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REVIEW

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# The Role of Halogenative Stress in Atherogenic Modification of Low-Density Lipoproteins

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**Abstract**—This review discusses formation of reactive halogen species (RHS) catalyzed by myeloperoxidase (MPO), an enzyme mostly present in leukocytes. An imbalance between the RHS production and body's ability to remove or neutralize them leads to the development of halogenative stress. RHS reactions with proteins, lipids, carbohydrates, and antioxidants in the content of low-density lipoproteins (LDLs) of the human blood are described. MPO binds site-specifically to the LDL surface and modifies LDL properties and structural organization, which leads to the LDL conversion into proatherogenic forms captured by monocytes/macrophages, which causes accumulation of cholesterol and its esters in these cells and their transformation into foam cells, the basis of atherosclerotic plaques. The review describes the biomarkers of MPO enzymatic activity and halogenative stress, as well as the involvement of the latter in the development of atherosclerosis.

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**Keywords:** myeloperoxidase, reactive halogen species, halogenative stress, low-density lipoproteins, modification of low-density lipoproteins, biomarkers of halogenative stress, atherosclerosis

The main transport form of cholesterol and cholesterol esters in human blood is low-density lipoproteins (LDLs). Increased cholesterol and LDL concentration in the plasma is a diagnostic criterion for the risk of atherosclerosis [1]. Atherosclerosis is a chronic vascular disease associated with the development of atheromatous plaques due to progressive accumulation of cholesterol. The resulting narrowing of the vessels lumen leads to the development of cardiovascular diseases (CVDs). In the light of classic works of N. N. Anichkov, the founder of

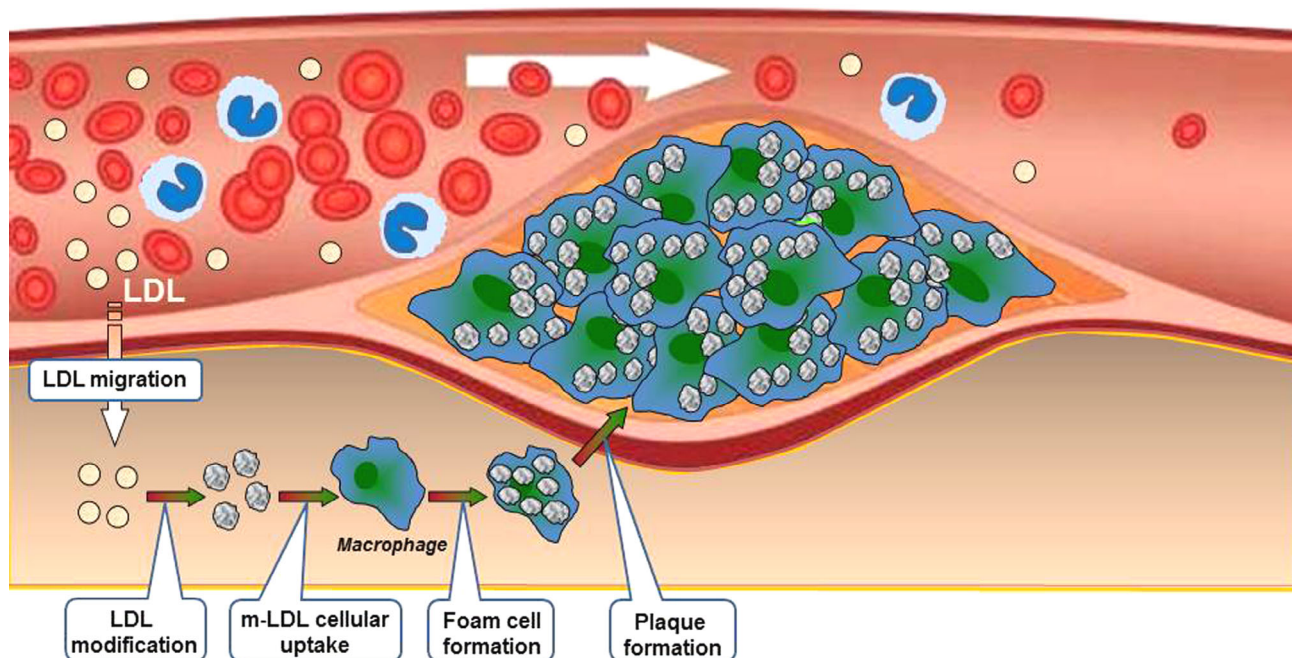
the “cholesterol theory” of atherosclerosis, who postulated that “there is no atherosclerosis without cholesterol” [2], the main efforts in atherosclerosis therapy and prevention have been directed at normalizing the concentration of total cholesterol in the plasma. In 1960s, J. W. Gofman used analytical ultracentrifugation for isolating cholesterol-transporting lipoproteins from the blood plasma [3, 4], and found an increase in the LDL concentration in heart attack patients. By the end of 1970s, it had been proven that LDLs are the source of lipids accumulating in the blood vessels. Hypercholesterolemia associated with a high content of LDLs is still considered the most serious risk factor in CVDs [5].

First studies aimed on the characterization of physicochemical properties of lipoproteins (and not only at determination of their individual classes quantities) appeared in 1970s-1980s. Later, the concept of modified LDLs (mLDLs) different from the LDLs in the plasma of healthy individuals with a known content and ratio of

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**Abbreviations:** ApoB100, apolipoprotein B-100; CP, ceruloplasmin; CVD, cardiovascular disease; HDL, high-density lipoprotein; LDL, low-density lipoprotein; LDL-Br, low-density lipoprotein modified by HOBr; LDL-Cl, low-density lipoprotein modified by HOCl; LPO, lipid peroxidation; mLDL, modified low-density lipoprotein; MPO, myeloperoxidase; RHS, reactive halogen species; VLDL, very low-density lipoprotein.

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**Fig. 1.** The role of mLDLs in the formation of atherosclerotic plaques and atherosclerosis development (see the text). (Colored versions of Figs. 1 and 7 are available in electronic version of the article on the site <http://sciencejournals.ru/journal/biokhsm/>)

protein and lipid components, was generated [6, 7]. Vascular cells and many blood cells capture native LDLs through special ApoB and ApoE receptors. This physiological mechanism of lipid reserve replenishment by the cells is regulated by the negative feedback mechanism and prevents cell overloading with cholesterol [8]. Modifications of the LDL surface result in the generation of mLDLs unrecognized by the ApoB and ApoE receptors. However, mLDLs can be captured by the unregulated scavenger receptor on macrophages and vascular smooth muscle cells. Cell oversaturation with cholesterol esters results in its transformation into the foam cell that serves as a basis of future atherosclerotic plaque [9, 10]. These findings gave rise to a new concept on the leading role of mLDLs in atherogenesis associated with various types of atherogenic LDL modifications. Transformation into foam cells via activity of scavenger receptors takes place regardless of the mechanism of LDL modification: oxidation [11, 12], proteolysis [13, 14], lipolysis [15], desialylation [16, 17], or glycosylation [18].

The role of mLDLs in atherogenesis (in a simplified scheme) is shown in Fig. 1. LDLs migrate from the bloodstream to the subendothelial space, where they become potential targets for various enzymatic and non-enzymatic modifications. The capture of the generated mLDLs by scavenger receptors leads to the overload of phagocytes with cholesterol. Transformation of phagocytes into foam cells initiates formation of atherosclerotic plaques.

The blood of CVD patients contains proatherogenic LDL subfractions that differ from native LDLs in their physicochemical properties. These subfractions include oxidized, electronegative, desialylated, small dense, or aggregated LDLs [19-21]. The main reason for the proatherogenic modification of LDLs has not yet been elucidated. It was suggested that the essential role of atherogenic LDL modifier belongs to myeloperoxidase (MPO), based on the elevated levels of this enzyme in atherosclerotic areas of the aorta [22] and in the blood plasma of CVD patients [23]. MPO is expressed by neutrophils and monocytes and secreted into the extracellular space upon their activation to provide the antimicrobial defense [24]. MPO catalyzes oxidation of halogen ions to the corresponding hypohalous acids. Hypochlorous acid (HOCl) is the most important natural precursor of free radicals produced as a result of chloride oxidation by MPO [25]. At the same time, HOCl and other hypohalous acids trigger the formation of active halogen-containing compounds interacting with protein and lipids in LDLs. These compounds have been named in recent years as *reactive halogen species* (RHS) [25, 26] by analogy with widely used terms *reactive oxygen species* and *reactive nitrogen species* [27, 28].

This review summarizes the data on the effects of MPO, RHS, and halogenative stress (response induced by these compounds) on LDL properties increasing their atherogenicity and responsible for the formation of atherosclerotic lesion in the vascular wall.

# ENZYMATIC ACTIVITIES OF MYELOPEROXIDASE. REACTIVE HALOGEN SPECIES. HALOGENATIVE STRESS

Mature MPO is a glycosylated homodimer ( $M_w \sim 145$  kDa) consisting of two light  $\beta$ -subunits and two heavy  $\alpha$ -subunits containing heme and cysteine residues involved in the formation of disulfide bond between the subunits [29]. The enzyme is found mainly in azurophilic granules of neutrophils and in lysosomes of monocytes. The content of MPO in neutrophils is estimated as 2–5% of total cell protein or 2–4  $\mu\text{g}$  per  $10^6$  cells. The content of MPO in monocytes is  $\sim 0.9\%$  of cell weight [30–32]. Degranulation of neutrophils and monocytes during their activation results in the MPO release into the extracellular space ( $>20\%$  of the MPO total content in neutrophils [33]). MPO can appear outside of neutrophils as a result of necrosis or NETosis [34]. MPO itself is a strong autocrine regulator of neutrophil function. It activates tyrosine kinases by binding to the CD11b/CD18 integrin on the surface of neutrophils and increases cytosolic  $\text{Ca}^{2+}$  concentration, which ultimately promotes degranulation and contributes to further release of granule proteins (including MPO) to the extracellular space [35–38].

Figure 2 shows major MPO-catalyzed reactions. At the first stage (reaction 1), the native enzyme reacts with the  $\text{H}_2\text{O}_2$  forming the so-called Compound I. The latter has two oxidative equivalents and, similarly to any peroxidase, performs sequential single-electron oxidation of various substrates (nitrite, ascorbate, tyrosine, etc.) through the formation of Compound II followed by return to the native enzyme form and completion of the peroxidase

cycle (reactions 1–3). Unlike conventional peroxidases, Compound I MPO is also capable of immediate two-electron oxidation of halides (reaction 4), such as chloride ( $\text{Cl}^-$ ), bromide ( $\text{Br}^-$ ) or iodide ( $\text{I}^-$ ), with the formation of hypochlorous ( $\text{HOCl}$ ), hypobromous ( $\text{HOBr}$ ), or hypoiodous ( $\text{HOI}$ ) acids, respectively, and completion of the halogenation cycle (reactions 1 and 4) [25]:



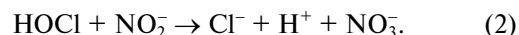
where Hal is a halogen atom.  $\text{HOCl}$  salts are called hypochlorites. The molecular form of  $\text{HOCl}$  and the dissociated hypochlorite anion are present in approximately equal concentrations, because  $\text{p}K_a$   $\text{HOCl}$  is  $\sim 7.5$  [39] in aqueous medium at physiological pH values:



Bromide (20–100  $\mu\text{M}$ ) and iodide ( $<0.6$   $\mu\text{M}$ ) that are present in the blood plasma in significantly lower concentrations than chloride, are also oxidized by MPO [40]. Halogen-containing reactive compounds formed by the enzymatic activity of MPO (first of all, hypohalous acids) are commonly called primary RHS [25, 26].

Due to the high reactivity, primary RHS interact with many functional groups of various compounds, leading to the formation of secondary RHS. The products of hypohalous acid interaction with amines and amides are the most important secondary RHS that damage the protein and lipid components of LDLs. The reaction of primary amines with  $\text{HOCl}$  and  $\text{HOBr}$  produces mono- and dichloramines and mono- and dibromamines, respectively ( $\text{R-NHCl}$ ,  $\text{R-NCI}_2$ ,  $\text{R-NHBr}$ , and  $\text{R-NBR}_2$ , where R is any molecule containing an amino group) [25, 41]. Low concentrations of  $\text{HOCl}$  and, especially  $\text{HOBr}$ , lead to the predominant formation of monohalogenamines under physiological conditions [25]. It is important to note that all major classes of biomolecules in the content of LDLs (proteins, phospholipids, carbohydrates) contain primary amino groups. A similar reaction of amides ( $\text{R-C(O)-NHR}_1$ ) with  $\text{HOCl}$  or  $\text{HOBr}$  leads to the formation of chloro- and bromamides ( $\text{R-C(O)-NClR}_1$  and  $\text{R-C(O)-NBrR}_1$ ), respectively [25, 41]. Secondary RHS also react with functional groups of proteins and other biomolecules, although at a slower rate than primary, most often by replacing the hydrogen atom with the halogen atom [25, 41].

Secondary RHS are also formed in the reactions of hypohalous acids with inorganic compounds. For example,  $\text{HOCl}$  oxidizes nitrite to nitrate in the following reaction ( $k = 7.4 \cdot 10^3 \text{ M}^{-1} \cdot \text{s}^{-1}$  at pH 7.2 and  $25^\circ\text{C}$ ) [42]:



Nitrile chloride ( $\text{NO}_2\text{Cl}$ ) is formed as an intermediate in this reaction [43]. Phenolic compounds, in particular

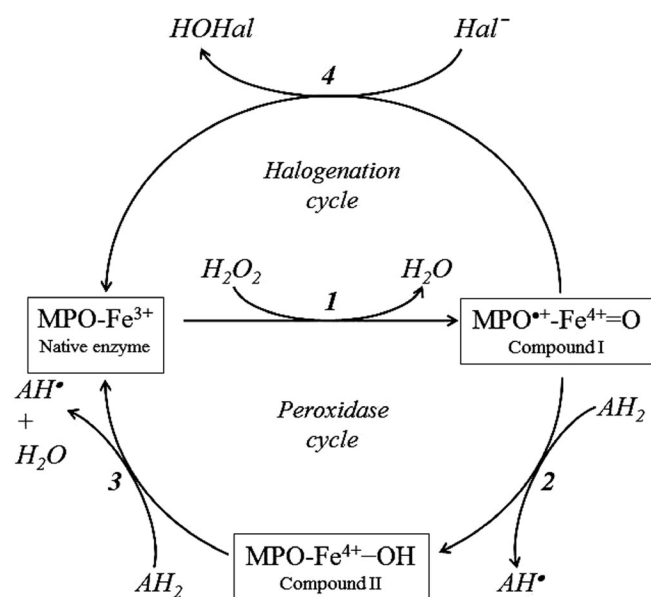


Fig. 2. Halogenation and peroxidase cycles of MPO. Hal, halogen;  $\text{AH}_2$ , peroxidase substrate.

tyrosine, are efficiently nitrated and, especially, chlorinated by  $\text{NO}_2\text{Cl}$  [42]. The latter decomposes into radicals [43]:



leading to tyrosyl radical formation [43] and lipid peroxidation (LPO) in LDLs [42, 44].

It should be emphasized that unlike reactive oxygen species ( $\text{H}_2\text{O}_2$ ,  $\cdot\text{O}_2^-$ ,  $\cdot\text{OH}$ , etc.), the oxidative capacity of RHS is determined by the halogen atom and not by the oxygen atom. It is the halogen atom that participates in the redox reaction by accepting electrons from the oxidized compound and turning into halide:



Due to this, hypohalous acids are strong two-electron oxidizing agents with the oxidizing ability decreasing in order:  $\text{HOCl} > \text{HOBr} > \text{HOI}$  (redox potentials for the  $\text{HOHal}/\text{Hal}^-/\text{H}_2\text{O}$  pairs are 1.08, 0.93, and 0.57 V, respectively) [45].

It is clear that attributing RHS to reactive oxygen species, as it has been done by some authors [46, 47], is fundamentally erroneous. Other names used in the literature, such as chlorinating/brominating oxidants [48], chlorinating intermediates [49], and reactive chlorinating/brominating species [50, 51], do not reflect the board reactivity of the halogen atom manifested in oxidative and other reactions (substitution, addition, etc.) [25, 26, 41]. In our opinion, the term *reactive halogen species* (RHS) proposed in our earlier works [25, 26, 52] and used by other authors [53, 54], is more informative and correct.

The pronounced antimicrobial activity of RHS used in nature for the rapid destruction of infectious pathogens in the inflammation foci is also potentially dangerous, since an uncontrolled efflux of RHS can damage biomolecules and supramolecular structures [25]. This does not occur in a healthy organism, where excessive production of RHS is controlled by a number of factors preventing MPO release by neutrophils, suppression of MPO activity, and RHS scavenging. However, depletion of the anti-halogenative system with simultaneous increase in the RHS production can provoke a so-called halogenative stress characterized by the imbalance between the RHS formation and organism's ability to remove or to neutralize them [25, 26]. Apparently, halogenative stress cannot exist by itself. It is closely associated with the oxidative, nitrosative, and carbonyl stresses, since RHS reactions with various compounds and functional groups leads to the formation of reactive oxygen and nitrogen species and carbonyl compounds [26].

## RHS INTERACTION WITH LDL COMPONENTS

Based on the kinetic models and experimental data, it was found that the main targets of  $\text{HOCl}$  and  $\text{HOBr}$  in

LDLs are Met, Cys, His, Trp, Lys, and Tyr residues, disulfide bonds and terminal primary amino groups of proteins, polar heads of phosphatidylethanolamine and phosphatidylserine, cholesterol unsaturated bonds, and antioxidants (carotenoids, tocopherol, and ubiquinol). Arginine residues and unsaturated bonds of aliphatic fatty acid chains of phospholipids react with  $\text{HOBr}$  faster than with  $\text{HOCl}$ . The rate constants of the second-order reactions of  $\text{HOCl}$  and  $\text{HOBr}$  with the functional groups of these compounds are shown in the table; as a rule, they range from  $10^3$  to  $10^8 \text{ M}^{-1}\text{s}^{-1}$ . Primary RHS react with Gln and Asn residues and protein peptide bonds at a much lower rate and virtually do not react with Ala, Val, Leu, Ile, Pro, and Phe and phosphocholine group of phosphatidylcholine [55-59]. Below, we will discuss in detail the RHS-induced changes in the LDL components (proteins, lipids, carbohydrates, antioxidants) that can be the cause of LDL proatherogenic modification.

**Modification of ApoB100.** ApoB100 ( $M_w \sim 512 \text{ kDa}$ ) is the main protein component of LDLs (up to 98%) recognized by the ApoB and ApoE receptors on the cell surface. Incubation of LDLs in the presence of various  $\text{HOCl}$  concentrations leads to the decrease in the content of amino acid residues in the following order:  $\text{Met} \sim \text{Cys} > \text{Lys} \sim \text{Tyr} > \text{His} \sim \text{Arg}$  [57]. A number of studies have compared the loss of ApoB100 functional groups in LDLs under the action of  $\text{HOCl}$  and in the presence of  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$ . Most of these studies have confirmed similar effects of  $\text{HOCl}$  and  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$ , thereby proving the key role of MPO-generated  $\text{HOCl}$  as the modifying agent [60, 61].

However, in some cases, ApoB100 modifications were different after MPO binding to the LDL surface and  $\text{HOCl}$  addition [62, 63]. The effects of  $\text{HOCl}$  and functioning MPO ( $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$ ) on the modification of amino acid residues in LDL were investigated in [63]. The authors identified 97 peptides containing modified residues (mostly Met, Trp, and Tyr) after trypsinolysis of ApoB100 and found significant differences between peptides modified by  $\text{HOCl}$  and MPO. It was suggested that this difference was due to the site-specific modification of ApoB100 after MPO binding to the LDL surface (in contrast to the LDL modification by  $\text{HOCl}$ ) or by a considerable increase (up to 90%) in the MPO activity upon enzyme binding to the LDL. Analysis of peptides obtained by trypsinolysis of ApoB100 in the content of LDLs isolated from the blood of patients with a high risk of CVD revealed a certain similarity between peptides modified *in vitro* and *in vivo*. Interestingly, some residues (Met4, Met785, Met1873, Met2015, Trp2468, Trp3943, and Met4192) were already modified in the native LDLs, which indicates their hypersensitivity to RHS [63].

According to the kinetics of modification of ApoB100 amino acid residues,  $\text{HOBr}$  is generally more active than  $\text{HOCl}$ .  $\text{HOBr}$  reacts with His, Trp, Lys, Tyr,

The content of functional groups in ApoB100, lipids, and antioxidants in LDLs and the rate constants of the second-order reactions of these groups with HOCl and HOBr

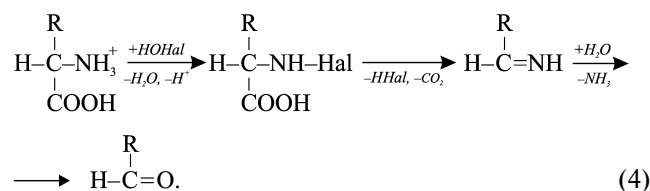
LDL component	Number of molecules per LDL particle	$k_{\text{HOCl}}, \text{M}^{-1}\cdot\text{s}^{-1}$	$k_{\text{HOBr}}, \text{M}^{-1}\cdot\text{s}^{-1}$
<i>ApoB100</i>	[58]	[55]	[56]
Met	78	$3.8\cdot 10^7$	$3.6\cdot 10^6$
Cys	9	$3.0\cdot 10^7$	$1.2\cdot 10^7$
Disulfide bond	8	$1.6\cdot 10^5$	$1.1\cdot 10^6$
His	115	$1.0\cdot 10^5$	$3.0\cdot 10^6$
Lys	357	$5.0\cdot 10^3$	$2.9\cdot 10^5$
Trp	37	$1.1\cdot 10^4$	$3.7\cdot 10^6$
Tyr	151	44	$2.3\cdot 10^5$
Arg	148	26	$1.8\cdot 10^3$
Asn	247	0.03	1.9
Gln	230	0.03	1.9
Terminal $\text{NH}_2$ -group	1	$1.0\cdot 10^5$	$2.0\cdot 10^6$
Peptide bond	4535	10	50-900
<i>Lipids</i>	[58]	[58]	[59]
Phosphatidylserine*	5	$3.3\cdot 10^4$	$9.3\cdot 10^5$
Phosphatidylethanolamine*	15	$1.8\cdot 10^4$	$8.8\cdot 10^5$
Phosphatidylcholine*	610	0.018	—
Double bonds	5095	8.7	$1.1\cdot 10^4$
Sphingomyelin (amide)	171	18.7 [93]	—
<i>Antioxidants</i>	[144]		[59]
Tocopherol	7	$1.3\cdot 10^3$ [58]	$6.4\cdot 10^4$
Ubiquinol	0.10	$1.3\cdot 10^3$ [58]	$2.5\cdot 10^6$
$\beta$ -Carotene	0.29	$2.3\cdot 10^4$ [146]	—

\* Rate constant for the reaction with the phospholipid polar heads.

Arg, terminal  $\text{NH}_2$ -groups, and disulfide bonds faster than HOCl [55, 56].

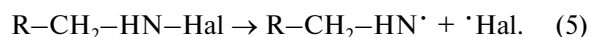
Interaction between primary RHS and proteins often leads to protein aggregation due to the formation of intermolecular cross-links. There are three mechanisms for the formation of such crosslinks associated with the MPO activity.

The first mechanism is HOCl (HOBr) reaction with  $\text{NH}_2$ -groups with the formation of aldehyde that can proceed by two independent ways. The first one is the formation of unstable chlor- and bromamines in the reaction with the terminal  $\alpha$ - $\text{NH}_2$  group with their following conversion into aldehydes through decarboxylation and deamination:

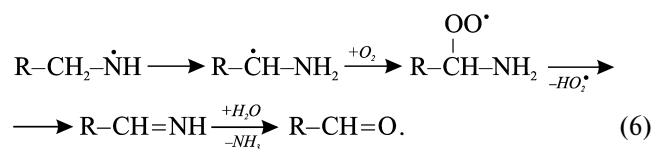


The second one is formation of more stable chlor- and bromamines formed in the reaction of HOCl and

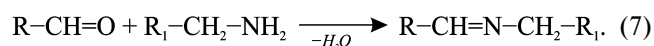
HOBr with the  $\text{NH}_2$ -groups of Lys side chains with the following homolytic decomposition of the N–Hal bond:



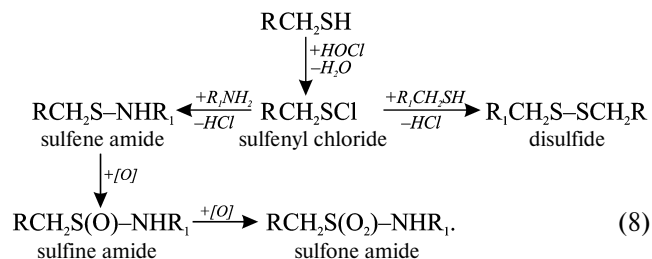
Under aerobic conditions, the N-centered radical migrates to the next C-atom with the aldehyde formation [64]:



Aldehyde reaction with the  $\text{NH}_2$ -group (mainly of Lys) results in the formation of both intra- and intermolecular cross-links:



The mechanism of the formation of intermolecular crosslinks is the interaction between HOCl and the SH-group of Cys residue according to the scheme:



The resulting unstable sulfenyl chloride reacts with the SH-group of Cys with the formation of disulfide bond (–S–S–) or with the amino group of Lys with the formation of sulfene amide (–NH–S– covalent crosslinking). The latter undergoes further oxidation to sulfine amide and then to sulfone amide [65, 66].

Finally, the third mechanism of the cross-link formation in protein is MPO-dependent but not associated with the RHS formation. Tyrosyl radical formed by the one-electron oxidation of Tyr residue in the MPO peroxidase cycle dimerizes into di-Tyr, resulting to the cross-linking [67].

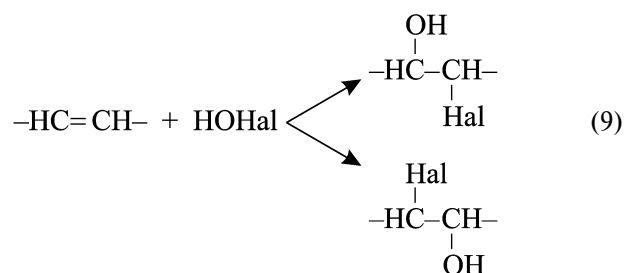
Protein interaction with primary RHS can result in the destruction of peptide bonds and protein fragmentation. Interaction of peptide bond amide group with HOCl or HOBr can lead to the replacement of the hydrogen atom by the halogen atom with the formation of chloro- or bromamides, respectively [25, 55, 56]. The rate constants of the bimolecular reactions of HOCl and HOBr with compounds mimicking the peptide bond strongly depend on the compound chemical structure and vary within 3–4 orders of magnitude. Moreover, the rate of amide group reaction with HOBr is about 30–40 times higher than with HOCl [55, 56]. In aqueous medium, chloramides are slowly hydrolyzed, resulting in peptide bond cleavage and protein fragmentation [68]. Using the spin trap method, it was shown that the N–Cl(Br) bond can undergo homolytic cleavage with the formation of an N-centered radical, e.g., in the presence of ions of transition metals. The N-centered radical is transformed via intramolecular rearrangement into the C-centered radical, leading to the polypeptide chain fragmentation [64, 69]. However, some authors believe that due to the fact that the rate constant for the HOCl reaction with the amide bond does not exceed  $25 \text{ M}^{-1}\text{s}^{-1}$  [55], its occurrence in the LDL *in vivo* is unlikely [57].

ApoB100 can be also modified by nitrile chloride ( $\text{NO}_2\text{Cl}$ ). Nitrile chloride is formed as an intermediate in the nitrite reaction with HOCl and decomposes into free radicals (reaction 3). It was found that Tyr residues are chlorinated and nitrated by  $\text{NO}_2\text{Cl}$  with the formation of tyrosyl radical [42–44]. Nitrated ApoB100 in LDL particles is selectively recognized by the CD36 scavenger receptor on the vascular cells [70]. Similar process can take place in the inflammation foci characterized by increased concentrations of both nitrite and HOCl [57].

**Modification of lipids.** The major targets of RHS in the LDL lipids are unsaturated  $-\text{CH}=\text{CH}-$  bonds of fatty acids and cholesterol and  $\text{NH}_2$ -groups of polar heads of some phospholipids.

**RHS reactions with unsaturated bonds.** There are two main mechanisms of RHS reactions with unsaturated  $-\text{CH}=\text{CH}-$  bonds: molecular mechanism occurring without participation of free radicals and free-radical mechanism (LPO) proceeding through the formation of peroxides and radical intermediate.

The molecular mechanism involves electrophilic addition of HOCl or HOBr to the double bond according to the reaction (9) with the formation of chloro- or bromohydrin isomers, respectively [71–73].



The same products are formed when unsaturated phospholipids are incubated even in the presence of MPO +  $\text{H}_2\text{O}_2$  + halide. Formation of chloro- or bromohydrin is completely prevented by the absence of MPO or one of the substrates ( $\text{H}_2\text{O}_2$  or halide) in the incubation medium or by addition of sodium azide (MPO inhibitor), taurine, or Met (scavenger of HOCl and HOBr) to the reaction. This proves that HOCl and HOBr formed in the MPO-catalyzed reaction participate in the synthesis of chloro- and bromohydrins, respectively [74–81].

If a fatty acid chain contains several unsaturated bonds, any of them can be attacked by HOCl (or HOBr). In the presence of excessive amounts of hypohalous acids, all double bonds enter the reaction [72, 73, 82]. Interaction of polyunsaturated phospholipids containing arachidonic or docosahexaenoic acids with hypohalous acids or MPO +  $\text{H}_2\text{O}_2$  +  $\text{Cl}^-/\text{Br}^-$  results in the formation of lysophospholipids as main reaction products. This is related to the negative inductive effect ( $-I$  effect) caused by the introduction into acyl chains of electron acceptor substituents, such as Cl, Br, OH, during the synthesis (reaction 9) of chloro- or bromohydrins. The  $-I$  effect leads to the hydrolysis of ester bond due to the formation of the electron density deficit on the ester bond carbon atom [77, 79–83].

No formation of lysophospholipids in the reaction of MPO +  $\text{H}_2\text{O}_2$  +  $\text{Cl}^-/\text{Br}^-$  with polyunsaturated phospholipids occurs if the incubation medium lacks the enzyme or one of the substrates ( $\text{H}_2\text{O}_2$  or halide), or contains sodium azide (MPO inhibitor), taurine, or Met (scavenger of HOCl and HOBr). It indicates that polyunsaturated phospholipids are converted into lysoderivatives

under the action of HOCl or HOBr formed in the MPO-catalyzed reaction [77, 80].

The reaction of primary RHS with the unsaturated bond of plasmalogen is of particular interest. Plasmalogens comprise ~4.5% of all phospholipids in LDLs [84]. In plasmalogen, the glycerol hydroxyl group in the sn-1 position is connected through the ether bond with the long aliphatic chain containing the  $-\text{HC}=\text{CH}-$  double bond in the  $\alpha$ -position, which distinguishes plasmalogens from other phospholipids (Fig. 3). This alkenyl ether bond is a target for primary RHS. Plasmalogen interaction with HOCl (or HOBr) results in the formation of lysophosphatidylcholine like that via the phospholipase  $\text{A}_2$ -independent mechanism and aldehyde halogenated at the second position (the first stage in Fig. 3) [51, 85, 86].

A similar reaction was observed when RHS were generated by  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-/\text{Br}^-$  or activated neutrophils. HOBr reacted with the plasmalogen alkenyl ether bond at neutral pH, unlike HOCl that reacted only at acidic pH values. If  $\text{Cl}^-$  and  $\text{Br}^-$  were simultaneously present in the reaction mixture as MPO substrates, the main product was 2-bromoaldehyde [51, 85].

Simultaneous presence of plasmalogen and unsaturated phosphatidylcholine in the reaction mixture led to the preferential generation of lysophosphatidylcholine

from plasmalogen followed by the formation of the unsaturated bond chlorohydrin in the lysophosphatidylcholine acyl chain [87]. When plasmalogen contained a monounsaturated fatty acid in the sn-2 position, lysophosphatidylcholine and 2-halogenoaldehyde were formed at the first stage. And only at the second stage, the mono-unsaturated lysophosphatidylcholine was converted to its halogenohydrin (Fig. 3) [86, 87]. Reactions of HOCl and HOBr with the plasmalogen alkenyl ether bond are much faster than with the unsaturated bond of the phospholipid acyl chain [88]. However, if a polyunsaturated (e.g., docosahexaenoic) acid is present in the sn-2 position of plasmalogen, the reaction with HOCl results not only in the formation of 2-chloroaldehyde but also in subsequent cleavage of the polyunsaturated acid with the generation of glycerophosphocholine (probably via the mechanism described above for polyunsaturated phosphatidylcholine) [86].

Similar reactions take place in LDLs. Incubation of LDLs in the presence of  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  results in accumulation of 2-chlorohexadecanal and 2-chlorooctadecanal, as well as chlorohydrins of unsaturated lysophosphatidylcholines. When LDLs were incubated with activated monocytes, the content of 2-chloroaldehydes increased severalfold in both LDLs and the cells [50].

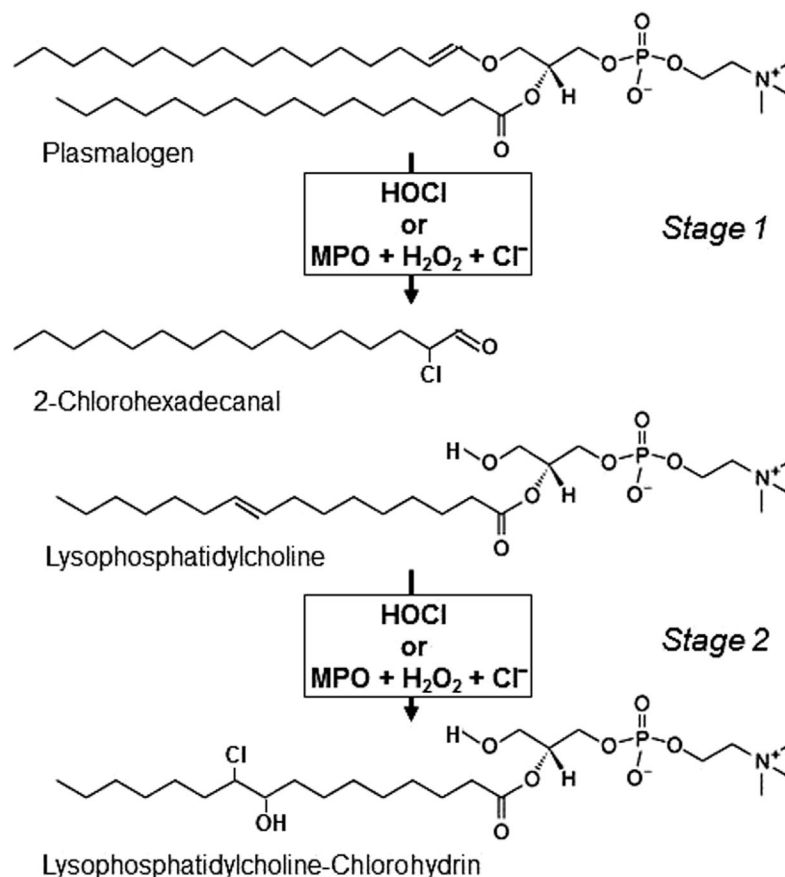


Fig. 3. Plasmalogen modification by HOCl or  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$ .



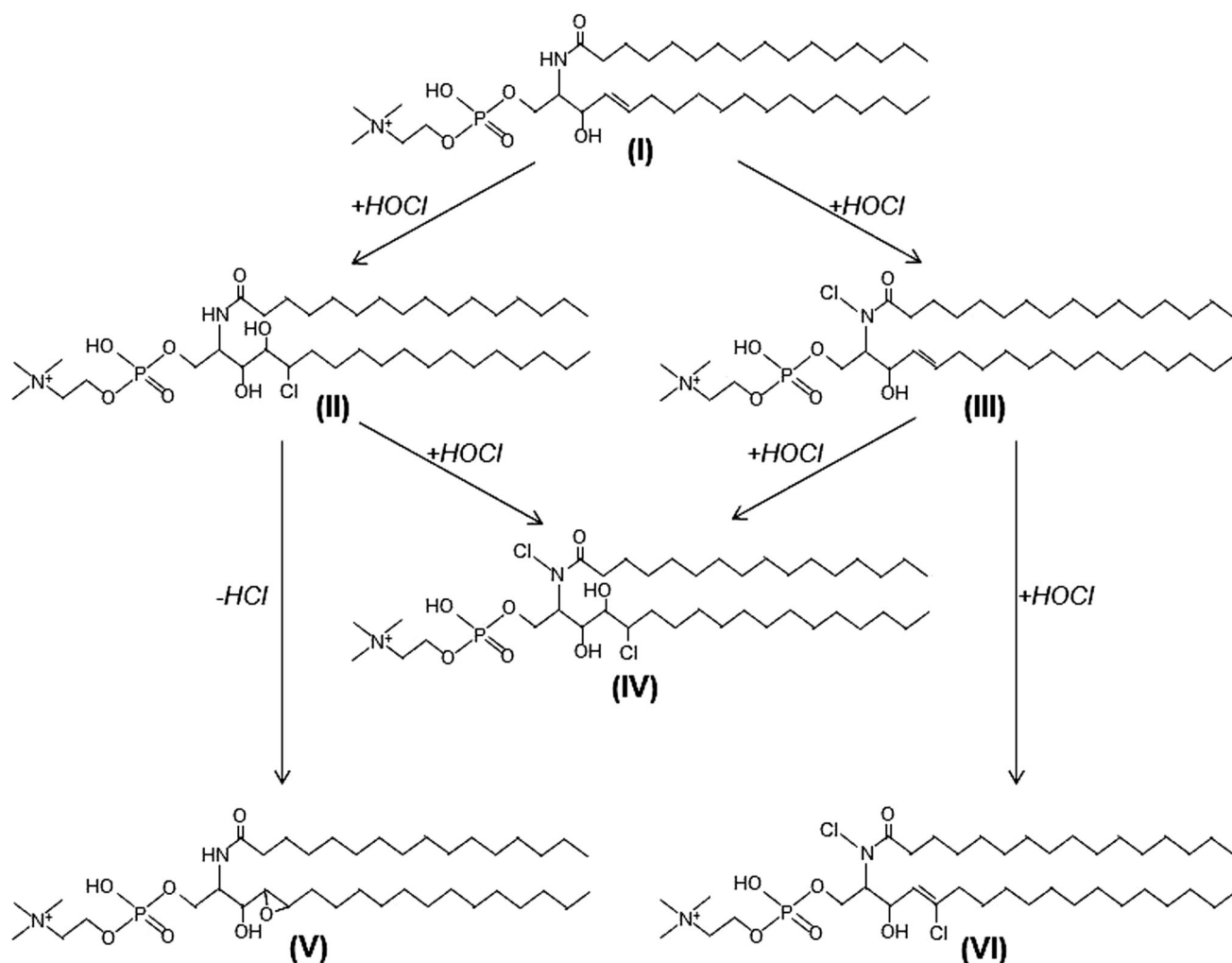


Fig. 4. Sphingomyelin modification by HOCl.

These results demonstrate that by producing HOCl/HOBr, MPO mimics the function of phospholipase through the formation of chloro- or bromohydrins from polyunsaturated phospholipids, followed by spontaneous hydrolysis of the ester bond with the generation of lysophospholipids. The latter are also formed as a result of cleavage of the alkenyl ether bond in plasmalogen under the action of HOCl/HOBr (Fig. 3). Using mass spectrometry and  $^{31}\text{P}$ -NMR, it was shown that the content of lysophosphatidylcholine increased when LDLs were incubated in the presence of HOCl [89]. These reactions can have biological significance. On one hand, it is known that LDLs contain lysophosphatidylcholine in sufficiently large quantities ( $\sim 50 \mu\text{g}$  per mg protein), and its content in LDLs increases with atherosclerosis progression [90, 91]. On the other hand, lysophosphatidylcholine is involved in the regulation of the activity of platelets and endothelial cell, which makes it an atherothrombotic molecule [92].

The second most prevailing phospholipid in LDLs after phosphatidylcholine ( $\sim 900 \mu\text{g}$  per mg protein) is sphingomyelin ( $\sim 400 \mu\text{g}$  per mg protein). Sphingomyelin is the only human phospholipid that does not contain glycerol but has the amino alcohol sphingosine. The fatty acid in sphingomyelin is attached to the sphingosine amino group, while the phosphocholine polar group is attached to the sphingosine hydroxyl group (Fig. 4, (I)). The RHS targets in the sphingomyelin molecule are the sphingosine double bond and the hydrogen atom of the amide bond. The rate constant of the HOCl reaction with sphingomyelin was estimated as  $18.7 \pm 3.05 \text{ M}^{-1}\cdot\text{s}^{-1}$  [93], which is faster than the reaction between HOCl and unsaturated bonds of acyl chains ( $0.56 \text{ M}^{-1}\cdot\text{s}^{-1}$  [94]). Using MALDI-TOF-mass spectrometry, it was shown that HOCl added as a reagent or produced by MPO transforms sphingomyelin into its chlorine derivative as a result of HOCl addition to the sphingosine double bond (Fig. 4, stage (II)) or substitution of the amide bond

hydrogen atom with the chlorine atom (Fig. 4, stage (III)). Under the action of two HOCl molecules, both reactions can proceed with the formation of compound (IV) (Fig. 4). Epoxide (V) resulting from chlorohydrin dehydrochlorination (Fig. 4) and chlorine derivative  $-\text{CH}=\text{CCl}-$  (VI) formed by the replacement of hydrogen atom of the  $-\text{CH}=\text{CH}-$  double bond with chlorine (Fig. 4) were minor products. Traces of (V) and (VI) were also detected in the reaction of unsaturated bonds of phospholipids acyl chains with primary RHS [78, 81, 82]. Chlorinated sphingomyelin derivatives produce significant effects on the cells, including initiation of reactive oxygen species production, uncoupling of the mitochondrial transmembrane potential, apoptosis induction, and DNA damage [93].

In cholesterol, the only target for hypohalous acids is the double bond at the 5,6 position of the steroid nucleus. Using thin-layer chromatography, NMR, and mass spectrometry, it was found that adding HOCl to the cholesterol-containing phospholipid liposomes leads to the formation of chlorine derivatives, mainly  $\alpha$ - and  $\beta$ -isomers of chlorohydrins [95, 96]. The same products were formed in LDLs treated with HOCl or in the presence of  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  [97]. Addition of catalase and HOCl scavengers prevented formation of such products, indicating involvement of MPO and HOCl in the reaction [95]. Unlike chlorohydrins of aliphatic hydrocarbon chains, cholesterol chlorohydrins are unstable and convert into epoxides by dehydrochlorination. Hydrolysis and further oxidation of  $\alpha$ - and  $\beta$ -epoxides leads to the formation of cholesterol hydroxy- and keto- derivatives [98, 99]. The reaction of HOBr with the cholesterol double bond proceeds similarly to analogous reaction with HOCl but requires smaller amounts of HOBr compared to HOCl [100].

The free radical mechanism of the unsaturated lipid modification by RHS is described in detail in reviews [25, 101]. Here, we discuss only the results confirming the existence of this mechanism in LDLs. The first works on the LPO induced in LDLs by RHS and  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  were conducted in early 1990s [102, 103]. The following studies have shown that LDL incubation with primary RHS or in the presence of  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  results in the formation of typical primary molecular products, such as diene conjugates and hydroperoxides, usually detected by traditional methods of LPO induction [102, 104], secondary products of carbonyl nature, that react with 2-thiobarbituric acid [101-103, 105-108], and end products that fluoresce in the visible light (e.g., Schiff bases). The latter are formed by the interaction of oxidized lipids with proteins (reaction (7)) [105, 108]. Free radical traps, such as  $\alpha$ -tocopherol and butylated hydroxytoluene, in micromolar concentrations completely blocked the RHS-induced LPO [42, 106], which confirmed its free-radical mechanism. By penetrating into the LDL lipid phase, HOCl caused degradation of lipid-

soluble antioxidants (carotenoids,  $\alpha$ -tocopherol), reducing LDL resistance to LPO [42, 71, 109, 110]. RHS can initiate LPO through the reaction of HOCl or HOBr with hydroperoxides (rate constants, 10.8 and  $8.9 \text{ M}^{-1}\text{s}^{-1}$  for HOCl and HOBr, respectively [111]). Hydroperoxides are always present in trace amounts in unsaturated lipid [112] and can form peroxy and alkoxy radicals in the reaction with hypohalous acids [111, 113-120]. Elevated levels of LDLs containing LPO products have been found in the blood of CVD patients [121, 122].

It is known that some compounds promote LPO in LDLs. For example, hemoglobin modified by HOCl increases accumulation of LPO products in LDLs by a factor of  $\sim 2$  compared to the native protein. The increase in the prooxidant capacity of hemoglobin after incubation with HOCl is associated with heme destruction, release of  $\text{Fe}^{2+}$  ion, and its subsequent oxidation by HOCl to  $\text{Fe}^{3+}$  [123]. The last reaction generates the  $\cdot\text{OH}$  radical, an extremely efficient LPO inducer [124].

Nitrite also promotes HOCl-induced LPO in LDLs [42] through the formation of nitrile chloride ( $\text{NO}_2\text{Cl}$ ) in reaction (2) with further decomposition into free radicals in reaction (3). Addition of butylated hydroxytoluene (free radical trap) to the reaction mixture significantly inhibited accumulation of LPO products in LDLs that indicates the radical mechanism of peroxidation in the presence of  $\text{HOCl} + \text{NO}_2^-$  [42]. A mixture of  $\text{HOCl} + \text{NO}_2^-$  causes tyrosyl radical formation [44] in proteins. Tyrosyl radicals are also formed in the peroxidase cycle of MPO by Tyr oxidation with Compounds I and II (Fig. 2). Tyrosyl radical is a known inducer of LPO free radical reactions in LDLs [125]. Another mechanism of the MPO-mediated LPO activation in LDLs is not associated with the formation of HOCl and  $\text{NO}_2\text{Cl}$ . It is  $\text{NO}_2^-$  oxidation to the  $\text{NO}_2\cdot$  radical in the MPO peroxidase cycle [44].

$\text{NH}_2$ -groups of Lys also contribute to the HOCl-induced LPO in LDLs by reacting with HOCl with the formation of Lys chloramine that decomposes in reaction (5) to the N-centered radical. The latter initiates LPO in LDLs [104]. Similar processes occur in the reaction of  $\text{NH}_2$ -group of phosphatidylethanolamine with HOCl, which also results in the formation of N-centered radical [126].

*Reactions of phospholipid polar heads with RHS.* The polar heads of phospholipids in LDLs are mainly choline phosphate, ethanolamine phosphate, and serine phosphate. Neither HOCl nor HOBr react at a noticeable rate with the quaternary nitrogen of phosphatidylcholine [71, 72, 75, 77, 78, 82, 83, 127, 128]. In contrast, the  $\text{NH}_2$ -groups of ethanolamine phosphate and serine phosphate easily react with HOCl and HOBr with the formation of chlor- and bromamines, respectively (the table). Chloramines of phosphatidylethanolamine and phosphatidylserine differ considerably in their stability due to presence of carboxyl group in the position next to the  $\text{NH}_2$ -group in serine phosphate. Such chloramines

quickly decompose with the formation of aldehydes. Figure 5 shows the conversion of phosphatidylethanolamine and phosphatidylethanolamine amino groups in the interaction with HOCl. The latter transforms phosphatidylethanolamine into mono- and then dichloramine [126, 129] which undergoes dehydrochlorination accompanied by the homolytic cleavage of the N–Cl bond with the formation of the N-centered radical (Fig. 5) [126]. The difference in the mechanisms of conversion of phosphatidylethanolamine and phosphatidylserine chloramines led to the hypothesis on the MPO-dependent regulation of LPO of unsaturated phospholipids [126]. It was found that the presence of phosphatidylethanolamine in unsaturated phosphatidylcholine liposomes contributes to the accumulation of LPO products under the action of HOCl due to the formation of N-centered radical acting as an LPO inducer. In contrast, inclusion of phosphatidylserine in the phosphatidylcholine liposomes inhibited HOCl-induced LPO as a result of decrease in HOCl concentration due to its interaction with the amino groups. In this case, the generated products are incapable of LPO initiation (Fig. 5 and [126]).

The reaction of unsaturated phosphatidylethanolamine with HOCl or HOBr involves the attack on the polar head  $\text{NH}_2$ -groups and the  $-\text{CH}=\text{CH}-$  bonds of acyl chains. The  $\text{NH}_2$ -groups enter the reaction with HOCl first, and only after all amines are converted to

dichloramines (Fig. 5), chlorohydrin formation begins [130], that is due to a significant difference (by 3–4 orders of magnitude) in the rate constants of the HOCl reactions with the  $\text{NH}_2$ -group and unsaturated  $-\text{HC}=\text{CH}-$  bond of phosphatidylethanolamine (the table and [58]). Formation of bromamines and bromohydrins in the reaction of unsaturated phosphatidylethanolamine with HOBr starts almost simultaneously [130]. The rate constants of these reactions differ insignificantly (the table and [59]).

**Modification of carbohydrates.** Carbohydrates are found in LDLs as components of glycolipids and ApoB100. ApoB100 is a glycoprotein containing N-acetylglucosamine, galactose, mannose, and sialic acid at a molar ratio of  $\sim 4 : 2 : 5 : 2$  [131]. ApoB100-associated glycoconjugates are represented by two main types: bi-antennal sialylated chains and highly mannosylated chains containing N-acetylglucosamine. The ApoB100 molecule contains 5–6 highly mannosylated and 8–10 sialylated conjugates bound to the protein through Asn residues (according to [132]). The carbohydrate composition of the LDL lipid fraction differs from the composition of protein-associated carbohydrates. Thus, LDL glycolipids contain glucose and N-acetylgalactosamine, but not mannose [131].

Primary RHS either do not react with sugars lacking nitrogen atoms in their molecules (e.g., ribose, glucose,

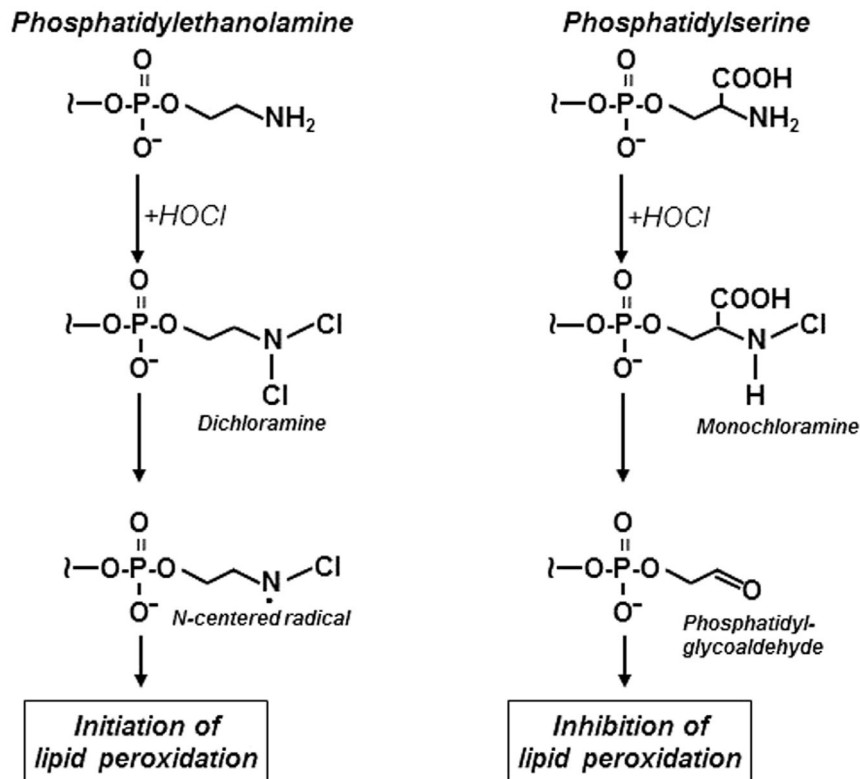


Fig. 5. Modification of phosphatidylethanolamine and phosphatidylserine by HOCl formed in the MPO-catalyzed reaction.

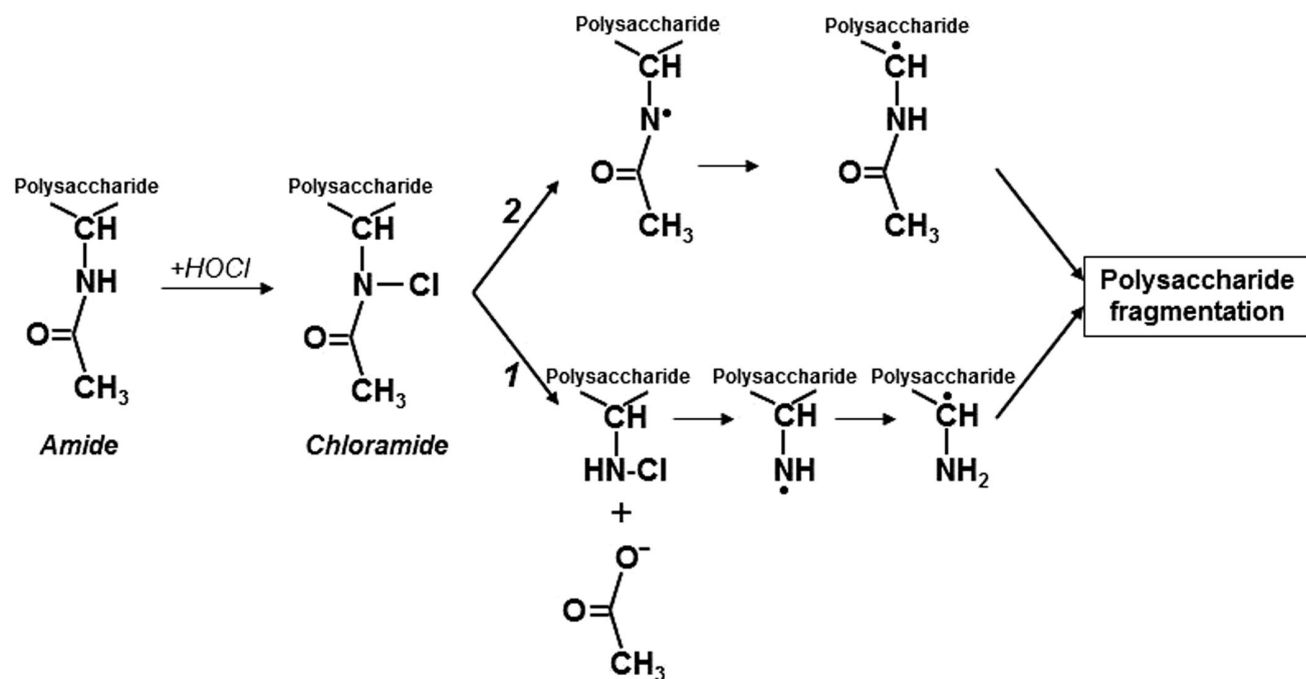


Fig. 6. HOCl reaction with N-acetylated amino sugar.

glucuronic acid) or react extremely slowly with them [133, 134]. At the same time, RHS react at a high rate with N-acetylated amino sugars [41, 134]. N-acetylglucosamine, N-acetylgalactosamine, and sialic acids are the RHS targets in ApoB100 and LDL glycolipids. It was established that at the first stage of the reaction, the N-acetamide group hydrogen atom is replaced by a halogen atom with the formation of chlor- or bromamide. Figure 6 shows the reaction of N-acetylated amino sugars with HOCl. The resulting chloramide is slowly hydrolyzed to monochloramine and acetate (reaction 1 in Fig. 6) [135] or transformed to the N-centered amidyl radical as a result of homolytic cleavage of the N–Cl bond (reaction 2 in Fig. 6) [136, 137]. As mentioned above (reaction 1 in Fig. 6 and reaction (3)), the N–Cl bond in monochloramine undergoes homolytic cleavage with the formation of N-centered radical. Then, both N-centered radicals are isomerized to C-centered radicals on the C-atom in the second position of the ring and on the C-atom of the nearby ring, which leads to the ring fragmentation of the polysaccharide chain [135–137]. Similar processes were observed in the reaction of glycosaminoglycans with HOBr [138]. Ions of transition metals contribute to the homolytic cleavage of the N–Cl bond [137, 138]. Superoxide anion radical significantly accelerates the break of the N–Cl bond presumably due to the reduction of transition metal ions [139, 140].

It was suggested that desialylation transforms LDLs into their pro-atherogenic form [141, 142]. Desialylated LDLs are actively captured by subendothelial cells, turning them into the foam cells [20]. Desialylated LDLs were

isolated from the blood of patients with atherosclerosis [20, 141]. The authors of this hypothesis suggest that LDL desialylation is catalyzed by the *trans*-sialidase enzyme [143]. The results described in this section indicate that at least partially, the cleavage of glycoconjugates from the LDLs can occur under the action of RHS formed in the MPO-catalyzed reactions.

**RHS interaction with antioxidants.** LDLs contain many lipid-soluble antioxidants. On average, 1 mol of LDL contains 6.37 mol of  $\alpha$ -tocopherol, 0.93 mol of  $\gamma$ -tocopherol, 0.1 mol of ubiquinol-10 (reduced coenzyme Q10), and carotenoids, such as  $\beta$ -carotene (0.53 mol),  $\alpha$ -carotene (0.22 mol), lycopene (0.29 mol), cryptoxanthin (0.25 mol), and trace amounts of lutein, anhydrolutein, zeaxanthin, and canthaxanthin [110, 144]. All these antioxidants are destroyed to a different degree by adding HOCl to the LDLs. First, ubiquinol-10 disappears, then lycopene, then  $\alpha$ -tocopherol and  $\beta$ -carotene [145]. Comparison of carotenoids (*cis*- and *trans*-lycopenes,  $\alpha$ - and  $\beta$ -carotenes) and their oxy-derivatives (zeaxanthin,  $\alpha$ - and  $\beta$ -cryptoxanthins, *cis*- and *trans*-anhydroluteins) showed that, the former are more efficient HOCl scavengers. The content of these carotenoids in LDLs is decreased by 25% by treatment with 36–52 HOCl molecules per one LDL particle [110]. However, considering the number of antioxidant molecules per LDL particle shows that 50% oxidation of  $\alpha$ -tocopherol required ~125-fold, ubiquinol-10 and lycopene ~2000-fold, and  $\beta$ -carotene ~2500-fold molar excess of HOCl. The second-order rate constants of the reaction of Trolox ( $\alpha$ -tocopherol water-soluble analog), ubiquinol-0

(ubiquinol-10 analog), and  $\beta$ -carotene with HOCl are significantly lower than those for the reaction of functional groups of some amino acid residues (the table and [55, 58, 146]).

Nitrite promote the HOCl-induced degradation of  $\alpha$ -tocopherol and  $\beta$ -carotene in LDLs [42], which might be explained by the generation of  $\cdot\text{NO}_2$  and  $\text{Cl}\cdot$  radicals in reaction (2) and (3). These radicals react with  $\alpha$ -tocopherol and  $\beta$ -carotene faster than HOCl [42].

The rate constants of the similar reaction of Trolox and ubiquinol-0 with HOBr are  $\sim 50$  and  $\sim 2000$  times higher, respectively, than with HOCl [59]. Even at the HOBr/LDL molar ration of 2000, only 0.4% of HOBr reacts with antioxidants, while the rest HOBr molecules react with lipids (56.6%) and proteins (43.0%) [59]. Therefore, it is unlikely that antioxidants are the major targets of RHS. This suggestion was confirmed by the study [147] that demonstrated that neither  $\sim 8$ -fold enrichment nor  $\sim 6$ -fold depletion of LDLs with  $\alpha$ -tocopherol produced effect on the HOCl-induced modification of Trp and Lys residues of ApoB100.

#### MODIFICATION OF LDL STRUCTURE IN THE REACTIONS WITH RHS

The reactions of RHS with the functional groups of ApoB100, lipids, antioxidants, and carbohydrates in LDLs are unavoidably accompanied by changes in the physicochemical properties and structural organization of the LDL particles. Spin-labeled analogues of stearic acid with a paramagnetic fragment at the 5th, 12th, or 16th C-atoms were used to assess the changes in the physicochemical properties of the phospholipid monolayer on the LDL surface. These probes made it possible to obtain the information on the physicochemical characteristics of the lipid phase at different distances from the LDL particle surface. Adding HOCl to the LDLs caused a decrease in the mobility of the phospholipid fatty acid chains and, at the same time, increased the polarity of their microenvironment up to the 12th  $\text{CH}_2$ -group of the fatty acid chain (which corresponded to the depth of  $\sim 1.7$  nm from the LDL surface). Only at the HOCl concentration exceeding 1 mM, significant changes in the studied parameter were observed in the LDL deeper regions (in the region of the 16th  $\text{CH}_2$ -group) [148].

HOCl-induced changes in the ApoB100 structure were studied using the spin-tag 3-maleimidoproxil covalently linked to the protein SH-groups. The EPR spectrum of LDLs labeled with this spin tag represented a superposition of two signals, wide and narrow, corresponding to the strongly and weakly immobilized label, respectively, which indicated the presence of two populations of SH-groups in the LDL, with higher and lower mobility. Adding HOCl to the LDLs decreased the fraction of strongly immobilized spin label and increased the

fraction of weakly bound label, suggesting an increase in the mobility of ApoB100 labeled sites upon reaction with HOCl [149].

LDL particles are characterized by the negative surface potential of  $-16$ – $-17$  mV [150, 151]. Using electrophoresis and EPR with paramagnetic  $\text{Mn}^{2+}$  cations, it was shown that reaction with primary RHS increases the negative surface charge and electrophoretic mobility of LDLs [145, 147, 151]. This can be mainly due to the transformation of positively charged amino acid residues Lys, His, and Arg [55, 56], as well as  $\text{NH}_2$ -groups of the ethanolamine phosphate and serine phosphate of phospholipids located at the surface of LDL particles [58, 59] into halogenamines or other neutral products. The alterations in the local charge distribution on the LDL surface can induce aggregation of LDL particles. Incubation of LDLs with HOCl or  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  is accompanied by an increase in the average LDL particle size, as shown in many *in vitro* experiments [152, 153]. Cultured smooth muscle cells from the intima of healthy human aorta accumulate cholesterol from HOCl-modified LDLs [153]. As a rule, lipid accumulation by the cells increases with an increase in the average size of LDL particles [17, 20, 142]. Similar spherical lipid inclusions with a diameter of 100–400 nm have been repeatedly found in the arterial intima affected by atherosclerosis [154, 155].

Three possible mechanisms for the MPO-induced formation of intermolecular crosslinks in ApoB100, which can be the cause of the LDL particle aggregation, were described in the previous chapter. Hazell et al. showed [60] that LDL aggregation in the human blood induced by HOCl or  $\text{MPO} + \text{H}_2\text{O}_2 + \text{Cl}^-$  was accompanied by the increase in the content of carbonyl compounds and simultaneous disappearance of Lys residues. When the amino groups of Lys in ApoB100 in LDLs were preventively blocked by reducing methylation, no aggregation of LDL particles was observed. It was found that LDL aggregation occurs according to the first described mechanism (reactions (4), (6) and (7) in the previous chapter) through the conversion of Lys chloramines into aldehydes with subsequent formation of Schiff bases type intermolecular crosslinks.

High-molecular-weight protein complexes resistant to heat denaturation in the presence of sodium dodecyl sulfate and 1,4-dithiothreitol were found in the human blood LDLs after reaction with HOCl. Such resistance indicated formation of covalent but not disulfide crosslinks most likely via the first mechanism (see previous chapter). An increase in the HOCl concentration was accompanied by the increase in the content of di-Tyr (third mechanism, previous chapter). However, the observed amount of di-Tyr was not enough to explain all the existing intermolecular crosslinks in the LDLs [156].

The possibility of formation of the intra- and intermolecular S–S and S–N crosslinks in LDLs by the second mechanism (reaction (8) in the previous chapter) was

demonstrated using synthetic peptides containing Cys and Lys residues [66]. Incubation of these peptides with HOCl resulted in the generation of intra- and intermolecular sulfene amides according to reaction (8), which were then oxidized to sulfine amides and sulfone amides. Hence, the observed S–N crosslinks were formed by the HOCl-mediated interaction between the SH-groups of Cys and NH<sub>2</sub>-groups of Lys. Disulfides were also found among the products of sulfenyl chloride reaction with the SH-groups of Cys (reaction (8) in previous chapter) [66].

Therefore, RHS formed in the MPO-catalyzed reaction can modify the physicochemical properties and structural organization of the surface proteolipid layer of LDL particles and stimulate LDL aggregation and their transformation into proatherogenic form.

### INTERACTION BETWEEN MPO AND LDL

Given that MPO is extremely polycationic ( $pI \sim 10$ ) and LDL surface is negatively charged [150, 157], it can be assumed *a priori* that they are capable of interacting with each other. The interaction between MPO and LDLs was for the first time demonstrated by co-precipitation with phosphotungstate and studying the electrophoretic mobility of the coprecipitates [158]. The presence of the cationic detergent cetrimide, as well as an increase in the NaCl concentration to 0.3 M or pH decrease to 3.6, prevented the interaction, which suggests the electrostatic nature of MPO binding to LDLs [159]. The binding of MPO to the surface of LDLs and very low-density lipoproteins (VLDLs) was confirmed by electrophoresis, gel filtration, and photon correlation spectroscopy. It was determined that the average diameter of MPO–LDL and MPO–VLDL complexes was  $28.0 \pm 1.9$  and  $85.9 \pm 8.9$  nm, respectively [160].

The absence of lipids in the contact area between MPO and LDL was demonstrated using spin-labeled derivatives of fatty acid that incorporated into the LDL phospholipid monolayer and provided positioning of the paramagnetic center at different distances from the surface. MPO interaction with LDLs did not significantly affect the parameters of the EPR spectra of these probes, which indicated that the interaction between MPO and LDLs does not involve lipids. Studying the distribution of paramagnetic Mn<sup>2+</sup> cations between the LDL surface and water environment in the dependence of MPO adding demonstrated that the interaction between MPO and LDLs does not lead to significant changes in the negative surface charge of LDL particle. These results suggest that the contact area between MPO and LDL is extremely small and does not involve phospholipids, the main carriers of negatively charged groups on the LDL surface [159].

Interestingly, MPO did not interact with high-density lipoproteins (HDLs). Since LDLs and VLDLs differ from HDLs in the ApoB100 content on their surface, we

believe that MPO binds to this protein. The fact that antibodies against ApoB100 displace MPO from the complex with LDLs or VLDLs confirms this hypothesis [160]. Analysis of the ApoB100 amino acid sequence for the potential MPO-binding sites enriched with negatively charged residues and lacking positively charged amino acids identified three peptides: <sup>1</sup>EEEMLEN<sup>7</sup>, <sup>53</sup>VELEV<sup>59</sup>Q<sup>59</sup>, and <sup>445</sup>EIQDDCTGDED<sup>456</sup>. We demonstrated that only the <sup>445</sup>EIQDDCTGDED<sup>456</sup> peptide disrupted the MPO–LDL complex and, therefore, could represent the MPO-binding site in ApoB100. The specific interaction of this peptide with MPO was also confirmed by gel filtration. Therefore, <sup>445</sup>EIQDDCTGDED<sup>456</sup> causes dissociation of the MPO–LDL complex by interacting with MPO and not with the LDL [159].

The association of MPO with the LDL surface is accompanied by the increase in the enzyme activity [63], which contributes to the LDL surface modification by RHS formed in the MPO-catalyzed reactions. This modification reduces the affinity of ApoB100 for the ApoB- and ApoE-receptors and increases the affinity of mLDLs for the scavenger receptors of macrophages and monocytes. The latter leads to the progressive accumulation of intracellular cholesterol and its esters and transformation of macrophages and monocytes into foam cells [161].

The plasma protein ceruloplasmin (CP) interacts with MPO and inhibits both peroxidase and halogenating activities of the enzyme [162–164]. CP retains its ability to inhibit MPO even when MPO is associated with the LDL surface and forms the CP/MPO/LDL(VLDL) [160]. We showed that the <sup>445</sup>EIQDDCTGDED<sup>456</sup> peptide (which uncouples the interaction of MPO with LDL), as well as physiological modulators of the MPO halogenating activity, such as thiocyanate and CP, decreased the accumulation of cholesterol by monocytes and macrophages in the presence of LDLs modified by HOCl and HOBr [161].

Based on the data obtained, we propose a mechanism for the participation of MPO and produced RHS, as well as the <sup>445</sup>EIQDDCTGDED<sup>456</sup> peptide and inhibitors/modulators of MPO halogenating activity (CP and thiocyanate), in the proatherogenic modification of LDLs, activation of monocytes, and formation of foam cells (Fig. 7). The binding of MPO to LDL leads to the site-specific modification of the latter (1 in Fig. 7). LDL-Cl (LDL-Br) stimulates leukocytes (2 in Fig. 7), provokes respiratory burst, and MPO release by neutrophils and monocytes (3 in Fig. 7). Secreted MPO binds to the LDLs and promotes their damage, thus closing the “vicious circle” of the LDL-Cl (LDL-Br) generation. mLDLs in a complex with MPO are captured by monocytes and macrophages, leading to the accumulation of cholesterol in these cells and their transformation into foam cells (4 in Fig. 7). Inhibitors/modulators of the MPO halogenating activity (CP and thiocyanate), as well

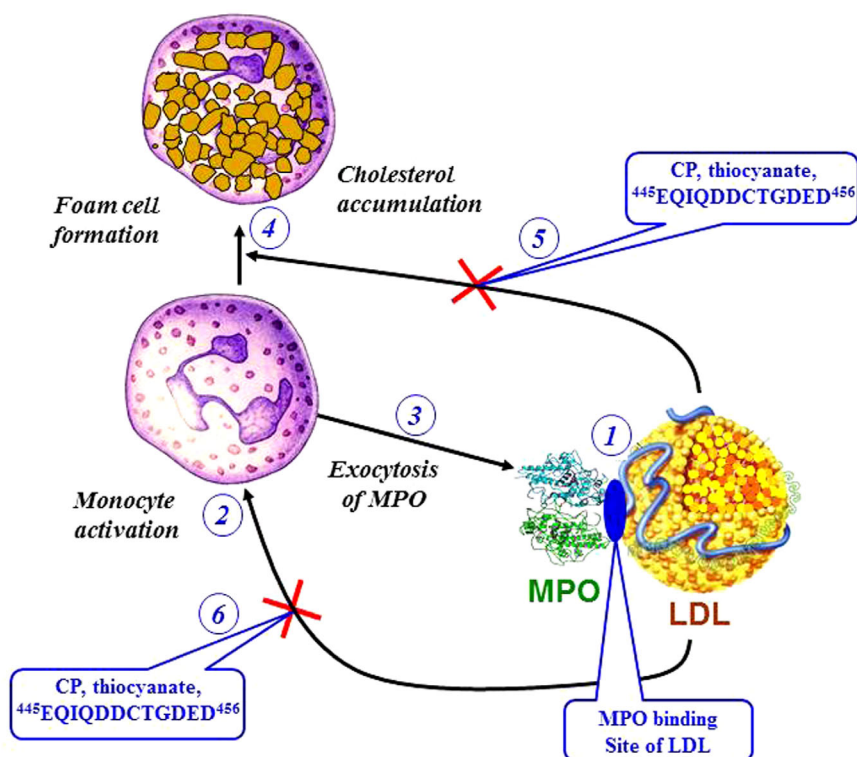


Fig. 7. Participation of MPO,  $^{445}\text{EQIQDDCTGDED}^{456}$  peptide, and inhibitors/modulators of MPO halogenating activity (CP and thiocyanate) in the atherogenic modification of LDLs, activation of monocytes, and formation of foam cells.

as the inhibitor of the LDL–MPO complex  $^{445}\text{EQIQDDCTGDED}^{456}$ , prevent the oxidative/halogenative modification of LDLs. On one hand, this reduces LDL uptake and accumulation of intracellular cholesterol (5 in Fig. 7); on the other hand, it breaks the vicious circle of monocyte (neutrophil) activation and LDL–Cl (LDL–Br) formation (6 in Fig. 7).

These results allow to make an important addition to the scheme describing the role of mLDLs in the formation of atherosclerotic plaques (Fig. 1). The binding of MPO to LDLs and proatherogenic modification of LDL surface start already in the bloodstream. The transformation of LDLs into the proatherogenic form is aggravated after migration of the MPO–LDL complexes into the subendothelial space. This hypothesis is confirmed by detection of ApoB100-containing lipoprotein complexes with MPO in the blood of patients with atherosclerosis [160].

#### BIOMARKERS OF MPO ACTIVITY AND HALOGENATIVE STRESS IN CARDIOVASCULAR DISEASES

Epidemiological studies of recent years have shown that high MPO concentration in the human blood is associated with an increased risk of CVD, regardless of classi-

cal risk factors, such as troponin T and C-reactive protein (reviews [23, 165]). Indeed, an increase in the MPO concentration or activity in the blood plasma [166–168], leukocytes [166], and atherosclerotic plaques [169], as well as in the complexes with ApoB100-containing LDLs and VLDLs, has been repeatedly registered in CVD patients [160]. As a rule, an increase in the concentration and/or activity of MPO was accompanied by the accumulation of specific biomarkers in the patient's blood and tissues.

The “ideal” biomarker of the MPO-induced protein modification is 3-chlorotyrosine. This highly stable product is formed in the body by the HOCl reaction with tyrosine. 3-Chlorotyrosine can be detected by various combinations of methods, such as chromatography, mass spectrometry, NMR, etc. The elevated levels of 3-chlorotyrosine have been registered in the human blood plasma during myocardial infarction [170, 171], atherosclerotic plaques and LDLs isolated from them [172, 173], HDLs [174, 175] and apolipoprotein A-I isolated from HDLs and aorta from CVD patients [176], and cardiac tissue homogenate from individuals with atrial fibrillation [177]. It is important to note that the amount of MPO [170, 171] and, often, the severity of the disease [176] correlated with the recorded 3-chlorotyrosine levels.

HOCl/HOBr-modified proteins can be detected by ELISA using monoclonal antibodies. These antibodies

are very specific and sensitive to the RHS-modified epitopes and do not bind to proteins modified by aldehydes or  $\text{Cu}^{2+}$  ions generated in the oxidative stress [178]. RHS-modified proteins were found in atherosclerotic plaques in the aorta [179] and iliac artery [180]. In the latter case, the level of RHS-modified protein correlated with the content of MPO and ApoB100 in the vessel and with the extent of its damage [180].

Chloro- and bromohydrins of aliphatic chains of lipids are specific markers of lipid damage in the halogenative stress. These compounds are very stable and can accumulate in the body in detectable amounts, since the concentration of unsaturated bonds of aliphatic chains is high. The disadvantage of these biomarkers is a relatively low rate of their formation (the table). Using liquid chromatography/mass spectrometry (LC-MS), the authors [181] registered the formation of chlorohydrins (but not hydroperoxides) of oleic, linoleic, and arachidonic acids in the phospholipid fraction of LDLs after incubation with HOCl. Messner et al. [87] detected lysophosphatidylcholine chlorohydrins in human arterial endothelial cells after their treatment with HOCl using the ESI-MS method. Later, the same authors found a 60-fold increase in the content of lysophosphatidylcholine chlorohydrin in human atherosclerotic plaques compared with the tissue unaffected by atherosclerosis [182]. An alternative approach for detecting tissue damage by RHS formed in the MPO-catalyzed reactions is the use of monoclonal antibodies against chlorohydrins of oleic acid [183].

Aliphatic 2-chloroaldehydes formed in the HOCl reaction with plasmalogen are other important markers of the RHS-induced lipid damage (Fig. 3). These compounds are less stable but can be detected in very low concentrations by mass spectrometry. 2-Chlorohexadecanal was found to accumulate in the rat heart during experimental myocardial infarction [184]. More than a 1400-fold increase in the 2-chlorohexadecanal content was detected in the atherosclerotic plaques of human aorta compared with the aorta of healthy donors [91]. 2-Chloraldehyde and lysophosphatidylcholine are formed as result of the plasmalogen double bond break (Fig. 3). The 20- and 34-fold increase in the content of lysophosphatidylcholine with arachidonic and linoleic acids, respectively, was observed in the atherosclerotic plaques compared with the non-diseased aorta [91].

Halogen derivatives of purine and pyrimidine bases can play the role of markers of the MPO-dependent degradation of nucleotides and nucleic acids. Thus, a 10-fold increase in the level of 5-chlorouracil, which is formed in the body by the MPO-dependent mechanism, was found in the atherosclerosis-damaged areas of human aorta [185]. The amount of 5-bromouracil was higher in the atherosclerotic plaques compared with the unaffected areas of the aorta, but this difference was statistically insignificant [185].

Recently, we obtained specific antibodies against halogenated LDLs (LDL-Cl and LDL-Br) and chlorinated CP (CP-Cl) [186, 187]. Using these antibodies, we were able to detect LDL-Cl, LDL-Br, as well as CP-Cl in the blood of CVD patients by an improved ELISA method. Interestingly, the content of LDL-Cl, LDL-Br, and CP-Cl in the patients' blood correlated with the MPO level registered by ELISA and with the MPO halogenating activity ( $r \sim 0.80$ ). A high value of positive correlation was found for LDL-Cl and LDL-Br ( $r = 0.91$ ;  $p < 0.0001$ ), although they were not associated with the peroxidase activities of either plasma or MPO [186]. These results confirm an important role of halogenative stress in the atherogenic modification of LDLs. Halogenated LDLs and CP can be used as potential biomarkers of halogenative stress in CVD.

The results presented in this review demonstrate an important role of halogenative stress in atherogenic modification of LDLs. The key element of the LDL modification is the site-specific binding of MPO to the LDL surface. Inhibition of MPO or dissociation of the MPO-LDL complex prevents MPO-dependent modification of LDLs, accumulation of cholesterol in the vascular cells, and early stages of atherosclerosis development. It has become obvious that one of the most important tasks in preventing and slowing down atherosclerosis is the development of tools and approaches for reduction of the MPO-dependent production of RHS. There are at least three different approaches. The first one is searching for methods that would limit the release of MPO during activation and degranulation of neutrophils. For example, this effect was found for the glucocorticoids  $\alpha$ -methylprednisolone and hydrocortisone [188]. The second approach is inhibition of the MPO halogenating activity, e.g., by MPO inhibition with CP [163, 164] or switching the enzyme to the peroxidase cycle by changing the medium [189]. The third approach is scavenging and neutralization of already formed RHS. Another possibility to prevent or to slow down atherosclerosis is the use of efficient, non-toxic components capable of disrupting the MPO-LDL complex, like the  $^{445}\text{EQIQDDCTGDED}^{456}$  peptide that mimics the MPO-binding site of ApoB100 on the LDL surface. Identification of new specific, sensitive, and informative biomarkers of MPO activity and halogenative stress is of particular importance, since it can provide the information for the development of modern methods for early diagnostics of atherosclerosis and CVDs. Further clarification of issues discussed in our review will allow to develop fundamentally new approaches aimed at identification and timely prevention of halogenative stress, and hence, atherosclerosis.

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