

Fe(III)–Lipoxygenase Converts Its Suicide-Type Inhibitor Octadeca-9,12-diynoic Acid into 11-Oxo-octadeca-9,12-diynoic Acid

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ABSTRACT: Triple bond analogues of polyunsaturated fatty acids irreversibly inactivate lipoxygenases. During the inactivation the inhibitors are converted enzymatically [Kühn, H., *et al.* (1984) *Eur. J. Biochem.* 139, 577–583]. Since the converted inhibitor molecules may hold important information about the inactivation mechanism, we have determined the structure of the product that is formed during the irreversible inactivation of soybean lipoxygenase-1 by octadeca-9,12-diynoic acid (ODYA), the triple bond analogue of linoleic acid. This product is formed only in the presence of Fe(III)–lipoxygenase-1 and O₂. It was purified by C18 solid phase extraction and reversed phase HPLC and was identified with UV, IR, and NMR spectroscopic and mass spectrometric techniques as the novel lipoxygenase product, 11-oxo-octadeca-9,12-diynoic acid (11-oxo-ODYA). It is estimated that each lipoxygenase molecule produces 8–10 11-oxo-ODYA molecules before it is inactivated. Furthermore, we have shown that in a secondary reaction 3–4 molecules of 11-oxo-ODYA are covalently attached per lipoxygenase molecule, most likely, to solvent-exposed amino groups. This leads to the formation of a *N*-penten-4-yn-3-one chromophore, $\text{RC}(\text{NHX})=\text{CHC}(\text{O})\text{C}=\text{CR}^1$, in which X stands for the protein and R or R¹ for $\text{CH}_3(\text{CH}_2)_4$ - or $-(\text{CH}_2)_7\text{COOH}$, respectively. Fe(II)– and Fe(III)–lipoxygenase remain active upon reaction with purified 11-oxo-ODYA. It is concluded that (a) several enzymatic turnovers are required for the complete inactivation of lipoxygenase by ODYA and (b) covalent attachment of 11-oxo-ODYA occurs outside the active site and is not the cause of the inactivation.

Lipoxygenases (linoleate:oxygen oxidoreductase, EC 1.13.11.12) are iron-containing dioxygenases that convert unsaturated fatty acids possessing one or more 1Z,4Z-pentadiene systems into Z,E conjugated hydroperoxy fatty acids. Mammalian lipoxygenases catalyze the first step in the formation of leukotrienes and lipoxins from arachidonic acid (Yamamoto, 1992). These products play an important role in the immune response and in inflammatory processes. The physiological role of plant lipoxygenases is less clear, but possible roles have been suggested in plant growth and development, biosynthesis of regulatory molecules, senescence, and response to wounding and pathogens (Siedow, 1991). Because of their abundance, plant lipoxygenases, and in particular lipoxygenase-1 from soybeans, are easily obtained in quantities that allow extensive structural (*e.g.*, Shibata *et al.*, 1988; Van der Heijdt *et al.*, 1992; Boyington *et al.*, 1993; Minor, 1993) and mechanistic studies (*e.g.*, de Groot *et al.*, 1973; Verhagen *et al.*, 1978; Ludwig *et al.*, 1987; Schilstra *et al.*, 1993). Because of structural and functional similarities between lipoxygenases from various sources, soybean lipoxygenase-1 can serve in many respects as a model system for other lipoxygenases.

There is considerable interest in food chemistry and in medicine (Nuhn *et al.*, 1991; Ford-Hutchinson *et al.*, 1994) in compounds that specifically inhibit lipoxygenase catalysis, either reversibly or irreversibly. Analogues of natural fatty acids which contain one or more triple bonds instead of the Z-double bonds are well-known inhibitors of lipoxygenase and cyclooxygenase (Schewe *et al.*, 1986). They cause irreversible inactivation of lipoxygenases. The inactivation mechanism and the inactivation potential of a number of these compounds have been investigated by several authors (Blain & Shearer, 1965; Downing *et al.*, 1970, 1972; Hammerström 1977; Corey & Munroe, 1982; Corey & Park, 1982; Kühn *et al.*, 1984, 1991; Shieh *et al.*, 1985; Corey, 1987; Borel *et al.*, 1993). The mechanism by which triple bond analogues of natural substrates inactivate lipoxygenases has not yet been unambiguously established. However, it is generally accepted that the inactivation is caused by an O₂-dependent, suicide-like action of the enzyme in which the inhibitor is enzymatically converted. In a study on the action of ¹⁴C-labeled ETYA¹ methyl ester on soybean lipoxygenase, quite large amounts of a radioactively labeled product were observed (Kühn *et al.*, 1984). The reaction product was not characterized any further, nor was the product formation quantified.

In order to identify the processes that eventually result in the irreversible inactivation, we studied the enzymatic conversion of the linoleic acid analogue octadeca-9,12-diynoic acid (ODYA) by soybean lipoxygenase-1. In this

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paper, we present the characterization of the product that is formed during the irreversible inactivation of soybean Fe(III)–lipoxygenase-1 by ODYA.

MATERIALS AND METHODS

Materials. ODYA (octadeca-9,12-dienoic acid) was synthesized as described (Otsuki *et al.*, 1986); tetradeca-5,8-diyn-7-ol was prepared as described (Brandsma, 1988) and oxidized with MnO_2 to tetradeca-5,8-diyn-7-one (Dinwoodie *et al.*, 1962). A 1:1 mixture of 5-(*N*-methylamino)tetradeca-5-en-8-yn-7-one and 6-(*N*-methylamino)tetradeca-6-en-9-yn-8-one was prepared from tetradeca-5,8-diyn-7-one and methylamine as described (Metler *et al.*, 1968).

Linoleic acid, (9*Z*,12*Z*)-octadeca-9,12-dienoic acid (99% pure), was from Sigma. 13-HPOD (13(*S*)-hydroperoxy-(9*Z*,12*E*)-octadeca-9,12-dienoic acid) was prepared enzymatically as described (Schilstra *et al.*, 1992). Solutions of 13-HPOD (200 mM), linoleic acid (300 mM), and ODYA (300 mM) were stored under nitrogen in methanol (Merck, gradient grade) until use. Acetic acid and sodiumborohydride were from Merck. Pyridine, hexamethyldisilazane, and trimethylchlorosilazane were from Fluka. All reagents were of the purest grade available. Lipoxygenase-1 from soybeans (White Hilum) was purified as described (Finazzi Agrò *et al.*, 1973). Its concentration was estimated from the absorbance at 280 nm ($\epsilon_{280} = 1.6 \times 10^5 \text{ mol}^{-1} \cdot \text{L} \cdot \text{cm}^{-1}$). The enzyme (Fe(II)–lipoxygenase) was stored at 4 °C at a concentration of 17 mg/mL in 0.05 M sodium acetate buffer, pH 5.5. Fe(III)–lipoxygenase was prepared by incubation of Fe(II)–lipoxygenase with a 5-fold excess of 13-HPOD for approximately 3 min, followed by dialysis against a 0.1 M sodium borate buffer, pH 10 at 4 °C. In all experiments Fe(III)–lipoxygenase was used immediately after preparation.

Equipment. UV absorption spectra were recorded with a Hewlett Packard HP 8452A diode array spectrophotometer. IR spectra of the compounds were recorded with a Mattson RS 5 FT-IR spectrometer equipped with a DTGS detector, using a HATR accessory with a 45° ZnSe crystal (SpecAc). Infrared spectra of all samples were measured from 4000 to 600 cm^{-1} with a resolution of 2 cm^{-1} and coaddition of 200 scans in a nitrogen atmosphere at 20 °C. Interferograms were apodized with a triangular function.

^1H -NMR spectra were recorded with a Bruker AC 300 (300 MHz) spectrometer in CDCl_3 at 27 °C. ^1H NMR chemical shifts are given in ppm (δ) relative to internal TMS.

EPR spectra of native lipoxygenase (20 mg/mL) were recorded with a Varian E-9 spectrometer at 5 K.

GC-MS analysis was performed on an Interscience GC 800 Series gas chromatograph, equipped with a DB-1 column (30 m \times 0.31 mm i.d.) using a temperature gradient from 140 °C, rising at 4 °C/min to 300 °C (2 min). The mass

spectra were recorded with a Fisons Instruments MD 800 MassLab spectrometer under electron impact with an ionization energy of 70 eV.

FAB-MS analyses were carried out on a JEOL JMS SX/SX 102A four-sector mass spectrometer, operated at 10 kV accelerating voltage, equipped with a JEOL MS-FAB 10 D FAB gun operated at a 10 mA emission current, producing a beam of 6 keV xenon atoms. Data acquisition and processing were performed using JEOL Complement software. MS/MS CID mass spectra were acquired using a collision cell in the third field-free region of the instrument, using air as collision gas. The pressure of the collision gas was adjusted to obtain a 50% reduction of the beam. Resolution of the instrument was maintained at 1000 throughout all experiments. The underivatized samples were dissolved in CHCl_3 and a *m*-nitrobenzyl alcohol/sodium iodide matrix was used.

Oxygen uptake was measured in 1 mL of 0.1 M sodium borate buffer, pH 10.0 at 25 °C, using a standard Hansatech Clarke-type electrode disk and a Hansatech CB1 control box.

Lipoxygenase activity measurements were carried out using a Hi-Tech SF-51 stopped-flow apparatus equipped with an LS-10 light source and PM-60 photomultiplier.

HPLC analyses were performed using a Hewlett Packard HP 1090 liquid chromatograph, equipped with an HP 1040A diode array detector and an HP 79994A analytical workstation. HPLC analyses and purification of fatty acids were carried out on CP-Spher C18 (5 μm ; 250 \times 4.6 mm i.d.; Chrompack) or Cosmosil 5C18-AR (5 μm ; 250 \times 10 mm i.d.; Nacalai Tesque) columns, using a methanol/water/acetic acid mixture (80/20/0.01 v/v) as the eluent at flow rates of 1 mL/min (Chrompack column) or 3 mL/min (Nacalai Tesque column). Sephadex G-25 (25 cm \times 1 cm i.d.; Pharmacia; Gilson Minipuls 3 pump, elution rate 0.5 mL/min) or Supradex G-200 (100 cm \times 25 mm i.d.; Pharmacia; Pharmacia FPLC P-500 pump and Liquid Chromatography Controller LCC-500, elution rate 0.4 mL/min) columns were used for size exclusion chromatography. The eluates were passed through a flow cell (300 μL), and every 5 s spectra were recorded.

Preparation and Purification of the Reaction Products. In order to obtain sufficient amounts of reaction product, typically 2 μM Fe(III)–lipoxygenase was incubated for 90 min at 25 °C in air-saturated 0.1 M sodium borate buffer, pH 10.0, with 50 or 75 μM ODYA in a total volume of 100 mL. The reaction was terminated by acidifying the mixture to pH 5 with 3 M HCl and then the fatty acids were extracted as described (Van Aarle *et al.*, 1991). Further purification was carried out by preparative HPLC (Nacalai Tesque column, see Equipment section). The pure compound was dissolved in 250 μL of methanol and stored under nitrogen at –25 °C.

Chemical Modifications. After purification, the enzymatically converted ODYA was reduced, methylated, hydrogenated, and silylated as described (Van Aarle *et al.*, 1991).

RESULTS

To study the mechanism of inactivation of lipoxygenase by triple bond analogues of natural fatty acid substrates, Fe(III)– and Fe(II)–lipoxygenases were incubated with the linoleic acid analogue octadeca-9,12-dienoic acid (ODYA), and the reaction products were analyzed.

¹ Abbreviations: ETYA, eicosa-5,8,11,14-tetraynoic acid; EPR, electron paramagnetic resonance; FAB-MS, fast atom bombardment mass spectrometry; FAB-MS/MS, FAB ionization, collision induced decompositions (CID) in the third field-free region (3rd FFR); FT-IR, Fourier transform infrared spectroscopy; GC-MS, gas chromatography–mass spectrometry; (RP-)HPLC, (reversed phase) high performance liquid chromatography; 13-HPOD, 13(*S*)-hydroperoxy-(9*Z*,11*E*)-octadeca-9,12-dienoic acid; 11-hydroxy-ODYA, 11-hydroxyoctadeca-9,12-dienoic acid; 11-oxo-ODYA, 11-oxooctadeca-9,12-dienoic acid; ^1H -NMR, proton nuclear magnetic resonance; ODYA, octadeca-9,12-dienoic acid; TMS, tetramethylsilane.

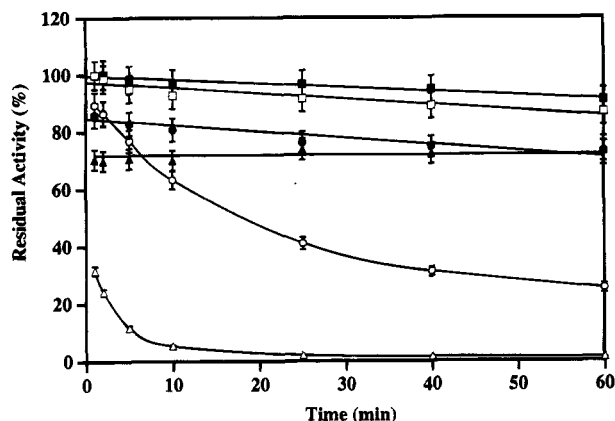


FIGURE 1: Residual activities, measured in a stopped-flow apparatus, of 100 nM Fe(II)- (closed symbols) and Fe(III)-soybean lipoygenase-1 (open symbols) after treatment with 0 (squares), 2 (circles), or 20 μM ODA in 0.1 M sodium borate buffer, pH 10.0 at 25 °C. The reactions were initiated by the addition of ODA at $t = 0$.

Inactivation of Fe(III)- and Fe(II)-Lipoygenase by ODA. In order to investigate whether the effect of ODA on Fe(III)-lipoygenase is different from that on Fe(II)-lipoygenase, 2 or 20 μM ODA was added (at $t = 0$) to 100 nM solutions of either Fe(III)-lipoygenase or, native, Fe(II)-lipoygenase in 0.1 M sodium borate buffer, pH 10.0. The residual activities were measured in a stopped-flow apparatus by mixing the enzyme solutions with equal volumes of 100 μM linoleic acid in 0.1 M sodium borate buffer, pH 10.0. The fraction of active enzyme was calculated from the rate of 13-HPD formation in the test solution (Schilstra *et al.*, 1992). All reactions were carried out at 25 °C. The results are shown in Figure 1. The activities are expressed as percentages of the 13-HPD formation rate observed in the absence of ODA (absolute value: 9.1 μM/s; 10% decrease in activity over 1 h of incubation at 25 °C). Over a 1 h period of incubation, the enzymatic activity decreased by approximately 25% when 2 or 20 μM ODA was added to Fe(II)-lipoygenase, but by 70% (2 μM ODA) or 99% (20 μM ODA) when Fe(III)-lipoygenase was used. The duration of the lag phase in the incubation mixtures containing Fe(II)-lipoygenase increased from 0.4 s (in the absence of ODA) to 1.2 s (2 or 20 μM ODA). From this experiment it can be concluded that ODA is able to inactivate Fe(III)-lipoygenase, but has no effect on the activity of Fe(II)-lipoygenase. The small decrease in enzyme activity in the Fe(II)-lipoygenase samples may be explained by the presence of some Fe(III)-lipoygenase in the Fe(II)-lipoygenase preparation. The EPR spectrum of a concentrated native enzyme preparation indicated the presence of some Fe(III)-lipoygenase (data not shown). The increase of the lag phase of the, native, Fe(II)-lipoygenase preparations after addition of ODA also points to the presence of some Fe(III)-lipoygenase (see Discussion) (Schilstra *et al.*, 1994).

Spectral Changes during the Reaction of ODA with Lipoygenase. Spectral changes or reaction products were not detected when 100 nM Fe(II)-lipoygenase was incubated with ODA for periods up to 5 h. However, after mixing ODA with Fe(III)-lipoygenase, chromophores with absorbance maxima at 258 and 340 nm are formed. The higher the concentration of Fe(III)-lipoygenase at the start of the reaction, the more of these chromophores are

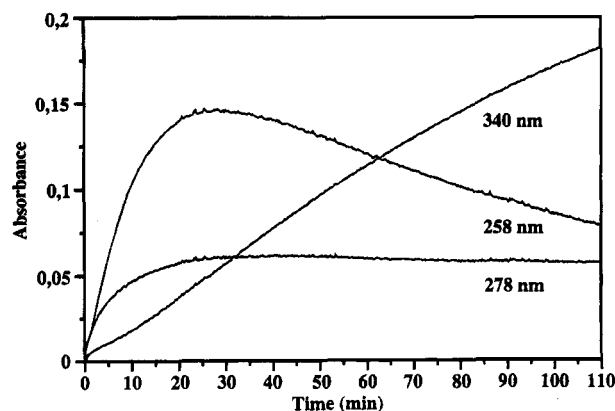


FIGURE 2: Time dependence of the absorbance at 258, 278, and 340 nm of the reaction between 2 μM Fe(III)-lipoygenase and 75 μM ODA in 0.1 M sodium borate buffer, pH 10.0. The values were corrected for the values of A_{258} , A_{278} , and A_{340} of a 2 μM Fe(III)-lipoygenase solution.

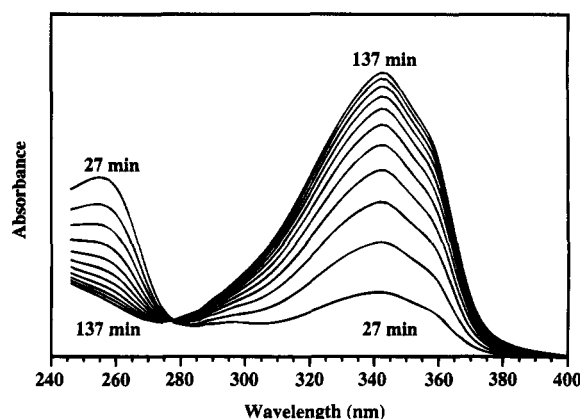


FIGURE 3: Spectra, corrected for the protein absorbance, of the reaction between 2 μM Fe(III)-lipoygenase and 75 μM ODA in 0.1 M sodium borate buffer, pH 10.0, recorded every 10 min during the late stages of the reaction (where A_{278} was constant). Absorption below 240 nm is not shown because of protein interference.

formed. Neither of the chromophores is formed under anaerobic conditions or after incubation of ODA with heat-inactivated Fe(III)-lipoygenase.

In Figure 2, the time course of a typical reaction between Fe(III)-lipoygenase (2 μM) and ODA (75 μM) is shown. In Figure 3 the spectra (corrected for the protein absorbance) of the reaction mixture at 10 min intervals, starting at $t = 27$ min, are shown. In Figures 2 and 3 the isosbestic point is present at 278 ± 2 nm from 27 min after the start of the reaction onward. The data indicate that the chromophore at 258 nm is formed in a primary reaction and that it is converted into the chromophore at 340 nm in a secondary reaction. The primary reaction stops when the dioxygenase activity has ceased, but the secondary reaction continues. Since the 340 nm chromophore is formed in the secondary reaction at the expense of the chromophore at 258 nm, the ratio of the molar absorption coefficients ($Q = \epsilon_{258}/\epsilon_{340}$) can be calculated as follows: $Q = -\Delta A_{258}/\Delta A_{340}$. ΔA is the absorbance change measured over 15 min since this period of time afforded a sufficient signal to noise ratio. In Figure 4 it is shown that Q remains constant (0.58 ± 0.02) from approximately 30 min after the start of the reaction. The value of the absorbance that would have been observed at 258 nm if there had been no conversion into the chromophore

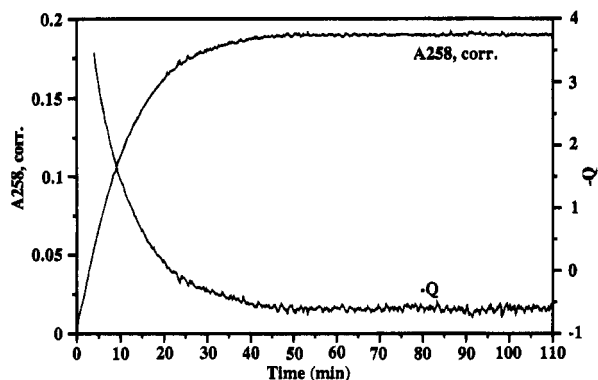


FIGURE 4: Time course of Q and A_{258}^{corr} in the reaction between 2 μM Fe(III)–lipoxygenase and 75 μM ODAYA in 0.1 M sodium borate buffer, pH 10.0. Curve 1 (right vertical axis): values of Q , calculated at time = t (min) as $Q = (A_{258,t} - A_{258,t+15}) / (A_{340,t} - A_{340,t+15})$. Curve 2 (left vertical axis): values of A_{258}^{corr} , calculated from A_{258} and A_{340} in Figure 2 as $A_{258}^{\text{corr}} = A_{258} + 0.58A_{340}$.

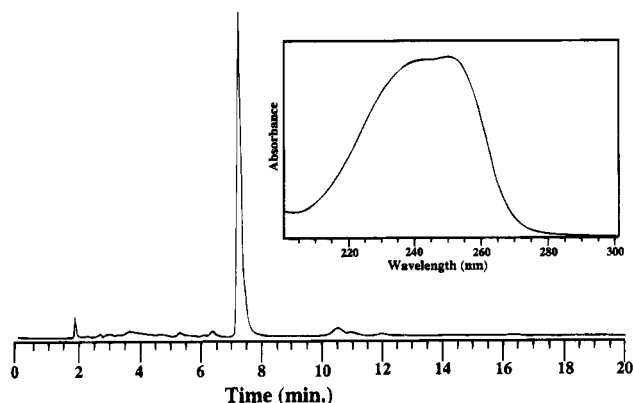


FIGURE 5: HPLC chromatogram ($\lambda = 242$ nm) and UV spectrum (inset) of the product formed by the reaction of Fe(III)–soybean lipoxygenase-1 with ODAYA.

at 340 nm ($A_{258}^{\text{corr}} = A_{258} + 0.58A_{340}$) is also indicated in Figure 4.

During the conversion of ODAYA by Fe(III)–lipoxygenase, O_2 is consumed. A_{258}^{corr} , after the primary reaction had stopped, and the total O_2 uptake in the reaction between 2 μM Fe(III)–lipoxygenase and varying amounts of ODAYA (6, 15, 30, and 60 μM , 0.3, 0.6, and 1.2 mM) were measured. The relationship between the total O_2 uptake and A_{258}^{corr} was linear, with an intercept of $0.01 \pm 0.02 \text{ cm}^{-1}$ and a slope of $(1.0 \pm 0.1) \times 10^4 \text{ mol}^{-1} \text{ L} \cdot \text{cm}^{-1}$. The linear correlation coefficient was 0.97. Assuming that one molecule of O_2 is consumed in the formation of one 258 nm chromophore (hereafter referred to as P1), ϵ_{258} , the molar absorption coefficient of P1 at 258 nm, is equal to the slope of the line. Using this ϵ_{258} , it was calculated that 8–10 P1 chromophores per lipoxygenase molecule were formed during inactivation. The chromophore at 340 nm will be referred to as P2.

Isolation of P1. P1 could be isolated via C18 solid phase extraction provided that the reaction was terminated before P1 was converted into the chromophore at 340 nm (P2). The extract was submitted to analytical and preparative RP-HPLC analysis. The HPLC elution pattern and UV spectrum are shown in Figure 5. No peaks with an absorption maximum at 340 nm were observed in this elution pattern.

UV and FTIR Spectroscopy of P1. The UV absorption spectrum of purified P1 has maxima at 242 and 258 nm with relative intensities of 1.00:1.06 (see inset, Figure 5). The

value of ϵ_{258} is estimated to be approximately $1.0 \times 10^4 \text{ mol}^{-1} \text{ L} \cdot \text{cm}^{-1}$ (see above). This chromophore is absent in the spectrum of NaBH_4 -reduced P1. The UV spectrum of P1 strongly resembles that of tetradeca-5,8-diyn-7-one, which has absorption maxima at 240 and 250 nm, with relative intensities of 1.02:1.00 and an ϵ_{250} of $1.17 \times 10^4 \text{ mol}^{-1} \text{ L} \cdot \text{cm}^{-1}$ in methanol. Similar values were reported for trideca-5,8-diyn-7-one, which has absorption maxima at 239.5 and 250 nm, with relative intensities of 1.01:1.00 and an ϵ_{250} of $1.33 \times 10^4 \text{ mol}^{-1} \text{ L} \cdot \text{cm}^{-1}$, when measured in ethanol (Metler *et al.*, 1968; Migliorese *et al.*, 1974).

Strong absorptions at 1626 and 2208 cm^{-1} in the IR spectrum of P1 indicate the presence of a conjugated carbonyl group or a double bond and an asymmetrically substituted triple bond in the molecule. These IR absorptions correlate very well with those at 1624 ($\text{C}=\text{O}$) and 2205 ($\text{C}\equiv\text{C}$) cm^{-1} of tetradeca-5,8-diyn-7-one. The spectrum of NaBH_4 -reduced P1 shows absorptions at 3380 and 2222 cm^{-1} , indicating the presence of a hydroxyl group and a triple bond, respectively. These values are in good agreement with those found at 3356 (OH), and 2198, 2222, and 2254 ($\text{C}\equiv\text{C}$) for trideca-5,8-diyn-7-ol (Metler *et al.*, 1968).

Signals characteristic for allenic bonds are completely absent in both the UV and IR spectra. The disappearance of the absorptions at 242 and 258 nm (UV) and at 1626 cm^{-1} (IR) and the appearance of a signal at 3380 cm^{-1} (IR) upon reduction with NaBH_4 provide strong evidence for the presence of a carbonyl group in P1.

Mass Spectrometry of P1. FAB-MS in the negative mode was used to determine the molecular mass of underivatized fatty acids because exclusively $[\text{M} - \text{H}]^-$ ions are formed without appreciable fragmentation taking place (Tomer *et al.*, 1983; Jensen *et al.*, 1985). The FAB spectrum of ODAYA shows a $[\text{M} - \text{H}]^-$ ion at m/z 275, while FAB analysis of P1 shows a $[\text{M} - \text{H}]^-$ ion at m/z 289. The FAB spectrum of NaBH_4 -reduced P1 contains a $[\text{M} - \text{H}]^-$ ion at m/z 291. The increase in molecular mass from 276 (ODYA) to 290 (P1) and from 290 to 292 (NaBH_4 -reduced P1) points to the presence of a carbonyl group in P1 and the conversion into a hydroxyl group in NaBH_4 -reduced P1 (see above). This was confirmed by the elimination of H_2O (m/z 319) from the molecular ion at m/z 337 in the FAB-MS/MS CID spectrum of NaBH_4 -reduced P1 (see Figure 6).

Determination of the positions of unsaturated bonds in fatty acids was achieved by following the collision induced decomposition (CID) in the third field-free region (3rd FFR) of alkali metal cationized ions (Adams & Gross, 1987). The best results were obtained with sodium cationized ions $[\text{M} + 2\text{Na} - \text{H}]^+$. In these FAB-MS/MS CID measurements, remote-charge site fragmentations, interrupted by unsaturated bonds, take place. In the spectra (see Figure 6) of ODAYA, P1, and NaBH_4 -reduced P1, the ions at m/z 90, 104, 117, 131, and 173 corresponding to $[\text{CO}_2\text{Na}_2]^{++}$, $[\text{C}_2\text{H}_2\text{O}_2\text{Na}_2]^{++}$, $[\text{C}_3\text{H}_3\text{O}_2\text{Na}_2]^+$, $[\text{C}_4\text{H}_5\text{O}_2\text{Na}_2]^+$, and $[\text{C}_7\text{H}_{11}\text{O}_2\text{Na}_2]^+$, respectively, prove that the part of the molecule adjoining the carboxyl group has not changed and that the triple bond between C9 and C10 is unaffected in P1 and in NaBH_4 -reduced P1. In the high mass regions in the spectra of all three compounds (ODYA, P1, and NaBH_4 -reduced P1) the fragments $[\text{M} - \text{CH}_4]^+$ (m/z 305, 319, and 321), $[\text{M} - \text{C}_2\text{H}_6]^+$ (m/z 291, 305, and 307), $[\text{M} - \text{C}_3\text{H}_8]^+$ (m/z 277, 291, and 293), and $[\text{M} - \text{C}_4\text{H}_9]^+$ (m/z 264, 278, and 280) are observed. These fragmentations indicate that the conversion

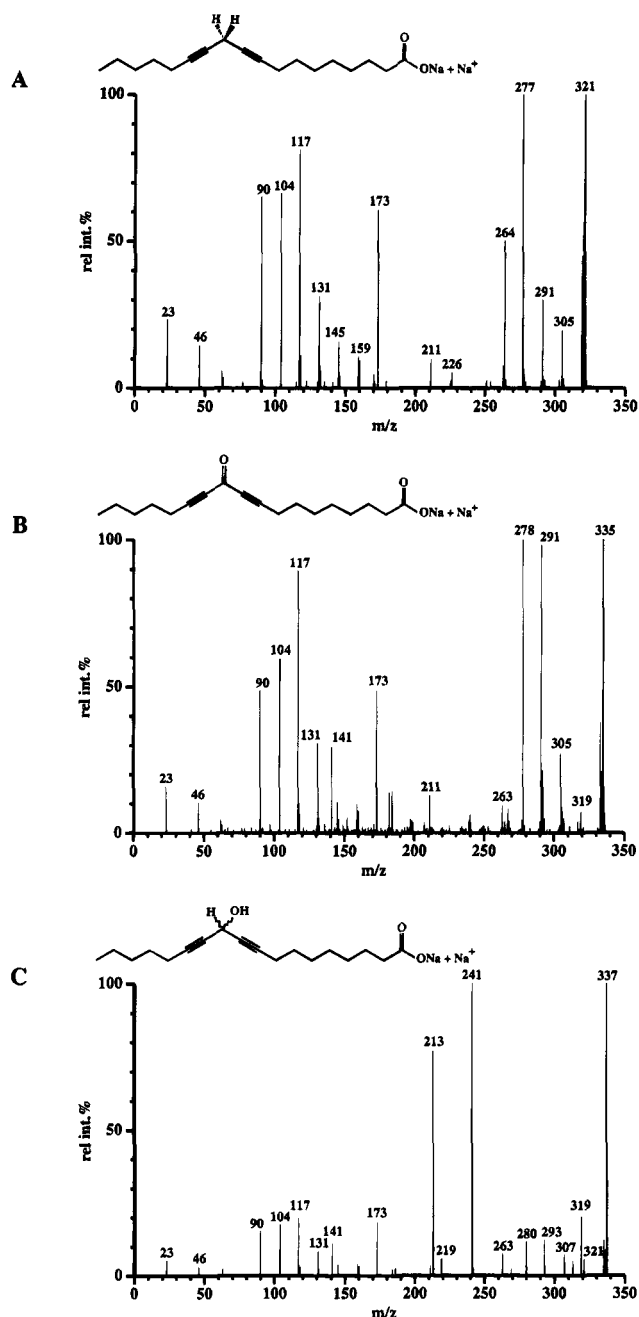


FIGURE 6: FAB-MS/MS spectra of the disodium salts of ODYA (A), 11-oxo-ODYA (B) and 11-hydroxy-ODYA (C).

of ODYA into P1 and the subsequent NaBH_4 reduction of P1 did not change the methyl end of the molecule and that the triple bond between C12 and C13 is unaltered. The FAB-MS/MS CID data of NaBH_4 -reduced P1 also show, apart from the ions described above, major ions at m/z 213 and 241 which are assigned to $[\text{HC}\equiv\text{C}(\text{CH}_2)_7\text{COONa}_2]^+$ and

$[\text{H}(\text{C}=\text{O})\text{C}\equiv\text{C}(\text{CH}_2)_7\text{COONa}_2]^+$, respectively. These fragments are formed *via* a H-shift and are formed only when the hydroxyl is located at C11 in NaBH_4 -reduced P1. On the basis of the FAB and FAB-MS/MS CID spectra of ODYA, P1, and NaBH_4 -reduced P1, it can be concluded that the C11 methylene group of ODYA has been oxidized to a carbonyl group.

The GC-MS (EI ionization) mass spectrum of fully reduced, methylated, and silylated P1 contains major ions at m/z 201 and 287 which are assigned to $[\text{CH}_3(\text{CH}_2)_6(\text{OTMS})\text{CH}]^+$ and $[\text{CH}(\text{OTMS})(\text{CH}_2)_9\text{COOCH}_3]^+$, respectively, providing further evidence that the carbonyl group of P1 is located at C11.

NMR Spectroscopy of P1. Purified P1 was also characterized using ^1H -NMR spectroscopy. Comparison of the spectra of ODYA and P1 (Table 1) reveals that the two protons of the 1,4-diyn system (3.12 ppm, H-11) in the spectrum of ODYA do not occur in the spectrum of P1. Furthermore, the small couplings of 2.4 Hz of H-11 with H-8 and H-14, present in the spectrum of ODYA, are absent in the spectrum of P1, proving that no protons in P1 separated by a triple bond are present. The protons H-8 and H-14 of P1 are chemically identical and have shifted downfield (2.39 ppm) relative to H-8 and H-14 of ODYA, showing the presence of an electronegative substituent near these protons.

The ^1H -NMR spectrum of NaBH_4 -reduced P1 resembles that of ODYA. The resonances of the chemically identical protons H-8 and H-14 have shifted upfield (2.22 ppm) compared with those in the spectrum of P1, and a small coupling of 2.0 Hz, just like in the spectrum of ODYA, reappeared. A signal at 5.1 ppm with a coupling of 1.9 Hz indicates the presence of a proton next to a triple bond close to an electronegative substituent. The fact that the protons H-8 and H-14 have a coupling with a proton at 5.1 ppm after NaBH_4 reduction proves that both H-8 and H-14 are situated next to a triple bond, and according to the results presented above, the carbonyl group can only be situated at C11, between the two triple bonds.

We conclude, therefore, that P1 is 11-oxooctadeca-9,12-diynoic acid (11-oxo-ODYA) and that NaBH_4 -reduced P1 is 11-hydroxyoctadeca-9,12-diynoic acid (11-hydroxy-ODYA).

Characterization of P2. Attempts to extract P2 *via* C18 solid phase, ether, or Bligh & Dyer extractions (Bligh & Dyer, 1959) failed. The reason for this became clear when reaction mixtures were subjected to size exclusion chromatography. A mixture of $8\ \mu\text{M}$ Fe(III)-lipoxygenase and $75\ \mu\text{M}$ ODYA was allowed to react for 5 h at 25°C . When the reaction was terminated, the absorbances at 258 and 340 nm were 0.17 and 0.55, respectively, indicating that approximately $45\ \mu\text{M}$ ODYA was converted into 11-oxo-ODYA in the primary reaction and that almost $30\ \mu\text{M}$ (65%)

Table 1: 300 MHz ^1H -NMR Data of ODYA, 11-Oxo-ODYA, and 11-Hydroxy-ODYA^a

protons	ODYA	11-oxo-ODYA	11-hydroxy-ODYA
H-2	2.35 (t, $J_{2,3} = 7.4$)	2.36 (t, $J_{2,3} = 7.5$)	2.36 (t, $J_{2,3} = 7.4$)
H-8 + H-14	2.15 (tt, $J_{7,8} = J_{14,15} = 7.0$; $J_{8,11} = J_{11,14} = 2.4$)	2.39 (t, $J_{7,8} = J_{14,15} = 7.1$)	2.22 (tt, $J_{7,8} = J_{14,15} = 7.1$; $J_{8,11} = J_{11,14} = 1.9$)
H-11	3.12 (t, $J_{8,11} = J_{11,14} = 2.4$)		5.10 (t, $J_{8,11} = J_{11,14} = 2.0$)
H-18	0.89 (t, $J_{17,18} = 7.1$)	0.91 (t, $J_{17,18} = 7.1$)	0.90 (t, $J_{17,18} = 7.1$)
H-2, 3, 4, 5, 6, 7, 15, 16, 17	1.24–1.78 (m)	1.25–1.72 (m)	1.26–1.81 (m)

^a Chemical shifts (δ) are given in ppm at 27°C and were measured in CDCl_3 relative to internal TMS. Coupling constants (in parentheses) are given in Hz; t = triplet, tt = triple triplet, m = multiplet.

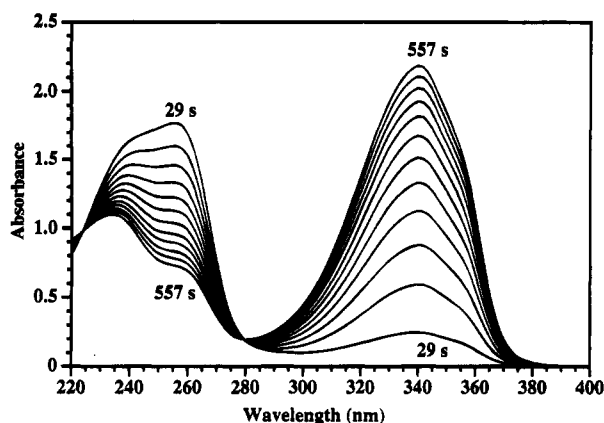


FIGURE 7: UV absorption spectra, recorded every 50 s, during the reaction between 170 μ M 11-oxo-ODYA (P_{c258} , $A_{258} = 1.85$) and 60 mM methylamine in 0.1 M sodium borate buffer, pH 10.0.

11-oxo-ODYA was converted into P2 in the secondary reaction. When this mixture was passed over a Sephadex G-25 column, P2 coeluted with the high-molecular-mass fraction. An identical reaction mixture was also fractionated on a Supradex G-200 column. Also in this experiment, P2 coeluted with the protein peak. Therefore, it seems likely that P2 is formed when 11-oxo-ODYA binds covalently to the enzyme.

In agreement with this suggestion it is observed that incubation of 12.7 μ M 11-oxo-ODYA for 3 h in 0.1 M sodium borate buffer, pH 10.0, does not result in any change in the UV absorption spectrum of 11-oxo-ODYA. However, when 11-oxo-ODYA is incubated with 2 μ M Fe(III)– or Fe(II)–lipoxygenase, the absorbance at 258 nm decreases in 3 h from 0.140 to 0.068, indicating that 6.5 μ M 11-oxo-ODYA is converted into P2. The absorbance at 340 nm increases from 0.015 to 0.142. The spectra taken during the reaction exhibit an isosbestic point at approximately 278 nm. The value of Q ($\epsilon_{258}/\epsilon_{340}$), calculated from these data, is 0.57 ± 0.2 . Both Fe(II)– and Fe(III)–lipoxygenases are fully active after treatment with 11-oxo-ODYA.

According to the experiments described above, each lipoxygenase molecule can covalently bind on average 3–4 molecules of 11-oxo-ODYA. This demonstrates, because of this stoichiometry, and in view of the fact that lipoxygenase remains fully active after treatment with purified 11-oxo-ODYA, that the covalent attachment of 11-oxo-ODYA does not take place in the active site of the enzyme. Compounds with a 1,4-pentadiyn-3-one or 1-propyn-3-one system react readily with primary amines (Chauvelier, 1954; Metler *et al.*, 1968) to form a variety of products, including Schiff bases, addition products, pyridones, and pyrones, depending on the reaction conditions. It is possible that P2 is formed in a reaction between solvent-exposed primary amino groups of the enzyme and 11-oxo-ODYA. To obtain evidence for this suggestion, a large excess (60 mM) of the primary amines methylamine or 3-amino-1-propanol was added to 170 μ M solutions of 11-oxo-ODYA ($A_{258} = 1.85$) in 0.1 M sodium borate buffer, pH 10.0. The course of the reaction of 11-oxo-ODYA and methylamine is shown in Figure 7. From the results it was calculated that Q is 0.54 ± 0.03 . The absorbance at 340 nm in the spectrum of P2 has a shoulder at 360 nm. The absorbance at 340 nm that emerges in the reaction between 11-oxo-ODYA and methylamine also has a shoulder at 360 nm. The product of the

reaction between 11-oxo-ODYA and methylamine was isolated via C18 solid phase extraction and gave two partially overlapping peaks with identical UV-spectra in RP-HPLC analysis. Satisfactory separation of the peaks was not achieved. The adducts were readily soluble in methanol and in chloroform and were stable under mild acidic conditions. FAB-MS analysis of the underivatized adduct mixture revealed only one $[M - H]^-$ ion at m/z 320. So, the two compounds have the same molecular mass of 321.

The adducts of methylamine and 11-oxo-ODYA have the same UV spectrum as a 1:1 mixture of 5-(*N*-methylamino)-tetradeca-5-en-8-yn-7-one and 6-(*N*-methylamino)tetradeca-6-en-9-yn-8-one, with an absorbance maximum at 340 nm and a shoulder at 360 nm. Absorbances at 2216, 2100 ($C\equiv C$), and 1600 ($C=O$) cm^{-1} in the IR spectrum of the methylamine adduct with 11-oxo-ODYA are close to those at 2213 ($C\equiv C$) and 1598 ($C=O$) cm^{-1} in the spectrum of the 1:1 mixture of 5-(*N*-methylamino)tetradeca-5-en-8-yn-7-one and 6-(*N*-methylamino)tetradeca-6-en-9-yn-8-one.

The 1H -NMR spectrum of the methylamine adduct with 11-oxo-ODYA shows two triplets at 2.20 ($J = 7.7$ Hz) and 2.21 ($J = 7.6$ Hz) ppm assigned to two methylene groups next to a double bond. Three triplets at 2.31 ($J = 7.1$ Hz), 2.35 ($J = 7.0$ Hz), and 2.36 ($J = 7.4$ Hz) ppm are assigned to two methylene groups next to a triple bond and the methylene group next to the carboxyl group, respectively. The doublet at 2.98 ($J = 5.2$ Hz) ppm and the singlet at 5.15 ppm are assigned to the *N*-methyl and a vinylic proton, respectively.

These data are in good agreement with the 1H -NMR spectrum of the 1:1 mixture of 5-(*N*-methylamino)tetradeca-5-en-8-yn-7-one and 6-(*N*-methylamino)tetradeca-6-en-9-yn-8-one, with triplets at 2.19 ($J = 7.8$ Hz), 2.20 ($J = 7.9$ Hz), 2.30 ($J = 7.2$ Hz), and 2.32 ($J = 7.0$ Hz) ppm, a doublet at 2.97 ($J = 5.4$ Hz) ppm, and the singlet at 5.14 ppm.

The methylamine adduct with 11-oxo-ODYA is therefore identified as a mixture of 11-oxo-13-(*N*-methylamino)-octadeca-12-en-9-ynoic acid ($CH_3(CH_2)_4C(NHCH_3)=CHC(O)C\equiv C(CH_2)_7COOH$) and 11-oxo-9-(*N*-methylamino)-octadeca-9-en-12-ynoic acid ($CH_3(CH_2)_4C\equiv CC(O)CH=C(NHCH_3)(CH_2)_7COOH$). On the basis of the spectroscopic and chemical properties of the reaction products of 11-oxo-ODYA and methylamine or lipoxygenase, we propose that 11-oxo-ODYA is bound to amino groups of the protein, forming 11-oxo-13-(*N*-X)octadeca-12-en-9-ynoic acid and 11-oxo-9-(*N*-X)octadeca-9-en-12-ynoic acid chromophores, wherein X is the protein.

DISCUSSION

Fe(III)–lipoxygenase converts octadeca-9,12-diynoic acid (ODYA) into a novel product, 11-oxooctadeca-9,12-diynoic acid (11-oxo-ODYA), in the presence of oxygen. During this conversion the enzyme is eventually irreversibly inhibited.

At concentrations of 75 μ M ODYA and 2 μ M Fe(III)–lipoxygenase, 8–10 molecules of 11-oxo-ODYA are estimated to be formed per molecule of lipoxygenase before all of the enzyme has become inactive. These results confirm earlier observations (Kühn *et al.*, 1984) that, to some extent, triple bond analogues of natural fatty acids function as normal substrates. The most likely pathway that leads to the

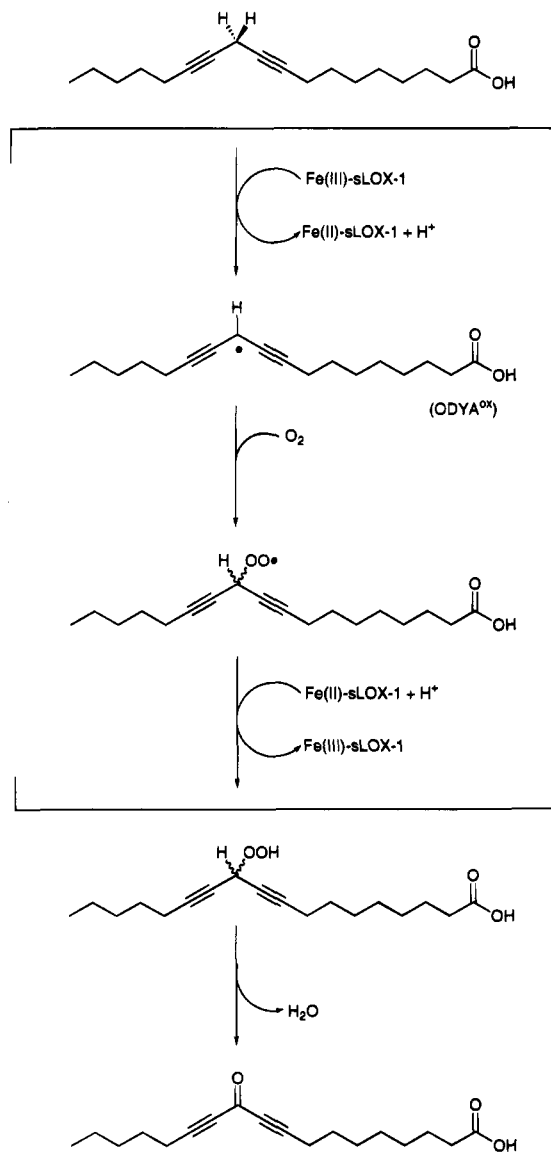


FIGURE 8: Proposed reaction scheme for the formation of 11-oxo-ODYA from ODA and O₂, catalyzed by Fe(III)-lipoxygenase. Fe(II)- and Fe(III)-sLOX-1 are Fe(II)- and Fe(III)-soybean lipoxygenase-1, respectively.

formation of 11-oxo-ODYA from ODA and O₂ is shown in Figure 8.

The fact that only Fe(III)-lipoxygenase catalyzes the conversion of ODA indicates that the first step in the reaction is the abstraction of, most likely, the *pro-S* hydrogen (Corey, 1987), leading to formation of the radical ODA^{ox}. Hydrogen abstraction is accompanied by reduction of Fe(III) to Fe(II). Since more than one molecule of product is formed per molecule of lipoxygenase, the iron cofactor must undergo several cycles of reduction and oxidation before the enzyme is inactivated. In analogy to the reaction with linoleic acid, we propose that Fe(II)-lipoxygenase is reoxidized in an O₂ addition step and that the first reaction product is 11-hydroperoxy-ODYA. The conversion of 11-hydroperoxy-ODYA into 11-oxo-ODYA involves a dehydration reaction. This dehydration may occur spontaneously, but could also be catalyzed by Fe(II)-lipoxygenase, analogously to the reaction of Fe(II)-lipoxygenase with HPOD in the formation of oxidienes. During this reaction Fe(II)-lipoxygenase is oxidized to Fe(III)-lipoxygenase. This is corroborated by the observation that soybean lipoxygenase-1

converts (11*Z*)-eicosa-11-en-14-ynoic acid into 11-hydroperoxy-(12*E*)-eicosa-12-en-14-ynoic acid and 11-oxo-(12*E*)-eicosa-12-en-14-ynoic acid. The hydroperoxide was found to irreversibly inhibit lipoxygenase (Shieh *et al.*, 1985).

As stated above, covalent binding of 11-oxo-ODYA to lipoxygenase does *not* result in inactivation of the enzyme. It is likely, therefore, that the covalent attachment of 0.89 molecule of 14,15-dehydroarachidonic acid to soybean lipoxygenase-1, detected by Corey and Park (1982), bears no relation to the inactivation process. Kühn *et al.* (1984) found no significant labeling of soybean lipoxygenase-1 after its reaction with the ¹⁴C-labeled ETYA methyl ester. However, under the conditions that were used (lipoxygenase and inhibitor concentrations of 22 and 300 nM, respectively), the rate at which the secondary reaction occurs may have been very low. They reported that the inactivation of lipoxygenase was accompanied by the oxidation of a single methionine residue, but this could not be confirmed in later experiments (Höhne *et al.*, 1991). Several authors have proposed that inactivation by triple bond analogues of natural substrates may occur *via* intermediate allenes (Downing *et al.*, 1970; Corey & Park, 1982; Corey *et al.*, 1984; Corey, 1987). Our experiments do not support this hypothesis, because the only enzymatic product that we have detected is 11-oxo-ODYA, and not the 9- or 13-substituted compounds that would be expected to be formed from allenic structures. The absence of allenic intermediates was also confirmed by Shieh *et al.* (1985). Preliminary calculations on the electron distributions in the 1,4-diyne radical system indicate that the radical character is located almost exclusively on C3 and that allenic electron distributions are improbable. Therefore, it seems more likely that the triple bonds remain intact during the reaction. Further quantum mechanistic calculations are currently performed.

In *true* suicide reactions such as the reactions of serine proteases with chloromethyl ketones, or xanthine oxidases with allopurinol, the substrates are enzymatically converted into a product that binds covalently to the active site of the enzyme. In these cases, suicide inactivation occurs with a 1:1 stoichiometry. In contrast, the irreversible inactivation of lipoxygenase by ODA occurs, on average, only after 8–10 turnover cycles. Compared to the suicide inhibitors mentioned above, ODA is, therefore, rather ineffective. In view of the data that have been collected so far on the effect of triple bond analogues of natural fatty acids on lipoxygenase, it is proposed that the inactivation occurs after 8–10 catalytic cycles of the enzyme and is caused by the hydroperoxy radical of the inhibitor. Studies on the nature of the reaction intermediates are in progress.

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