soluble proteins is the classical function of LRP<sup>3;11;12</sup>. Recently, we identified C4BP $\alpha$  as a new ligand for LRP<sup>73</sup>. C4BP is a component of both complement system and coagulation and has been found to bind several proteins, such as C4b, amyloid P component, bacterial components and protein S<sup>74-78</sup>. Heparin has been described to bind to C4BP as well<sup>79;80</sup> and to inhibit the interaction between C4BP and LRP<sup>73</sup>. This would suggest that positively charged residues in C4BP are involved in the interaction with LRP. Analyzing C4BP mutants in which positive amino acids are mutated into glutamine resulted in the finding that a triple mutant C4BP showed a severe reduction in LRP binding (chapter 5). In addition, the triple mutant is less efficiently degraded in CHO cells containing LRP. The observed residual binding suggests that other interactions are also involved. Indeed, experiments using high salt concentrations reveal that these high concentrations are not able to completely inhibit C4BP/LRP binding, suggesting that hydrophobic interactions may also play a role. In this respect, the binding between activated factor IX and LRP has been shown to be partially dependent on hydrophobic interactions<sup>81</sup>.

The importance of positively charged amino acids in LRP binding would suggest that heparan sulfate proteoglycans (HSPGs) also play a role in the binding of C4BP. CHO cells lacking functional HSPGs show strongly reduced binding of C4BP. Also for other LRP ligands it has been demonstrated that these HSPGs serve as a primary docking site<sup>82-84</sup>, thereby increasing their local concentration. Subsequently, ligands can efficiently be transferred to LRP, which mediate their internalization.

The mutated positively charged amino acids in the triple mutant have also been demonstrated to be important in binding of other C4BP-binding proteins<sup>76;85</sup>, suggesting that these proteins may interfere with the interaction between C4BP and LRP. Alternatively, because of C4BP's multivalency, a trimolecu-

lar complex can occur consisting of LRP, C4BP and the C4BP-binding protein S. Protein S binds to the  $\beta$ -chain of C4BP<sup>77;86</sup>. We demonstrated that LRP is still able to bind C4BP in complex with protein S. Since protein S does not bind to LRP itself<sup>73</sup>, C4BP is proposed to link protein S and LRP. Thus, LRP may indirectly regulate the plasma concentration of these C4BP-binding proteins.

## The Versatility of LRP

In chapters 2-5 we highlighted 4 distinct aspects of the multifunctional protein LRP. This receptor can exert many diverse functions in various cell types. Moreover, the function of LRP in one cell type may not be restricted to a single process. Thus, the question rises as to which regulatory mechanisms may specify LRP function. In the next paragraphs, we discuss to what extent the microenvironment plays a role in determining the specificity of LRP.

### LRP's ligands considered

LRP is a receptor that is able to recognize multiple structurally unrelated ligands<sup>9;10</sup>. These ligands can prompt various cellular events, like endocytosis<sup>3;11;12</sup>, cell adhesion and growth<sup>48;87</sup>, and neuronal transmission<sup>88;89</sup>. The spatial localization of LRP ligands may be one mode in which the function of LRP is specified. Some ligands show a highly cell-restricted expression pattern, like  $\beta$ 2-integrins that are only present on leukocytes<sup>22</sup>. In this regard, we showed that LRP directly binds to the  $\alpha$ -subunit of  $\beta$ 2-integrins (chapter 2). Interestingly, in CHO cells LRP mediates the uptake of soluble I-domain of  $\alpha_M$  (data not shown), whereas it does not regulate surface expression of  $\beta$ 2-integrins in monocytes (chapter 2). It cannot be excluded that delivery to the lysosomal degradation pathway is limited to soluble ligands (refs 12;90-92 and chapter 5) More likely, the endocytic process is cell-type-dependent in cells (see 'Cell-restricted functions of LRP').

The function of LRP may also be restricted by a temporarily availability of its ligands. One clear example hereof is coagulation factor IX. In its naïve state, factor IX is unable to ligate to LRP. However, factor XIa-mediated activation of this zymogen exposes a binding site for LRP<sup>82</sup>. Another profound case of temporarily present agents are proteins derived from bacteria and viruses during an inflammatory event, like HIV tat protein and pseudomonas exotoxin A<sup>93-96</sup>. To this end, LRP may serve as an entry receptor, thereby contributing to the hazardous properties of these compounds. At the same time, it is of interest to consider the presence of soluble LRP, which has the potential to act as a first defense barrier, preventing cellular penetration of these infective agents<sup>97</sup>.

Third, the structure of a ligand may allow dimerization or multimerization of LRP. Backsai et al<sup>88</sup> proposed that LRP/ $\alpha_2$ M-mediated NMDA-induced calcium influx may be due to LRP dimerization, since LRP-activating antibodies, but not their Fab-fragments caused elevated intracellular calcium levels.  $\alpha_2$ M is a multi-valent protein, which may be able to cluster LRP, and thereby inducing signaling. C4BP $\alpha$  can be internalized via LRP (ref 73 and chapter 5), but also shows other cellular effects. For instance, it stimulates B-lymphocyte proliferation via CD40<sup>98</sup>. Since LRP is also present at B-lymphocytes (chapter 3), it would be of interest to study the involvement of LRP in this process. Moreover, because of its multivalency, C4BP may be able to multimerize LRP to evoke cellular signaling, or it may cause the formation of heterologous receptor complexes that consist of C4BP-binding receptors.

Since LRP's ligands are structurally dissimilar, but have the potential to induce similar physiological processes, the structure of these proteins may not be a major determinant of LRP's function. Thus, a model in which the spatial and temporal occurrence of ligands may be preferable over their structure for influencing the role of LRP under specific circumstances.

### **Cell-restricted functions of LRP**

Hepatocytes are identified as master endocytic cells that mediate the clearance of various plasma proteins in an LRP-mediated fashion<sup>99-101</sup>. Since hepatocytes are located directly adjacent to the blood stream, hepatic LRP can efficiently internalize its ligands, thereby regulating their plasma concentration. Although studies on the presence of LRP in endothelial cells are inconclusive, LRP has been described to be expressed by brain endothelium<sup>102</sup>, another cell type that is constitutively exposed to blood. LRP is involved in rapid clearance of A $\beta$  peptides from the brain in these cells (chapter 4). Whereas other cell types, like adipocytes<sup>49;103;104</sup> and fibroblasts (chapter 5), are also competent to internalize LRP ligands, other cells are mainly involved in distinct cellular events<sup>10</sup>. One striking example of cell-restricted function is illustrated in chapter 4. Although neurons and brain endothelium both express LRP<sup>4;5;102;105</sup>, LRP-mediated transcytosis of A $\beta$  peptides is predominantly limited to brain endothelium.

In neurons, LRP shows various unrelated functions. Among them are modulation of N-methyl-D-aspartate (NMDA) receptor-mediated calcium signaling<sup>88</sup>, processing of APP<sup>62</sup> and stimulation of neurite outgrowth<sup>106</sup>. These functions may become apparent under distinct conditions of activation of neuronal cells. For example, modulation of NMDA receptor signaling may occur during synaptic transmission of an action potential<sup>89</sup>, while under conditions of neuronal damage LRP may be particularly involved in neurite outgrowth<sup>106;107</sup>.

LRP displays also multiple functions in monocytes. Activated monocytes adhere to the endothelium in an LRP-dependent mode (chapter 2). Upon subsequent differentiation into macrophages, LRP expression is increased dramatically<sup>108</sup>. In the macrophage, LRP contributes to phagocytosis<sup>15;16</sup> and to 12/15 lipoxygenase-mediated oxidation of LDL<sup>46</sup>. In contrast to neurons, the function of LRP in monocytes/macrophages is thus clearly spatially and temporarily determined. However, precise mechanisms that regulate LRP function are presently unknown. In both neurons and monocytes, it seems conceivable that cofactors that are (temporarily) active or present during these processes influence LRP function.

### **Receptor cross-talk**

LRP is able to efficiently act in concert with other proteins present at the cell surface. HSPGs are a group of receptors abundantly present at the cell surface<sup>109</sup>. Heparin-binding LRP ligands (*e.g.* factor VIII and tissue-factor pathway inhibitor (TFPI)) are efficiently captured from the plasma to the plasma membrane, resulting in a high concentration pool at the cell surface. Ligands can then be directed to LRP for endocytosis or other cellular events<sup>84;110;111</sup>. Chapter 5 illustrates that endocytosis of binding of C4BP (a protein with heparin-binding sites<sup>79;112</sup>) is also a multistep process, which first involves binding to HSPGs and subsequently binding to and endocytosis by LRP. Whether LRP and HSPGs co-exist in a heterologous complex or modulate each others function is unclear so far.

LRP is described to be involved in the modification of cellular signaling events by other receptors. For this function as coreceptor, LRP may indirectly interact with other cell surface receptors via intracellular adaptor proteins. For instance, LRP is able to cooperate with the platelet-derived growth factor (PDGF) receptor<sup>19;20</sup>. Stimulation of the PDGF receptor results in phosphorylation of the LRP cytoplasmic tail<sup>19;20</sup>, enabling its conjugation with the adaptor protein Shc, which also binds to the PDGF receptor. Via this bridge LRP is able to integrate with the PDGF-induced signaling in order to minimize the cellular responses provoked by PDGF.

Another example in which LRP limits signaling raised by other receptors is that of the NMDA receptor. Long-term exposure of  $\alpha_2$ M to LRP on neurons downregulates NMDA receptor expression and thereby diminishes calcium influx via the NMDA receptor<sup>21</sup>. LRP is involved in short-term stimulation of calcium influx as well<sup>88</sup>. To accomplish this receptor cross-talk, LRP and NMDA receptor are inter-linked via the intracellular adaptor protein PSD-95<sup>17;89</sup>.

Concerning leukocytes, we demonstrated that LRP regulates the clustering of  $\beta$ 2-integrins (chapter 2). Both receptors may form a complex at the cell surface

via an interaction between their extracellular domains. To exert a regulatory function, LRP may interact with  $\beta$ 2-integrins via their cytoplasmic tails as well. To this end, LRP may be directly involved in the redistribution of  $\beta$ 2-integrins. The process of relocation depends on the detachment of  $\beta$ 2-integrins from the actin cytoskeleton<sup>33;39;113;114</sup>. How or whether LRP is involved in this process, is still unclear. However, LRP has been recently described to be required for actin reorganization during the engulfment of apoptotic cells<sup>115</sup>. Both receptors contain several putative phosphorylation sites that may affect interactions with adaptor proteins<sup>16-18;44;116</sup>. Therefore, these adaptor proteins may serve to link the intracellular domains of  $\beta$ 2-integrins and LRP, thereby enabling LRP to modulate  $\beta$ 2-integrin clustering, probably via actin remodeling.

Besides an indirect interaction via adaptor proteins, LRP is also able to directly interact with transmembrane receptors. LRP mediates the internalization of APP via a direct interaction<sup>117</sup>. It is noteworthy that neuronal LRP is only able to efficiently internalize APP, but not A $\beta$  peptides<sup>102</sup>. Apparently, motifs in APP's intracellular tail are required for endocytosis by LRP, possibly via adaptor proteins. Interestingly, LRP's intracellular tail is connected with the intracellular domain of APP via the adaptor protein Fe65<sup>118;119</sup>. Whether or not this interaction is required for endocytosis or signaling has still to be investigated. Chapter 4 reports that brain endothelial cells are capable to internalize A $\beta$ . However, it is still unclear whether motifs in A $\beta$  peptides or the endocytosis machinery of these cells is responsible for this process.

Taken together, the microenvironment of LRP, and in particular the presence of other membrane receptors, may define LRP functions in a specific cell-type.

### **Subcellular localization-dependent function of LRP**

Initial studies on LRP showed that LRP is localized in clathrin-coated pits at the cell surface for its function as endocytic receptor. LRP interacts with adaptor protein-2 in the clathrin complex. Motifs in the intracellular tail are specifically involved in endocytosis. Whereas for the LDL receptor the NPXY motif is required for efficient endocytosis<sup>120</sup>, the main endocytosis motifs for LRP are a YXXL and a di-leucine motif<sup>121</sup>. In chapter 5 we described that LRP is required for the endocytosis of C4BP. It seems conceivable that clathrin-coated pits are also involved in C4BP endocytosis.

Apart from the clathrin-coated pits, LRP has also been detected in lipid rafts<sup>19</sup>. It is tempting to speculate that LRP is recruited to lipid rafts in its function as signaling receptor<sup>122</sup>. LRP functions as a modulator of PDGF receptor signaling<sup>20</sup>. To do so, LRP and PDGF receptor are localized in caveolae<sup>19</sup>. In addition, LRP is described to be present in rafts in complex with  $\beta$ 2-integrins in activated monocytes (chapter 2). We found that LRP does not downregulate cell surface expres-

sion of  $\beta$ 2-integrins, suggesting that LRP is not involved in endocytosis in this interaction in rafts in these cells. However, LRP localized in lipid rafts can also mediate internalization of ligands, as described for APP, of which the proper internalization and processing is dependent on its localization in these cell membrane domains<sup>123</sup>. Probably caveolae are involved, since this lipid raft subtype is known to play an important role in non-clathrin-mediated endocytosis<sup>124</sup>.

Localization of LRP at the cell surface may be directed by the interaction between cytoplasmic tail and intracellular adaptor proteins. This is illustrated by Marzolo et al<sup>125</sup> who demonstrated that the NPXY motif of LRP is responsible for its basolateral sorting in MDCK cells. In other polarized cells, like neurons, LRP localization appears to be restricted to only one plasma membrane domain<sup>105;126</sup>. In brain endothelial cells, LRP is supposed to be present at their basolateral membrane and to mediate clearance of A $\beta$  peptides (chapter 4 and ref 102), whereas RAGE is involved in internalization of A $\beta$  peptides at the apical membrane<sup>71</sup>.

Thus, subcellular localization of LRP appears to be an important factor in the regulation of its function. The presence of coreceptors and intracellular adaptor proteins, and LRP's association with the cytoskeleton may be key factors in localizing LRP in cell membrane domains. Due to this spatial dissection, LRP is exposed to a selective set of ligands, which may specify LRP's function. In addition, function can be determined by the phosphorylation state of LRP's intracellular tail. Serine/threonine phosphorylation leads to an increase in endocytosis rate<sup>127;128</sup>. In contrast, cellular signaling events may be dependent on tyrosine phosphorylation<sup>16-18;129</sup>. It would be interesting to study which nature of phosphorylation is preferred at a specific site at the cell surface.

### **Structural requirements of LRP in relation to its function**

We demonstrated that leukocytes equally adhere to immobilized cluster II and IV. In this respect, leukocytes are similar to most other LRP ligands, which show similar affinity for isolated recombinant clusters II and IV<sup>130</sup>. One of these ligands is the tPA/PAI-1 complex. Interestingly, studies using minireceptors only containing cluster II and IV revealed that the tPA/PAI-1 complex prefers binding to cluster II. Possibly, the covalent attachment to the transmembrane domain may limit ligand access to the minireceptor containing cluster IV<sup>131</sup>. The ligand binding repeats in a cluster show similarities, but display differential ligand recognition. Primary conformation and spatial arrangements between the individual repeats are thought to ensure ligand specificity<sup>1</sup>. Further molecular dissection of the clusters showed that ligands prefer binding to certain successive repeats, which may overlap with repeats that bind other ligands<sup>132</sup>. Future studies may provide evidence whether ligand binding to a specific cluster or repeat

evokes a particular LRP-mediated cellular function. In this aspect, it is of interest to mention that RAP admittedly is able to bind to LRP with high affinity<sup>133</sup> and can be internalized via LRP, but many studies demonstrated that RAP is unable to stimulate cellular signaling pathways. Thus, structural information within RAP may determine its effect on LRP. Alternatively, RAP may change the conformation of LRP, thereby making it impossible to transduce a signal. Regarding LRP conformation, it has been generally postulated that one single RAP molecule can interact with cluster II, III and IV of LRP simultaneously. It is conceivable that other ligands may induce a distinct conformational change in LRP, thereby transmitting a particular signal to LRP's intracellular tail and allowing binding of specific adaptor molecules.

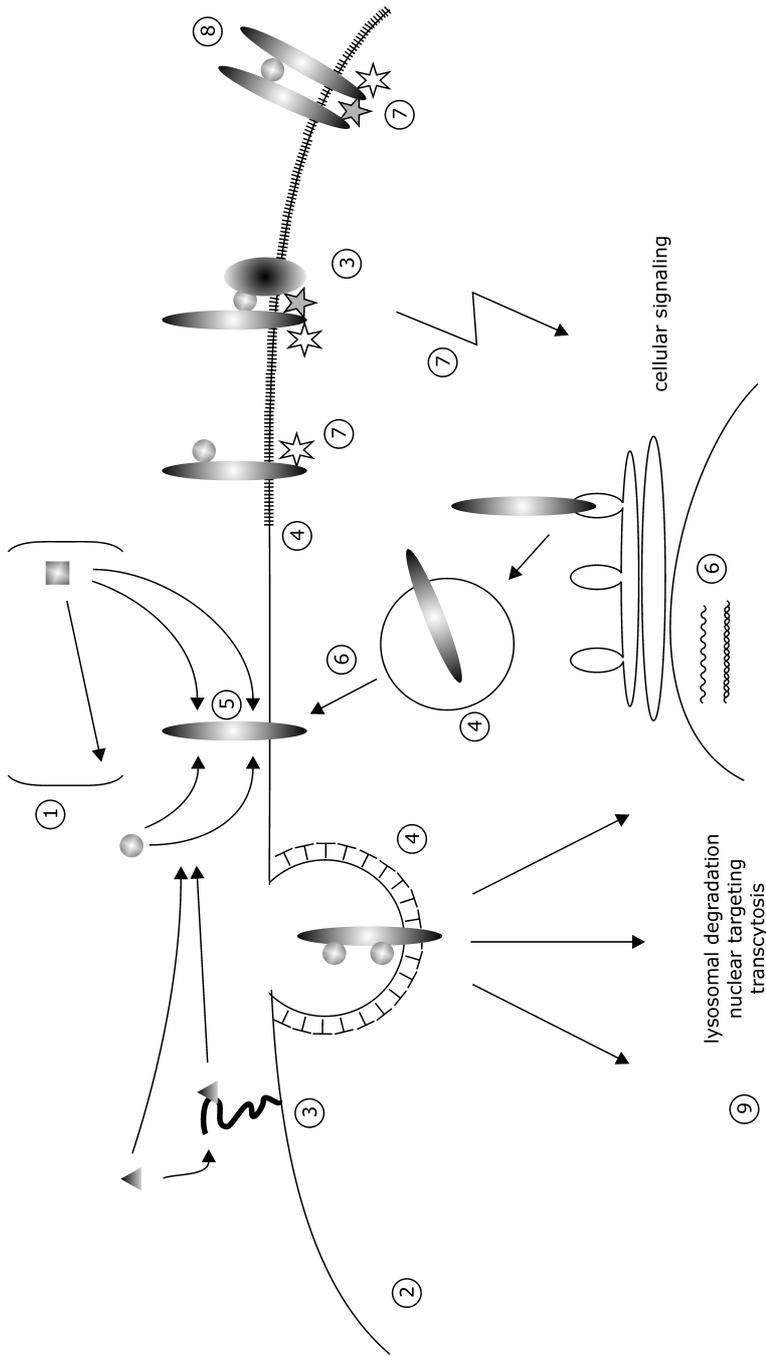
A tool to study apparently distinct functions of the clusters are LRP minireceptors that contain only one cluster (mainly II and IV). Though, the difference between the distance between cluster II and the cell membrane in full-length and truncated LRP may be a disadvantage of this method. Probably, the spatial occurrence of cluster II in full-length LRP may specify the function of LRP raised by binding of a ligand to this cluster compared to binding to cluster IV<sup>131;134;135</sup>.

### **Regulation of LRP expression**

In most cell types LRP is constitutively expressed at the plasma membrane, where its presence is required to exert its function. Under certain conditions LRP's cell surface expression is regulated by either *de novo* synthesis or translocation from intracellular storage pools.

Although exact mechanisms on how transcription is regulated are unknown<sup>136</sup>, NGF, vascular endothelial growth factor, transforming growth factor (TGF)- $\beta$ 1 and TGF- $\beta$ 2 are described to induce LRP mRNA expression<sup>137;138</sup>. A rapid induction of LRP protein expression at the cell surface can be accomplished by translocation of intracellularly stored LRP, as has been demonstrated for insulin- or NGF-stimulated cells<sup>49;137</sup>. This is further exemplified in chapter 3, which describes that LRP protein is rapidly recruited to the cell membrane in T-lymphocytes. These cells are the only cells described that maintain the total LRP pool intracellularly, in contrast to other cell types which show both cell surface and intracellular localization of LRP. The underlying mechanisms on how T-lymphocytes regulate this localization are unclear. In these examples signaling induced by another receptor may regulate LRP expression. In chapter 4 we described that the LRP ligand A $\beta$  peptide directly downregulates the expression of LRP by targeting it to the proteasomal degradation pathway.

These studies indicate that the function and activity of LRP may be temporally controlled by regulating its cell surface expression.



**Figure 1. Regulation of LRP function.** The versatility of LRP may be determined by its microenvironment. 1, occurrence and structure of ligands; 2, cellular expression patterns; 3, receptor cross-talk; 4, subcellular localization; 5, structure of LRP; 6, regulation of mRNA and protein expression; 7, intracellular adaptor proteins and cellular signaling; 8, receptor dimerization; 9, intracellular transport pathways.

## Summary and Future Prospects

In summary, the multifunctionality and the broad cellular expression of LRP prompted us to address the question whether its microenvironment play a pivotal role in directing cell-restricted functions. Information present in ligands and LRP structure may determine cellular events induced by receptor-ligand interaction. In addition, accessory proteins located either at the cell surface (coreceptors) or intracellularly (adaptor proteins) contribute to specifying functions of LRP at different steps of cellular responses (*Fig 1*).

Future research may be focused to consider details in ligand-receptor interactions in respect to components present in the environment, and how these interactions may influence LRP's functions in respect to intracellular transport pathways and cellular signaling. A proteomic approach may aid to identify LRP-binding proteins under specific (patho-)physiological conditions.

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## **Samenvatting in het Nederlands**

De meest fundamentele ideeën van de wetenschap zijn  
in wezen eenvoudig en kunnen in de regel worden uitgedrukt  
in een taal die voor iedereen begrijpelijk is  
*(Albert Einstein)*

## Inleiding

Cellen communiceren met hun omgeving via bepaalde eiwitten, zogenaamde receptoren, die zich bevinden op het oppervlak van de cel (= celmembraan). Vaak geldt dat receptoren alleen één soort eiwit kunnen herkennen aan het oppervlak van de cel. Een eiwit dat bindt aan zo'n receptor wordt ligand genoemd. Echter, er zijn ook receptoren bekend die meerdere soorten liganden kunnen binden. Dit is de groep van de zogenaamde 'multiligand receptoren'.

Een groep van aan elkaar verwante receptoren is de low-density lipoproteïne (LDL) receptor familie. De familie bestaat uit 12 leden die qua structuur veel op elkaar lijken. Een tweede overeenkomst is dat ze allemaal in staat zijn om meerdere soorten liganden te binden. De LDL receptoren zijn dus multiligand receptoren. De receptoren hebben ook verschillen ten opzichte van elkaar. Zo zijn enkele leden groter dan de rest. Bovendien komen ze niet allemaal in dezelfde celtypen voor en binden ze ook niet allemaal dezelfde liganden.

Het onderzoek zoals beschreven in het proefschrift is gericht op één receptor uit de familie, namelijk LDL receptor-related protein (LRP). Zoals de naam al doet vermoeden, lijkt deze receptor veel op de LDL receptor. Grote verschillen zijn dat LRP ruim 4x zo groot is en veel meer soorten liganden kan binden. Voorbeelden van liganden zijn stollingsfactoren, lipoproteïnen en groeifactoren. De liganden vertonen soms, maar meestal geen structurele overeenkomsten.

Het belang van LRP voor het leven is gebleken toen er transgene muizen werden gemaakt waarin LRP afwezig is. De muizen sterven vroeg in de embryonale ontwikkeling. De functie van LRP is net als zijn liganden erg divers. Voor de diverse functies is de staart van LRP die zich aan de binnenkant van de cel bevindt, erg belangrijk. Hieraan kunnen andere eiwitten binden die de functie van LRP kunnen bepalen. Een bekende functie van LRP is de opname van liganden uit het plasma of van het celoppervlak via het gedeelte van LRP dat uit de cel steekt. Dit proces heet endocytose en leidt uiteindelijk tot de afbraak van de liganden in zogenaamde lysosomen ('mini-maagjes' van de cel). Daarnaast kan LRP ervoor zorgen dat er een signaal in de cel ontstaat dat uiteindelijk bepaalde effecten teweegbrengt, zoals celgroei- en ontwikkeling. Een bijzondere vorm van signalering is dat LRP de functie van andere receptoren op het celoppervlak kan beïnvloeden.

LRP komt voor in verschillende cellen en weefsels in ons lichaam, zoals lever-, hersen- en vetcellen. Niet alle bovengenoemde functies komen in alle cellen voor. De combinatie van cellen en liganden kan de functie van LRP onder die bijzondere omstandigheden bepalen.

Dit proefschrift beschrijft een aantal nieuwe aspecten van LRP. Er is gekeken naar de functie van LRP in verschillende cellen. Een korte samenvatting van de gevonden resultaten is hieronder beschreven.

## **LRP en de regulatie van de functie van $\beta$ 2-integrinen**

LRP is dus een receptor die veel liganden kan binden. Er zijn ook andere groepen liganden bekend die dezelfde eigenschap hebben. Een voorbeeld zijn de  $\beta$ 2-integrinen. Dit zijn receptoren die op het oppervlak van witte bloedcellen voorkomen. Deze cellen zijn betrokken bij onze afweer.  $\beta$ 2-integrinen zijn onder andere betrokken bij de binding van de witte bloedcellen aan de vaatwand indien de cellen geactiveerd worden door andere onderdelen van het afweersysteem. Hierdoor kunnen de cellen door de vaatwand heen migreren en zo naar de plaats gaan waar ze nodig zijn om bijvoorbeeld de oorzaak van de infectie te bestrijden.

$\beta$ 2-integrinen zijn een complex van 2 eiwitten. Ze bestaan uit een  $\alpha$ -deel en een  $\beta$ 2-deel. Er zijn verschillende  $\alpha$ -delen bekend:  $\alpha_L$ ,  $\alpha_D$ ,  $\alpha_X$  en  $\alpha_B$ . In alle  $\alpha$ -delen is een bepaald gebied aanwezig dat betrokken is in de binding aan andere liganden, het zogenaamde I-domein.

In hoofdstuk 2 hebben we beschreven dat de ligand-bindende delen van LRP, de zogenaamde clusters II en IV, en het I-domein van het  $\alpha$ -deel van de  $\beta$ 2-integrinen direct aan elkaar konden binden. Ook waren LRP en  $\beta$ 2-integrinen elkaars liganden op het oppervlak van een groep witte bloedcellen, de monocyten. Het is gebleken dat LRP en  $\beta$ 2-integrinen aan elkaar bonden in zowel ongestimuleerde als gestimuleerde monocyten.

Het was al bekend dat als monocyten gestimuleerd worden,  $\beta$ 2-integrinen groepjes aan het celoppervlak vormen. Dit is noodzakelijk om efficiënte binding van de monocyten aan de vaatwand te krijgen. Wij hebben gevonden dat in deze groepjes ook LRP aanwezig was. De functie van LRP in deze situatie hebben we bekeken door monocyten te gebruiken waarin LRP niet aanwezig was. Zulke monocyten bleken dan niet meer in staat om de groepjes  $\beta$ 2-integrinen te vormen. Dit had als gevolg dat deze monocyten ook niet meer zo efficiënt aan de vaatwand konden binden.

Uit deze resultaten blijkt dus dat LRP de functie van  $\beta$ 2-integrinen kan beïnvloeden en dat LRP noodzakelijk is voor het goed functioneren van  $\beta$ 2-integrinen. In dit geval is LRP een zogenaamde co-receptor voor  $\beta$ 2-integrinen. Vervolgonderzoek moet uitwijzen hoe LRP precies de  $\beta$ 2-integrinen beïnvloedt.

## LRP en zijn expressie in lymfocyten

LRP komt in veel celtypen voor en in sommige celtypen heeft LRP een unieke functie. Omdat de aanwezigheid van LRP van vitaal belang is, is het noodzakelijk dat de expressie van LRP in celtypen goed wordt onderzocht en dat de (specifieke) functie van LRP in deze cellen opgehelderd wordt. In hoofdstuk 2 hebben we beschreven dat LRP belangrijk is voor de functie van monocyten. LRP komt ook voor in de witte-bloedceltypen granulocyten en dendritische cellen. Het is nog niet duidelijk of LRP ook aanwezig is in T-cellen. Er is nog niets bekend over de expressie in B-cellen en natural killer cellen (NK cellen). Elke celsoort heeft een unieke receptor op zijn celoppervlak waaraan deze cel te herkennen is. Deze heten CD-markers. T-cellen zijn te herkennen aan CD3. Er zijn 2 soorten T-cellen: helper T-cellen (CD4) en cytotoxische T-cellen (CD8). B-cellen hebben CD19 en NK cellen CD16 en CD56.

In hoofdstuk 3 wordt onderzoek naar de aanwezigheid van LRP beschreven in deze celtypen. Eerst werd gekeken naar mRNA. mRNA wordt afgelezen van DNA en mRNA wordt weer afgelezen om uiteindelijk eiwit te vormen. Zowel LRP mRNA als LRP eiwit was aanwezig in de witte-bloedceltypen T-cellen, B-cellen en NK cellen. Toen we keken naar de plaats waar LRP zit in de cellen, dan zagen we iets opmerkelijks. In B-cellen en NK cellen zat LRP op het celoppervlak. Echter, in T-cellen was LRP voornamelijk aan de binnenkant van de cel aanwezig.

Omdat LRP in het algemeen zijn functie aan het oppervlak van de cel heeft, hebben we gekeken of LRP vervoerd kan worden van de binnenkant naar het oppervlakte van de cel. Van andere celtypen is bekend dat dit kan gebeuren door cellen te stimuleren. We hebben gekeken of LRP ook naar buiten komt als T-cellen gestimuleerd worden. T-cellen kunnen op verschillende manieren gestimuleerd worden. Als eerste kan dat door het eiwit interleukine-2, dat is betrokken bij de celgroei die plaats vindt na stimulatie. Na stimulatie met interleukine-2 kwam er na 2 dagen meer LRP op het celoppervlak in 12% van de cellen. Als LRP al in de cel 'klaar ligt', dan zijn 2 dagen vrij lang voordat het op het oppervlak komt. Een andere manier van stimuleren is zoals ook gebeurt bij de afstoot van organen na transplantatie. De T-cellen van de donor kunnen dan cellen aanvallen van de ontvanger. Dit hebben we nagebootst door witte bloedcellen van één persoon samen te voegen met witte bloedcellen van een tweede persoon. Dan zagen we dat bij ongeveer 70% van de T-cellen LRP verschoof van binnen de cel naar de oppervlak in een periode van minder dan een half uur. Toen we verschillende witte bloedcellen zuiverden, zagen we dat als een T-cel van persoon 1 met een monocyten van persoon 2 mixten, dat dan LRP op het oppervlak kwam, maar niet als de gezuiverde T-cellen gemixt werden.

Uit deze resultaten kunnen we nog niet bepalen wat de rol van LRP is. Daarom moet er meer onderzoek gedaan worden om de rol van LRP in de afstoot van donorcellen te bepalen.

## **LRP en de rol in de ziekte van Alzheimer**

De ziekte van Alzheimer kenmerkt zich door afbraak van delen van de hersenen en de hierop volgende uitval van bepaalde hersenfuncties. De oorzaak is dat eiwitten die niet meer de goede structuren hebben worden afgezet en in de hersenen en daar dan de cellen beschadigen.

Op het oppervlak van hersencellen zit een eiwit, genaamd APP, dat door LRP opgenomen kan worden. In de cel wordt het dan geknipt en weer uitgescheiden. Deze uitgescheiden vorm (amyloid  $\beta$ ) kan dan schadelijk zijn als het gaat klonten en zogenaamde plaques gaat vormen. Ter bescherming kan het weer worden opgenomen door de omliggende cellen. Ook bij dit proces is LRP betrokken: het zorgt voor opname van amyloid  $\beta$  door de vaatwandcellen in de hersenen.

In hoofdstuk 4 wordt beschreven hoe LRP de opname van amyloid  $\beta$  op vaatwandcellen reguleert. Het bleek dat de slechte vorm van amyloid  $\beta$  ( $A\beta_{42}$ ) slechter bond aan LRP op de vaatwandcellen dan de goede vorm ( $A\beta_{40}$ ). Daardoor kon er meer slecht  $A\beta_{42}$  in de hersenen komen, wat dan leidde tot een grotere kans op de ziekte van Alzheimer. Amyloid  $\beta$  dat via LRP werd opgenomen, werd aan de andere kant van de vaatwandcellen uitgescheiden en kwam zo het bloed in. Zo kon amyloid  $\beta$  worden afgevoerd en was het niet schadelijk. Het was ook gebleken in muisexperimenten dat een hoge concentratie van amyloid  $\beta$  de concentratie van LRP op het oppervlak van de vaatwandcellen verminderde, doordat LRP in de cel werd afgebroken. Hierdoor kon er minder amyloid  $\beta$  worden afgevoerd en ontstonden er eerder schadelijke plaques. Patiënten met de ziekte van Alzheimer hadden ook minder LRP in de vaatwandcellen van hun hersenen. Amyloid kon dan niet via LRP afgevoerd worden en daardoor ontstond er een hogere concentratie amyloid  $\beta$  dat de plaques in de hersenen veroorzaakte.

LRP heeft dus in potentie de capaciteit om de ziekte van Alzheimer te voorkomen. In vervolggexperimenten kan gekeken worden hoe LRP amyloid  $\beta$  kan opruimen in patiënten, bijvoorbeeld door de opname van amyloid  $\beta$  te bevorderen of de afbraak van LRP tegen te gaan.

## LRP en de klaring van C4BP

Een van de best gekarakteriseerde functies van LRP is zijn rol in de endocytose van liganden. Als een ligand bindt aan LRP aan het celoppervlak, dan wordt zowel LRP als het ligand opgenomen door de cel. Het complex komt dan in zogenaamde endosomen. De zuurgraad hierin is lager dan buiten de cel, wat resulteert in het uiteenvallen van het complex. LRP kan dan herbruikt worden en gaat dan weer naar het celoppervlak. Het ligand wordt getransporteerd naar lysosomen en wordt daar afgebroken. Op deze manier kan de concentratie van een eiwit in bloed geregeld worden.

Recent is op ons laboratorium gevonden dat C4b-binding proteïne (C4BP) een ligand voor LRP is. In het bloed heeft het verschillende functies. Het bindt de stollingsfactor, proteïne S, dat door deze binding zijn functie niet meer kan uitvoeren. Het kan ook het eiwit C4b binden dat betrokken is in de afweer tegen bacteriën. LRP zorgt er uiteindelijk voor dat C4BP wordt afgebroken in de cel.

In dit proefschrift hebben we nader gekeken naar deze opname van C4BP door de cel (hoofdstuk 5). Voordat het aan LRP bindt, werd het eerst uit het plasma gevangen door heparan sulfaat proteoglycanen (HSPGs), eiwitten die in grote hoeveelheden op het oppervlak van cellen aanwezig zijn. We hebben een aantal mutanten van C4BP gebruikt om te kijken welke delen van C4BP nu precies aan LRP bonden. Die mutaties zijn gemaakt in het deel van C4BP dat ook aan heparine bindt. Heparine is een soort suiker dat de bloedstolling remt. Een vorm van heparine is ook aanwezig in de HSPGs. Onze resultaten geven aan dat die delen van C4BP die ook aan heparine/HSPGs bonden ook aan LRP bonden. Omdat C4BP ook aan proteïne S bindt, hebben we gekeken of het C4BP/proteïne S complex ook nog aan LRP kon binden. Het is gebleken dat het complex inderdaad nog aan LRP kon binden.

## Conclusie

In dit proefschrift hebben we 4 verschillende aspecten van LRP beschreven in 4 verschillende celtypen. In elke celsoort lijkt LRP een specifieke functie te hebben. De vraag die nu opkomt, is hoe het mogelijk is dat een en dezelfde receptor totaal verschillende functies heeft in diverse celtypen. Vervolgonderzoek moet uitwijzen hoe deze functies gereguleerd worden door te zoeken naar andere eiwitten in, op of rondom de cel die de specifieke functies van LRP kunnen beïnvloeden.





**Dankwoord**

*Ik moet oppassen dat dit hoofdstuk niet het langste van het boekje wordt. Heel wat mensen heb ik namelijk een plaatsje moeten beloven in het meest gewilde hoofdstuk van dit boekje. Dat betekent dus dat zo'n proefschrift nooit in je eentje tot stand zou zijn gekomen.*

*Peter, bedankt voor de mogelijkheid dat ik op dit project kon werken. Jij als eiwitmens, ik meer celmens. Ik moest gaan kantklossen en op reis (dit laatste heb ik soms iets te letterlijk opgevat...), en uiteindelijk heb ik met alle bagage mijn doel bereikt! Je was altijd erg kritisch, meestal niet onterecht. En natuurlijk ook enthousiast als we op het punt stonden belangrijke resultaten te krijgen. En tja, die confociaal plaatjes kosten nu eenmaal veel tijd.*

*Flip, Jan-Willem, Harry, Martijn, Ton, bedankt voor jullie kritische blik op mijn resultaten tijdens de werkbeprekingen.*

*Lenting-groepje, bedankt voor alle interesse gedurende deze 4 jaar. Ondanks dat we op veel verschillende onderwerpen werken, was er altijd goed overleg. Erik, bedankt voor je hulp bij de Biacore (moest je maar ergens anders gaan zitten). Sarah, bedankt voor je interesse in alles en voor alle moeite voor de laatste C4BP experimenten (Dit stukje mag toch wel in het Nederlands?). Ronan, finally your experiments will work. You only have to wait for the ultimate blunder. Good luck with all your future experiments! But when will you start talking Dutch? Carina, het was altijd een gezellig uitstapje naar Lab I. Nog niet genoeg van VWF? Even blijven doorzetten nog! Alles komt uiteindelijk op zijn pootjes terecht. En natuurlijk ook bedankt voor de gezelligheid buiten het lab. Caroline, we worked together for only a couple of weeks, but anyway: good luck with the sacred protein LRP! Lucy, bedankt voor al je hulp bij de laatste loodjes, ook al werkt niet alles zoals je zou willen (welkom op de research!).*

*Lab II (van voor en na de grote exodus). In willekeurige volgorde: Maarten (Ja ik ben eigenwijs, en ja ik kom uit Den Bosch), Cees ("nee, geen Foo Fighters"), Rolluf ("nee, geen Radiohead!"), Menno "Flummel"*

(je bent van me af!), Bettina, Barend, Mieke, Eszter, Remo, Laura, Caroline, Dafna, (kale)Bas ((ex)buurman, bedankt dat je het zo lang vol hebt kunnen houden), Winnie (ik kon alle verhalen over kinderen nog wel aan hoor), Jelle, Lucy, Marieke, Alex). Een hoop lol, een hoop rot-zooi, een hoop discussies (poldermodel?), maar ondanks alles was er toch nog te werken (dat is toch de oorspronkelijke functie van G03.601?). Bedankt voor al het gezeur dat jullie gewild en ongewild wilden/moesten aanhoren! En ook sorry voor mijn zangkwaliteiten (Bløf!). En vooral: zet jullie CD-verzameling weer op. En wie moet nu trouwens alle falconbuisen gaan halen?

Lab I. Even weer die falconbuisen dus gaan halen en ik bleef soms weer veel te lang hangen op het 'theekransje' (Janine (overloper naar lab III), Chantal (gedeeltelijke overloper) en Carina).

Lab III, voorbeeldlab, hoe houden jullie het zo schoon! Marjolijn, al jaren vriendin, later ook collega's en nu moest ik zelfs van je rafts leren isoleren. Suzanne, bedankt voor je gezelschap tijdens de vrijdagochtendritjes. Ik was toch nooit zo vervelend? Beloof je om nooit meer te stressen? Irlando, bedankt voor alle mental coaching sessies.

Alex, Lex voor vrienden, bedankt voor al je hulp als stagiaire. Tussen alle bakkies door heb je toch veel bijgedragen aan dit proefschrift. Had je ooit gedacht dat het zo zou uitpakken? En je kon als een van de weinigen met mijn chaotisch gedrag omgaan! Hoofdstuk 3 is van ons! Succes met je tweede stage en het AIO-zijn later (ze hebben je gewaarschuwd).

Joukje, bedankt voor alle administratieve dingen. Hoe vaak heb ik trouwens gezegd dat een pakketje echt het laatste pakketje naar Amerika was?

Alle oud-labgenootjes, bedankt voor al jullie hulp. Bianca, bedankt voor het wegwijs maken op het lab.

*Alle bewoners van andere labs: bedankt voor alle chemicaliën waarvoor ik bij jullie kwam shoppen. Hoeklab, daar kwam ik weer met mijn FACS problemen. Ik ben blij dat jullie nu pas verhuisd zijn, zodat ik niet zo ver hoefde te lopen voor de centrifuge. Jordan VI, bedankt voor de samenwerking op het CKL en voor het beantwoorden van al mijn leukovragen. Tuna, Saskia, Anton, Henk, Tineke en Elles, bedankt voor jullie hulp bij het lymfo-verhaal. Tja, wij als haemostaselab weten nu eenmaal niets van immunologie.*

*Niels, hier sta je dan in het dankwoord. Ik heb je het vaak genoeg moeten beloven! Hé, bedankt voor alles. Bedankt voor al je geduld (en jij hebt mij ook geleerd wat geduld is), theoretisch advies en gezelligheid. O nog iets: niet al te arrogant worden, hè?*

*Bedankt alle anderen in het ziekenhuis die ik vergeten ben. Niet boos worden hè?*

*Tanja-Maria, Eveliina, Leena, Maria, Liisa, Yvonne and Carl (and others!). Thank you for your hospitality during my stay in Helsinki. I really appreciate all technical advice, but, also advice for how to visit Helsinki in a few days! I love your city. Liisa, I hope you enjoyed your stay here in Utrecht. Thank you for your attempts to answer my e-mails about things you've never heard about.*

*Paula, I'm sorry I'm not so fast in perfusion experiments as you are. Many, many times I told you how much I'm sorry to ask you to perform these experiments. Here's my last sorry: Sorry! But anyway, it resulted in a really nice paper! Jaap-Jan, je kunt sneller denken dan dat wij kunnen bijhouden. Die halfuurtjes op het CLB waren daarom altijd erg vruchtbaar! Bedankt voor het brainstormen.*

*Ik heb eigenlijk altijd gezegd dat vrienden niet echt bijgedragen hebben aan dit boekje. Niemand snapt toch wat ik nou die 4 jaar heb gedaan. Maar ik kan het natuurlijk niet maken om jullie te vergeten! Al die etentjes en dagjes uit zijn natuurlijk net zo belangrijk als de hele*

dag pipetteren op het lab. Marjolijn, Marjolein (leuk, dezelfde namen, maar jullie wisten altijd precies over wie we het hadden! En leuk dat jullie mijn paranimfen willen zijn), Linda + alle aanhang: waarom wonen we nu zo ver van elkaar? Niet nog verder weggaan, hè? De vakantie in Frankrijk als een van de hoogtepunten. De eigenlijk vee-eeeeeeel te weinig etentjes in het jaar (ja, we gaan nu écht vaker afspreken), waar het werk op een of andere manier toch altijd een gespreksonderwerp was. Marjolijn, jij bleef in ieder geval dichtbij wonen en natuurlijk woonde je ook op het hematlab. Marjolein: leuk dat je weer in de buurt komt wonen. We moeten meer afspreken dan! Linda: niet rond de 20° bevallen hè? Bianca, Daniëlle en Judith: vanaf nu beloof ik dat ik jullie vaker mail en sms en bel! Richard, Schotse Hooglander, jij bent volgens mij de enige leek die ooit het lab heeft bezocht. En je hebt onverwachts toch nog iets bijgedragen aan het proefschrift (weet je nog wat? ;-)). Carina, Michiel, Casper, Esther, Astrid, Bart, Linda, wandelmaatjes, bedankt voor jullie gezelschap tijdens de weinige sociale contacten in de laatste maanden. Michiel, zie je: ook tijdens het MSN-en lukt het me om dit proefschrift te schrijven! Bedankt voor je praatjes.

Bien, "druk" met afstuderen, maar toch nog tijd om met me naar Bløf te (moeten) gaan. De voorkant is echt mooi geworden! Succes verder.

Pap en mam, tja hier heb ik het dan al die jaar voor gedaan. Sorry dat ik nooit kon uitleggen wat ik nu gedaan heb. Nou, speciaal voor jullie de 'Samenvatting in het Nederlands'. Mam, tja, ik kwam niet zo vaak meer langs, maar je viel me vaak genoeg lastig op MSN om contact te hebben. Pap, het is toch niet echt 'iets met weefsels'. En standaard 8 uur werken ("als je om 8 uur begint, mag je toch om half 5 naar huis"), dat zit er dus ook niet in. Maar toch bedankt hoor!

Groetjes,

Patricia



## **Curriculum Vitae**

De schrijfster van dit proefschrift werd geboren op 6 september 1978 te 's-Hertogenbosch. In 1996 behaalde zij het gymnasium diploma aan het Jeroen Bosch College te 's-Hertogenbosch. In hetzelfde jaar werd begonnen met de studie Medische Biologie aan de Universiteit Utrecht, waarvan in 1997 het propaedeuse diploma werd behaald (*cum laude*). Van dezelfde studie werd in 2001 het doctoraal diploma behaald met als hoofdvak Celbiologie (Prof. Dr. G. Strous, Dr. W. Stoorvogel en Dr. E.M. van Dam, Universitair Medisch Centrum Utrecht) en als bijvak Pediatrische Endocrinologie (Dr. S.C. van Buul-Offers en Dr. J.G. Koster, Universitair Medisch Centrum Utrecht). Van april 2001 tot juni 2005 was de schrijfster werkzaam als assistent in opleiding bij de vakgroep Haematologie van het Universitair Medisch Centrum Utrecht onder begeleiding van Prof. Dr. Ph.G. de Groot en Dr. P.J. Lenting. De resultaten van het onderzoek zijn beschreven in dit proefschrift, waarvan de verdediging op 20 september 2005 plaats zal vinden.