# Regulation of the Structural Stability of Erythrocytes by Hydrogen Peroxide: Mathematical Model and Experiment

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Abstract—In this study, the regulatory mechanisms induced by extracellular hydrogen peroxide were analyzed on the basis of a mathematical model that considers the key stages of the formation of methemoglobin and ferrylhemoglobin, as well as their binding to the erythrocyte membrane. Numerical modeling has shown that reversible binding of methemoglobin to the membrane is an adaptive mechanism aimed at stabilizing the lipid bilayer of the membrane. On the other hand, an increase in the concentration of ferrylhemoglobin and its binding to the membrane leads to an increase in pathophysiological processes that reduce the structural stability of cells. The quantity of methemoglobins and ferrylhemoglobins formed depends on the concentration of extracellular hydrogen peroxide and exposition time, the number of cells in the sample, the state of the antioxidant system of erythrocytes, the metabolic activity of cells and external metabolic conditions. Based on numerical modeling, optimal conditions (oxidant concentration and exposition time) have been determined, under which the activation of adaptive processes occurs. Experiments with erythrocyte hemolysis in vitro have shown that hydrogen peroxide at concentrations of 10–200  $\mu$ M causes an increase in the structural stability of the membrane and a decrease in the proportion of hemolyzed erythrocytes.

**Keywords:** erythrocytes, hydrogen peroxide, methemoglobin, adaptation, hormesis **DOI:** 10.1134/S1990747822010093

### **INTRODUCTION**

The formation of reactive oxygen species (ROS) is an important phenomenon in cellular respiration, since it results in a wide range of physiological and pathophysiological consequences. Until recently, the exclusively pathological role of ROS was the dominant point of view; the formation of ROS was associated with the development of chronic and degenerative diseases. However, it is known nowadays that ROS are participants in numerous physiological processes, including regulation of metabolism, activation of adaptive mechanisms, triggering apoptosis, etc. [1–4].

According to modern concepts, hydrogen peroxide is the main molecule of the ROS group involved in regulatory processes [2]. Basic features of the signals transmitted by hydrogen peroxide are the threshold concentration of activation and exposure duration. The effect of such signals is reversible, as evidenced by the presence in the cell of a special set of enzymatic systems, including catalase (CAT), glutathione peroxidase (GPX), peroxyredoxin (PRX), etc. [2, 5– 7]. A moderate increase in the concentration of hydrogen peroxide activates mechanisms associated with protection and adaptation [2–4]. These mechanisms control the balance of hydrogen peroxide formation and utilization, and violation of this balance leads to the development of various diseases [1, 3]. Thus, the effect of hydrogen peroxide on cells at low concentrations leads to adaptation and increased protection, while at high concentrations it causes damage to cellular structures. Such a property of living systems, described by a change in the sign of the biological effect with an increase in the activity/amount of the stressor, is termed hormesis and is explained by the presence of specific compensatory and adaptive mechanisms [8, 9].

Transcription factor Nrf2 (nuclear E2-related factor 2) plays the key role in the adaptation processes of most cells, the activity of which is regulated with the participation of the redox-dependent protein Keap1 (Kelch-like ECH-associating protein 1) [10, 11]. However, this system of maintaining redox homeostasis is absent in erythrocytes, and the nature of activation of adaptive processes of erythrocytes under oxidative stress remains unknown.

Erythrocytes are the most plentiful cells in the human body that perform a variety of functions related to homeostasis [9, 12, 13]. Changes in the functional state of the body, including those associated with a violation of redox homeostasis, would affect the state of erythrocytes; therefore, it is necessary to study the protective mechanisms of these cells and find ways to regulate their functioning under oxidative stress. Normally, the concentration of hydrogen peroxide in the blood does not exceed 10  $\mu$ M [14]. In pathophysiological processes, the extracellular concentration of hydrogen peroxide can increase up to 200–250  $\mu$ M [15, 16]. The study of mechanisms of the adaptive response of erythrocytes to hydrogen peroxide in the concentration range from 10 to 250  $\mu$ M was the purpose of this work.

Here we examined the role of hydrogen peroxide as a stimulating agent that increases the structural stability of human erythrocytes during oxidative hemolysis mediated by hypochlorous acid (HOCl). The cell destruction caused by HOCl is due to a colloidal osmotic mechanism with the formation of pores of high permeability [17]; therefore, an increase in the structural stability of the membrane may decrease the proportion of non-hemolyzed cells.

One of the most important mechanisms for increasing the structural stability of membranes is their interaction with the main protein of erythrocytes, hemoglobin [9]. Hemoglobin is a regulator of erythrocyte homeostasis and a key participant in intracellular oxidative processes [9, 12]. Binding of hemoglobin to the membrane regulates various intracellular processes, such as energy metabolism, membrane deformability, eryptosis, etc. Binding of hemoglobin with membrane proteins occurs due to covalent and non-covalent interactions and binding of hemoglobin with membrane lipids involves hydrophobic interactions [18–21]. The rate constants of formation and disruption of the hemoglobin-membrane complex depend on the state of hemoglobin [19, 21]. Binding constant of interaction of methemoglobin (MetHb), formed due to the oxidation of oxyhemoglobin, with the membrane components is the highest, suggesting a significant role of this reaction in regulating the stability of the membrane. The study of the molecular mechanisms of this interaction has shown that in the case of insignificant and reversible oxidation hemoglobin exerts a stabilizing effect on membranes, while the formation of ferrylhemoglobins (FerHB) under the oxidative stress impairs the interaction of the membrane with cytoskeleton [22]. Thus, the regulation of the oxidative metabolism of hemoglobin may underlie the hormesis effect produced by oxidants on erythrocytes.

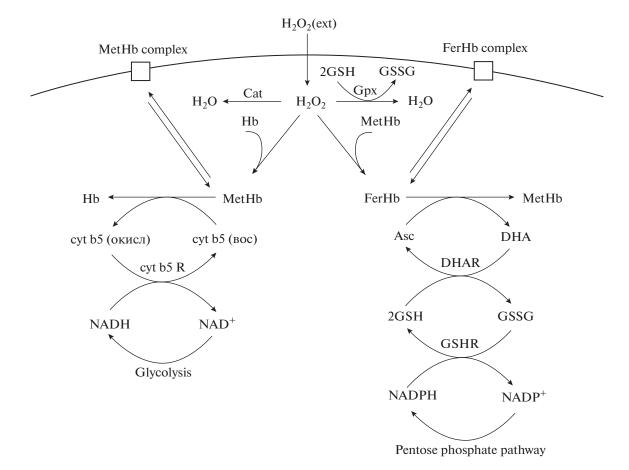
Confirmation of this hypothesis can be carried out by constructing a mathematical model of intracellular processes based on known data. The beginning of the study of redox metabolism of erythrocytes using mathematical modeling can be attributed to the 60s of the last century. Nicholls described the processes of transport and utilization of hydrogen peroxide based on diffusion and decomposition of the oxidant by catalase [23]. In the early 2000s, modeling of hydrogen peroxide metabolism in erythrocytes was supplemented by complex processes associated with the oxidation of hemoglobin and its transition to oxidized forms [6]. At the same time, Kinoshita et al. proposed models for reduction processes, which considered the contribution of several ways of reducing oxidized forms of hemoglobin [24]. The energy for the reduction can be replenished only through glycolysis processes, the first models of which were developed in the 1980s in the groups of Rappoport-Yakobash [25] in Germany and of Ataullakhanov [26] in the USSR. In the following decades, mathematical models of glycolysis were upgraded and refined [24, 27]. In this study, the effect of pre-incubation of cells with hydrogen peroxide on the structural stability of erythrocytes during oxidative hemolysis was explored and a quantitative contribution of participants in hemoglobin metabolism to the regulation of the structural stability of membranes was assessed.

#### DESCRIPTION OF THE MODEL

A mathematical model was created describing the key stages of the formation of methemoglobin and ferrylhemoglobin and their binding to the membrane under the action of extracellular hydrogen peroxide. The model includes the processes of diffusion and utilization of hydrogen peroxide in the cell, the formation and reduction of methemoglobin and ferrylhemoglobin, the formation and disruption of the hemoglobin– membrane complex and the formation of reducing equivalents due to glycolysis. The general scheme of these processes is shown in Fig. 1. All equations and rates are presented in Table 1. Constants and initial conditions are given in Tables 2 and 3.

The rate of change in the concentration of extracellular hydrogen peroxide depends on its diffusion into cells and is described by Fick's first law for membranes [33]. The diffusion of the oxidant into the cell and its utilization causes a decrease in the extracellular concentration of  $H_2O_2(v_1)$ , which is directly proportional to the number of cells in the sample, and a change in the concentration of hydrogen peroxide in the intracellular medium ( $v_2$ ).

In the cell, hydrogen peroxide is reduced to water with the participation of enzymes Cat and Gpx [7]. At high concentrations of hydrogen peroxide, Cat is the main participant in the  $H_2O_2$  utilization, as the rate constant of this reaction is higher than that of other enzymes and at the same time it does not require additional substrates. Upon interaction with a molecule of hydrogen peroxide, a Cat molecule enters the Comp I ( $v_3$ ) state; upon interaction with the next oxidizer molecule, it returns to its original state ( $v_4$ ). Gpx ( $v_5$ ) is an additional participant in the utilization; it catalyzes the reduction of hydrogen peroxide by intracellular glutathione (GSH), which is also used for the reduction of ascorbic acid (Asc) [31].



**Fig. 1.** The processes of erythrocyte metabolism with the participation of hydrogen peroxide considered in the model. Hb, hemoglobin; MetHb, methemoglobin; FerHb, ferrylhemoglobin;  $H_2O_2$ , hydrogen peroxide;  $H_2O_2$  (ext), extracellular hydrogen peroxide; Cat, catalase; Gpx, glutathione peroxidase; GSH, glutathione; GSSG, glutathione disulfide; cyt b5 (ox), oxidized form of cytochrome b5; cyt b5 (red), reduced form of cytochrome b5; cyt b5 R, cytochrome b5 reductase; NAD<sup>+</sup>/NADH, oxidized and reduced forms of nicotinamide adenine dinucleotide; Asc, ascorbic acid; DHA, dehydroascorbate; DHAR, dehydroascorbate reductase; GSHR, glutathione reductase; NADP<sup>+</sup>/NADPH, oxidized and reduced forms of nicotinamide adenine dinucleotide phosphate; MetHb complex, membrane complex with methemoglobin; FerHb complex, membrane complex with ferrylhemoglobin.

The oxidation of hemoglobin with the formation of methemoglobin ( $v_6$ ) [24, 34–36] and the oxidation of methemoglobin with the formation of ferrylhemoglobin ( $v_7$ ) are competing processes involving hydrogen peroxide [6]. The rates of these processes, according to the law of acting masses, are proportional to the concentration of participants.

It was shown that at elevated concentrations of methemoglobin, the main way of its reduction is interaction with cytochrome b5 ( $v_8$ ), which is then reduced by NADH with the participation of the cytochrome b5 reductase (cyt b5 R) ( $v_9$ ) [24]. The reduction of ferrylhemoglobin is carried out by cellular antioxidants without the participation of enzymes. The model considers the reaction of ferrylhemoglobin with Asc, as its reaction constant has the highest value ( $v_{10}$ ). When ascorbate is reduced, radicals are formed that rapidly dismutate to dehydroascorbate (DHA) in a direct reaction or with the participation of enzymes [37].

In this study, a complete model of glycolysis and the pentose phosphate pathway was constructed, the equations of which is presented in the Annex. The main contribution to the regulation of the NAD<sup>+</sup>/NADH redox state is associated with reactions catalyzed by glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and lactate dehydrogenase (LDH) [24, 32, 38]. The first reaction is one of the stages of glycolysis; it catalyzes the oxidation of glyceraldehyde-3-phosphate (gap) with the formation of 1,3diphosphoglycerate (13pg) and the release of NADH. The second catalyzes the conversion of pyruvate (pyr) to lactate (lac) with the release of NAD<sup>+</sup>. Under physiological conditions, this reaction proceeds in the direction of the formation of lactate and NAD<sup>+</sup> using the electrons released in the first reaction. Due to this, the ratio of concentrations [pyruvate] : [lactate] = 1 : 15 and of [NADH] :  $[NAD^+] = 1 : 600$  is observed in erythrocytes [24, 32]. When the NADH concentration

No.	Reaction	Velocity equation
1	$H_2O_2(ext) \rightarrow H_2O_2$	$v_1 = -P(H_2O_2 - H_2O_2(ext))\frac{S_{cell}}{V}N_{cell}$
2	$H_2O_2(ext) \rightarrow H_2O_2$	$v_2 = -P(H_2O_2 - H_2O_2(ext))\frac{S_{cell}}{V_{cell}}$
3	$H_2O_2 \xrightarrow{Cat} H_2O$	$v_3 = k_3 \cdot \text{Cat} \cdot \text{H}_2\text{O}_2$
4	$H_2O_2 \xrightarrow{CompI} H_2O + O_2$	$v_4 = k_4 \cdot \text{Compl} \cdot \text{H}_2\text{O}_2$
5	$H_2O_2 + 2GSH \xrightarrow{Gpx} 2H_2O + GSSG$	$v_5 = \frac{\text{Gpx} \cdot \text{GSH} \cdot \text{H}_2\text{O}_2}{K_1 \cdot \text{H}_2\text{O}_2 + K_2 \cdot \text{GSH}}$
6	$2Hb + H_2O_2 \rightarrow 2MetHb + H_2O$	$v_6 = k_6 \cdot \text{Hb} \cdot \text{H}_2\text{O}_2$
7	$MetHb + H_2O_2 \rightarrow FerHb + H_2O$	$v_7 = k_7 \cdot \text{MetHb} \cdot \text{H}_2\text{O}_2$
8	$MetHb + cytb5(red) \rightarrow Hb + cytb5(ox)$	$v_8 = k_8 \cdot \text{MetHb} \cdot \text{cytb5(red)}$
9	NADH + cytb5(ox) $\xrightarrow{cytb5R}$ NAD <sup>+</sup> + cytb5(red)	$v_{9} = k_{9} \cdot \text{cytb5R} \frac{\text{NADH}}{K_{M}^{\text{NADH}} + \text{NADH}} \times \frac{\text{cytb5(ox)}}{K_{M}^{\text{cytb5(ox)}} + \text{cytb5(ox)}}$
10	$FerHb + 2Asc \rightarrow MetHb + DHA + Asc$	$v_{10} = k_{10} \cdot \text{FerHb} \cdot \text{Asc}$
11	$DHA + 2GSH \xrightarrow{DHAR} Asc + GSSG$	$v_{11} = \frac{k_{13} \cdot \text{DHAR}}{1 + \frac{K_{\text{DHA}}}{\text{DHA}} + \frac{K_{\text{GSH}}}{\text{GSH}}}$
12	$GSSG + NADPH \xrightarrow{GSHR} NADP^+ + 2GSH$	$v_{12} = \frac{\text{GSHR} \cdot \text{GSSG} \cdot \text{NADPH}}{K_1 \cdot \text{NADPH} + K_2 \cdot \text{GSH}}$
13	$MetHb + Band3 \rightarrow MetHbcomplex$	$v_{13} = k_{13} \cdot \text{MetHb} \cdot \text{Band3}$
14	MetHbcomplex $\rightarrow$ MetHb + Band3	$v_{14} = k_{14} \cdot \text{MetHbcomplex}$
15	$FerHb + Band3 \rightarrow FerHbcomplex$	$v_{15} = k_{15} \cdot \text{FerHb} \cdot \text{Band3}$
16	$FerHbcomplex \rightarrow FerHb + Band3$	$v_{16} = k_{16} \cdot \text{FerHbcomplex}$

Table 1. Reactions and velocity equations in the model

is decreased, the equilibrium is violated; this leads to a reverse reaction catalyzed by LDH to restore the balance.

Regulation of the Asc/DHA redox pair is carried out with the participation of the electrons of GSH by dehydroascorbate reductase (DHAR) ( $v_{11}$ ) [31]. As a result of this reaction, glutathione disulfide (GSSG) is formed, which is reduced by glutathione reductase (GSHR) ( $v_{12}$ ) with the participation of the electrons from NADPH. NADPH is reduced in the pentose phosphate pathway with the participation of glucose-6-phosphate dehydrogenase (G6PDH) and phosphogluconate dehydrogenase (G06PDH) [33, 38].

The formation of methemoglobin leads to its interaction with the erythrocyte membrane, namely, binding to the band 3 protein (Band 3) [12, 20]. To simulate this process, the kinetics of ligand-receptor binding was considered [33]. Methemoglobin acts as a ligand, and the membrane Band 3 protein serves as a receptor. The formation of the complex is reversible and is characterized by the constants of formation and decomposition of the complex ( $v_{13}$  and  $v_{14}$ ). Excess of hydrogen peroxide cause the formation of ferrylhemoglobin, which is also able to bind to the membrane of erythrocytes. This process was considered similarly to the previous one with the rates ( $v_{15}$  and  $v_{16}$ ).

The constructed mathematical model was nonlinear and contained 41 differential equations. Considering the dynamics of the process was a fundamentally new approach, which made it possible to study not

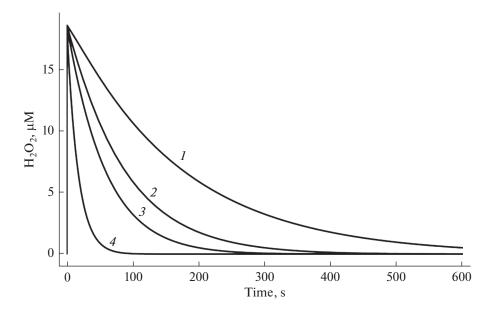
Table 2. Parameter values in the model

Reaction constants from Table 1	Reference	Reagent
		designation
$P = 6 \times 10^{-6} \text{ m s}^{-1}$ S <sub>cell</sub> = 1.09 × 10 <sup>-10</sup> m <sup>2</sup>	[28, 29]	$H_2O_2$ (ext)
$V = 10^{-6} \text{ m}^3$		$H_2O_2$
$N_{\rm cell} = 3 \times 10^7$		Cat
$P = 6 \times 10^{-6} \mathrm{m  s^{-1}}$	[28, 29]	Comp I
		Hb
$v_{\text{cell}} = 6.3 \times 10^{-11} \text{ m}^2$		MetHb
$k_3 = 6 \times 10^6 \mathrm{M}^{-1} \mathrm{s}^{-1}$	[6]	FerHb
$k_4 = 1.6 \times 10^7 \text{ M}^{-1} \text{ s}^{-1}$	[6]	cyt b5(red)
$Gnx = 1.4 \times 10^{-6} M$	[7]	cyt b5(ox)
$K_1 = 2.5 \times 10^{-5} \mathrm{M s}$		NADH
$K_2 = 2.4 \times 10^{-8} \text{ M s}$		$NAD^+$
$k_6 = 100 \text{ M}^{-1} \text{ s}^{-1}$	[6]	Asc
$k_7 = 98 \text{ M}^{-1} \text{ s}^{-1}$	[30]	DHA
		GSH
$k_8 = 6200 \text{ M}^{-1} \text{ s}^{-1}$	[24]	GSSG
$k_9 = 418 \text{ s}^{-1}$	[24]	NADPH
•		NADP <sup>+</sup>
14		Band 3
$\mathbf{K}_M = 1.3 \times 10^{-1} \mathrm{M}$		complex MetH
$k_{10} = 400 \text{ M}^{-1} \text{ s}^{-1}$	[30]	complex FerH
$k_{11} = 5.27 \text{ s}^{-1}$	[31]	
		only the amp nals. All con
		from literatur
		the system watche system wa
	[7]	matica softwa is presented i
$K_1 = 2.5 \times 10^{-5} \text{ M s}$ $K_2 = 2.5 \times 10^{-5} \text{ M s}$		Numerica
$k = 3 \mathrm{M}^{-1} \mathrm{s}^{-1}$	[12]	of methemog
15		made it possi of extracellul
$k_{14} = 7.8 \times 10^{-5} \mathrm{s}^{-1}$	[12]	vation of ada allowed expe
$k_{15} = 30 \text{ M}^{-1} \text{ s}^{-1}$	[21]	effect of hydroiden of membrane
$k_{16} = 0.8 \times 10^{-5} \mathrm{s}^{-1}$	[21]	with erythroc acid.
	$V = 10^{-6} \text{ m}^{3}$ $N_{cell} = 3 \times 10^{7}$ $P = 6 \times 10^{-6} \text{ m s}^{-1}$ $S_{cell} = 1.09 \times 10^{-10} \text{ m}^{2}$ $V_{cell} = 6.3 \times 10^{-17} \text{ m}^{3}$ $k_{3} = 6 \times 10^{6} \text{ M}^{-1} \text{ s}^{-1}$ $k_{4} = 1.6 \times 10^{7} \text{ M}^{-1} \text{ s}^{-1}$ $Gpx = 1.4 \times 10^{-6} \text{ M}$ $K_{1} = 2.5 \times 10^{-5} \text{ M s}$ $k_{2} = 2.4 \times 10^{-8} \text{ M s}$ $k_{6} = 100 \text{ M}^{-1} \text{ s}^{-1}$ $k_{7} = 98 \text{ M}^{-1} \text{ s}^{-1}$ $k_{8} = 6200 \text{ M}^{-1} \text{ s}^{-1}$ $k_{9} = 418 \text{ s}^{-1}$ $cytb5R = 7 \times 10^{-8} \text{ M}$ $K_{M}^{\text{NADH}} = 3.1 \times 10^{-7} \text{ M}$ $K_{M}^{\text{Cytb5(ox)}} = 1.5 \times 10^{-5} \text{ M}$ $k_{10} = 400 \text{ M}^{-1} \text{ s}^{-1}$ $k_{11} = 5.27 \text{ s}^{-1}$ $DHAR = 10^{-7} \text{ M}$ $K_{GSH} = 3.5 \times 10^{-3} \text{ M}$ $GSHR = 1.4 \times 10^{-6} \text{ M}$ $k_{1} = 2.4 \times 10^{-8} \text{ M s}$ $k_{2} = 2.5 \times 10^{-5} \text{ M s}$ $k_{13} = 3 \text{ M}^{-1} \text{ s}^{-1}$ $k_{14} = 7.8 \times 10^{-5} \text{ s}^{-1}$ $k_{15} = 30 \text{ M}^{-1} \text{ s}^{-1}$	$V = 10^{-6} \text{ m}^3$ $N_{cell} = 3 \times 10^7$ $P = 6 \times 10^{-6} \text{ m s}^{-1}$ $[28, 29]$ $S_{cell} = 1.09 \times 10^{-10} \text{ m}^2$ $V_{cell} = 6.3 \times 10^{-17} \text{ m}^3$ $k_3 = 6 \times 10^6 \text{ M}^{-1} \text{ s}^{-1}$ $[6]$ $k_4 = 1.6 \times 10^7 \text{ M}^{-1} \text{ s}^{-1}$ $[6]$ $k_4 = 1.6 \times 10^7 \text{ M}^{-1} \text{ s}^{-1}$ $[6]$ $Gpx = 1.4 \times 10^{-6} \text{ M}$ $[7]$ $K_1 = 2.5 \times 10^{-5} \text{ M}$ s $K_2 = 2.4 \times 10^{-8} \text{ M}$ s $k_6 = 100 \text{ M}^{-1} \text{ s}^{-1}$ $[6]$ $k_7 = 98 \text{ M}^{-1} \text{ s}^{-1}$ $[30]$ $k_8 = 6200 \text{ M}^{-1} \text{ s}^{-1}$ $[24]$ $k_9 = 418 \text{ s}^{-1}$ $[24]$ $cytbSR = 7 \times 10^{-8} \text{ M}$ $K_M^{NDH} = 3.1 \times 10^{-7} \text{ M}$ $K_M^{MDH} = 3.1 \times 10^{-7} \text{ M}$ $K_{M}^{Cos)} = 1.5 \times 10^{-5} \text{ M}$ $k_{10} = 400 \text{ M}^{-1} \text{ s}^{-1}$ $[30]$ $k_{11} = 5.27 \text{ s}^{-1}$ $[31]$ DHAR = $10^{-7} \text{ M}$ $K_{GSH} = 3.5 \times 10^{-3} \text{ M}$ $K_{0HA} = 2.1 \times 10^{-6} \text{ M}$ $[7]$ $k_{12} = 2.5 \times 10^{-5} \text{ M}$ s $k_{13} = 3 \text{ M}^{-1} \text{ s}^{-1}$ $k_{13} = 3 \text{ M}^{-1} \text{ s}^{-1}$ $[12]$ $k_{14} = 7.8 \times 10^{-5} \text{ s}^{-1}$ $[12]$

	*	
Reagent designation	Initial value	Reference
$H_2O_2$ (ext)	$0-2.5 \times 10^{-4} \mathrm{M}$	This study
$H_2O_2$	0	This study
Cat	$5.5 \times 10^{-6} { m M}$	[7]
Comp I	$5.5 \times 10^{-6} { m M}$	[7]
Hb	0.01 M	[24]
MetHb	0	[24]
FerHb	0	[30]
cyt b5(red)	$8.12 \times 10^{-7} \text{ M}$	[24]
cyt b5(ox)	0	[24]
NADH	$1.48 \times 10^{-7} \mathrm{M}$	[32]
$NAD^+$	$8.9 \times 10^{-5} \mathrm{M}$	[32]
Asc	$7.5 \times 10^{-5} \mathrm{M}$	[1]
DHA	0	[1]
GSH	$1.5 \times 10^{-3} \mathrm{M}$	[7]
GSSG	0	[7]
NADPH	$5 \times 10^{-5} \mathrm{M}$	[7]
NADP <sup>+</sup>	$0.2 \times 10^{-6} \mathrm{M}$	[7]
Band 3	$2.6 \times 10^{-5} \mathrm{M}$	[33]
complex MetHb	0	[12]
complex FerHb	0	[21]

only the amplitudes, but also the duration of the signals. All constants and initial conditions were taken from literature sources. An analysis of the stability of the system was performed. The numerical solution of the system was performed with the Wolfram Mathematica software. The full form of differential equations is presented in the Annex.

Numerical modeling of the processes of interaction of methemoglobin with the erythrocyte membrane made it possible to determine the concentration range of extracellular hydrogen peroxide, at which the activation of adaptive processes would be observed; this allowed experimental verification of the model. The effect of hydrogen peroxide on the structural stability of membranes was studied in the experiments in vitro with erythrocyte hemolysis induced by hypochlorous acid.



**Fig. 2.** Kinetics of utilization of intracellular hydrogen peroxide in cuvettes with different number of cells: cuvette (1),  $1 \times 10^7$  cells; cuvette (2),  $2 \times 10^7$  cells; cuvette (3),  $3 \times 10^7$  cells; and cuvette (4),  $10^8$  cells.

#### MATERIALS AND METHODS

Sodium hypochlorite (NaOCl) (Sigma-Aldrich) and hydrogen peroxide (Belmedpreparaty, Belarus) were used in the experiments.

The concentration of the NaOCl solution was determined spectrophotometrically as the concentration of OCl<sup>-</sup> at pH 12.0, which has the molar extinction coefficient ( $\epsilon_{292}$ ) of 350 M<sup>-1</sup> cm<sup>-1</sup>. Since pK<sub> $\alpha$ </sub> for HOCl is approximately of 7.5 at physiological pH, half of the compound was present in the protonated form of HOCl and half in the deprotonated form of OCl<sup>-</sup>. Thus, HOCl was a mixture of HOCl/OCL<sup>-</sup>. A working solution of HOCl was prepared immediately before analysis by dissolving the preparation in a 10 mM sodium-phosphate buffer (pH 7.4) containing 137 mM NaCl.

In the experiments, the blood of healthy donors obtained at the Republican Scientific and Practical Center of Transfusiology and Medical Biotechnologies was used. Erythrocytes were isolated and washed by centrifugation at 300 g in a phosphate buffered saline (PBS) containing 10 mM Na<sub>2</sub>HPO<sub>4</sub>/KH<sub>2</sub>PO<sub>4</sub>, 137 mM NaCl, 2.7 mM KCl, and 5 mM *D*-glucose (pH 7.4).

The kinetics of erythrocyte hemolysis was measured using a Solar CM-2203 spectrofluorimeter (SOLAR, Belarus) by recording the optical density of the cell suspension ( $3 \times 10^7$  cells/mL) at 640 nm. The concentration of erythrocytes was chosen in such a way that the initial optical density was of 0.5, which was optimal for measurements. A 1-mL aliquot of a suspension of washed erythrocytes was added to the optical cuvette and thermostated for 3-4 min at  $37^{\circ}$ C with constant stirring. Hemolysis was initiated by adding 0.2 mM HOCl to the suspension of erythrocytes in PBS after incubation with H<sub>2</sub>O<sub>2</sub> for 5 or 15 min at concentrations of  $10-250 \,\mu$ M.

The effect of the treatments (*I*) was evaluated as the relative difference of optical density after 10 min incubation with hypochlorous acid in the presence of hydrogen peroxide and optical density after 10 min incubation with hypochlorous acid without hydrogen peroxide.

The results were presented as mean values with standard deviation of the mean for three to five independent experiments. Statistical analysis was carried out using the Microsoft Excel software. The statistical significance of the difference between mean values was assessed using the Student's *t*-test, the significance level was taken as p < 0.05.

#### **RESULTS AND DISCUSSION**

The proposed model made it possible to calculate the dynamics of intracellular signaling processes involving methemoglobin and ferrylhemoglobin depending on the extracellular concentration of hydrogen peroxide. Numerical experiments based on the constructed model gave the following results.

Owing to diffusion and the action of catalase, a rapid (approximately for a second) establishment of the gradient of  $H_2O_2$  concentrations on the plasma membrane is generated. The intracellular concentra-

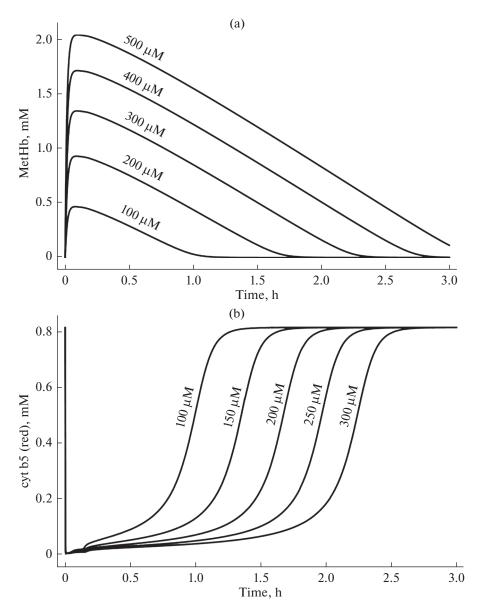


Fig. 3. Kinetics of formation and reduction of methemoglobin (MetHb) (a) and cytochrome b5 (red) (b) at different concentrations of extracellular hydrogen peroxide.

tion of hydrogen peroxide depends on the extracellular concentration, which decreases because of utilization of the  $H_2O_2$  by the cells. Utilization, and consequently, the dynamics of changes in extracellular and intracellular concentrations of hydrogen peroxide is proportional to the number of cells in the sample. Figure 2 shows the dynamics of intracellular hydrogen peroxide at an extracellular concentration of 200  $\mu$ M and at different number of erythrocytes in the sample. The consequence of the utilization of the oxidant is an exponential decrease in its intracellular and extracellular concentration. Figure 2 also shows that at a concentration of about 200  $\mu$ M and a cell count of  $3 \times 10^7$ ,

the total duration of the hydrogen peroxide signal does not exceed 300 s, and the intracellular concentration does not exceed 10% of the extracellular concentration, which is consistent with the data obtained in [23].

An increase in the intracellular concentration of hydrogen peroxide leads to an increase in the formation of methemoglobin (Fig. 3a), concentration of which reaches 2–3 mM. The obtained results were confirmed by the experimental studies [39]. Methemoglobin induces membrane stabilization and an increase in structural stability by forming complexes with cytoskeleton proteins [9, 40]. A decrease in the concentration of methemoglobin owing to its reduc-

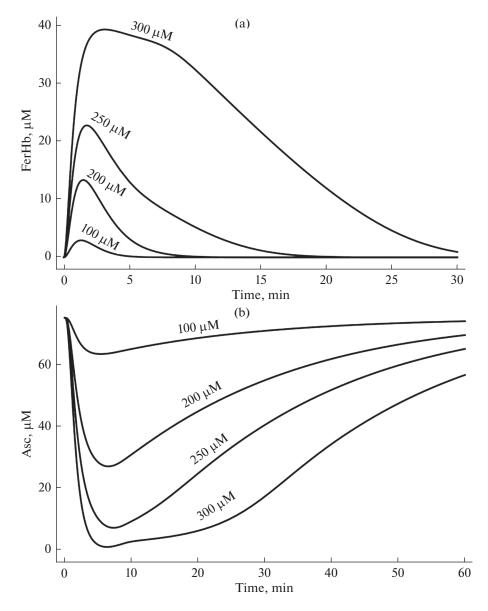
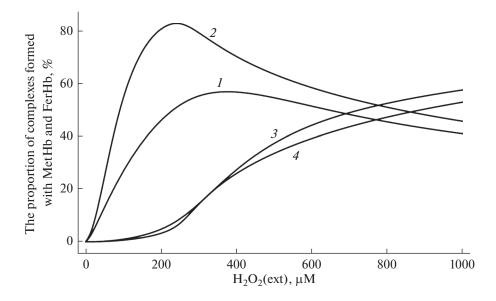


Fig. 4. Kinetics of formation and reduction of ferrylhemoglobin (FerHB) (a) and ascorbate (b) at different concentrations of extracellular hydrogen peroxide.

tion by cytochrome b5 leads to the breakdown of its membrane complexes. In the model, the total reduction time of methemoglobin for the  $H_2O_2$  concentrations of 200–300 µM is 2–3 h, which is comparable to the time presented in [39]. The dynamics of changes in the concentration of cytochrome b5 used for the reduction of oxidized hemoglobin depends on the concentration of  $H_2O_2$ ; at the  $H_2O_2$  concentrations exceeding 100 µM, the amount of reduced protein decreases to zero, which increases the time of reduction of methemoglobin (Fig. 3b). An increase in the rate of NADH reduction with an increase in extracellular lactate concentration (which can be observed with muscle exertion, hypoxia, and various pathological processes) will accelerate the process of methemoglobin reduction.

A further increase in the concentration of extracellular  $H_2O_2$  leads to the oxidation of methemoglobin and the formation of ferrylhemoglobin, which can also bind to the cell membrane. The amount of ferrylhemoglobin formed depends on the amount of hydrogen peroxide that entered the cells and the amount of methemoglobin formed. The paper [41] presented experimental data on the amount of ferrylhemoglobin at the ratio  $[H_2O_2]$  : [MetHb] = 10 : 1 and indicated complete oxidation of methemoglobin. In our experiments, this ratio did not rise above 1 : 1 and the model



**Fig. 5.** The proportion of methemoglobin–membrane complexes formed after 5-min (1) and 15-min (2) pre-incubation and the proportion of ferrylhemoglobin–membrane complexes after 5-min (3) and 15-min (4) pre-incubation depending on the concentration of extracellular hydrogen peroxide.

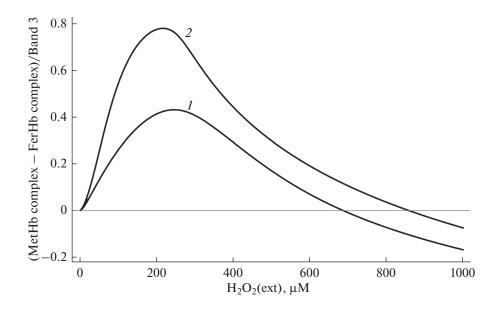


Fig. 6. Relative difference in the number of methemoglobin—membrane and ferrylhemoglobin—membrane complexes formed at different concentrations of extracellular hydrogen peroxide after 5-min (I) and 15-min (2) pre-incubation.

predicted the oxidation of 5% methemoglobin. The formation of a ferrylhemoglobin complex with the erythrocyte membrane disrupts interaction of cytoskeleton with the membrane, which leads to a decrease in the structural stability of the cell [21]. Figure 4a shows the dependence of the amount of ferrylhemoglobin formed on the concentration of extracellular hydrogen peroxide. Upon reaching 250  $\mu$ M of H<sub>2</sub>O<sub>2</sub>, an increase in the intensity of ferrylhemoglobin formation is observed, which is associated with depletion of the intracellular ascorbate pool (Fig. 4b). The model considering glycolysis shows that there would be a slowdown in the formation of NADH and NADPH under hypoglycemia, which leads to the

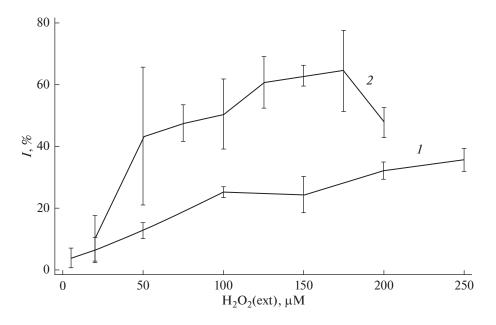


Fig. 7. The proportion of non-hemolysed erythrocytes depending on the concentration of extracellular hydrogen peroxide after 5-min (I) and 15-min (2) pre-incubation.

accumulation of oxidized forms of hemoglobin and an increase in the number of ferrylhemoglobin membrane complexes.

Within the framework of the proposed model of the adaptive mechanism, the degree of increase in the structural stability of the membrane depends on the number of membrane-methemoglobin and membrane-ferrylhemoglobin complexes formed. The proportion of receptors occupied by one or another protein, in turn, depends on the extracellular concentration of hydrogen peroxide and the incubation time. Figure 5 shows the theoretically calculated dependences of the fraction of the formed membrane complexes of proteins on the extracellular concentration of hydrogen peroxide. The dependence of the relative difference in the number of membrane-methemoglobin and membrane-ferrylhemoglobin complexes on the concentration of  $H_2O_2$  was bell-shaped (Fig. 6). At concentrations of hydrogen peroxide below 200 µM, an increase in the number of complexes with methemoglobin is observed in the cell with an increase in concentration. An increase in the incubation time from 5 to 15 min also leads to an increase in the number of membrane complexes formed by methemoglobin. At higher concentrations of the oxidant, the proportion of membrane complexes formed with methemoglobin decreases. Exceeding the concentration of 200 µM induces a sharp increase in the number of ferrylhemoglobin-membrane complexes, which is associated with a decrease in the number of reducing agent molecules. When the  $H_2O_2$  concentration reaches 800 µM, the number of complexes becomes equal.

Thus, according to this model, under the action of extracellular  $H_2O_2$  at concentrations from 10 to 200  $\mu$ M, a dose-dependent increase in the number of membrane-methemoglobin complexes, and consequently, an increase in the structural stability of the membrane is observed.

In the experimental study of the rate of erythrocyte hemolysis, it has been shown that pre-incubation of the cells with hydrogen peroxide at concentrations of 10-200 µM indeed reduced the proportion of hemolysed erythrocytes. The maximum protective effect, as predicted by the model (Fig. 6), depends on the incubation time of the cells with  $H_2O_2$ . Figure 7 shows the dependences of the proportion of unaffected erythrocytes on the extracellular concentration of hydrogen peroxide for incubation times of 5 and 15 min. With preliminary incubation of the cells for 5 min, a stabilizing effect was observed, reaching 30% of the control values. When the cells were incubated with hydrogen peroxide for 15 min, a more pronounced stabilizing effect was observed: the proportion of unaffected erythrocytes reached 60% for hydrogen peroxide at a concentration of 175 µM. However, at higher concentrations, the inverse relationship was manifested: with an increase in the concentration of  $H_2O_2$ , the proportion of unaffected erythrocytes decreased, which was probably due to an increase in the number of ferrylhemoglobin-membrane complexes. An increase in the number of these complexes could initiate the processes of lipid peroxidation and disrupt the interaction of the cytoskeleton with the membrane, which would

lead to a decrease of the structural stabilization effect at high concentrations of  $H_2O_2$ .

#### CONCLUSIONS

The construction of an adequate mathematical model of the adaptation mechanism of erythrocytes contributes to the analysis of optimal conditions for its regulation by external factors. In this study, a mathematical model has been constructed that considers the contribution of various participants in hemoglobin metabolism to the regulation of the structural stability of the membrane. The results of numerical modeling showed that reversible binding of methemoglobin to the membrane can be an adaptive mechanism aimed at stabilizing the lipid bilayer of the membrane. On the other hand, an increase in the concentration of an oxidized form of hemoglobin, ferylhemoglobin, and its binding to the membrane can lead to an increase in pathophysiological processes (lipid peroxidation, disruption of the interaction of the membrane with the cytoskeleton, etc.), which can reduce the structural stability of the cells. Apparently, the ratio of membrane complexes of various oxidized forms of hemoglobin determines the hormesis dependence of the response of erythrocytes to the action of hydrogen peroxide, that is, regulatory at low concentrations and damaging at high concentrations. To analyze optimal conditions, the model considered the extracellular concentration of hydrogen peroxide, the number of cells in the sample, the state of the erythrocyte antioxidant system (the content of ascorbate and other reducing agents), the metabolic activity of cells (glycolysis and pentose phosphate pathway), and external metabolic conditions (lactate content). As a result of theoretical and experimental studies, it has been shown that under the action of hydrogen peroxide at concentrations of 10-200 µM, the adaptive mechanism of erythrocytes was activated due to an increase in the number of membrane-bound methemoglobin, which increased the structural stability of the membrane under oxidative stress. The obtained data expand the understanding of the mechanisms of protection of erythrocytes and make it possible to predict the protective properties of the cells under oxidative stress.

#### ACKNOWLEDGMENTS

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#### COMPLIANCE WITH ETHICAL STANDARDS

The authors declare that they have no conflict of interest.

All procedures performed in studies involving human participants were in accordance with the ethical standards

of the 1964 Helsinki Declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all individual participants involved in the study.

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## Translated by E. Puchkov

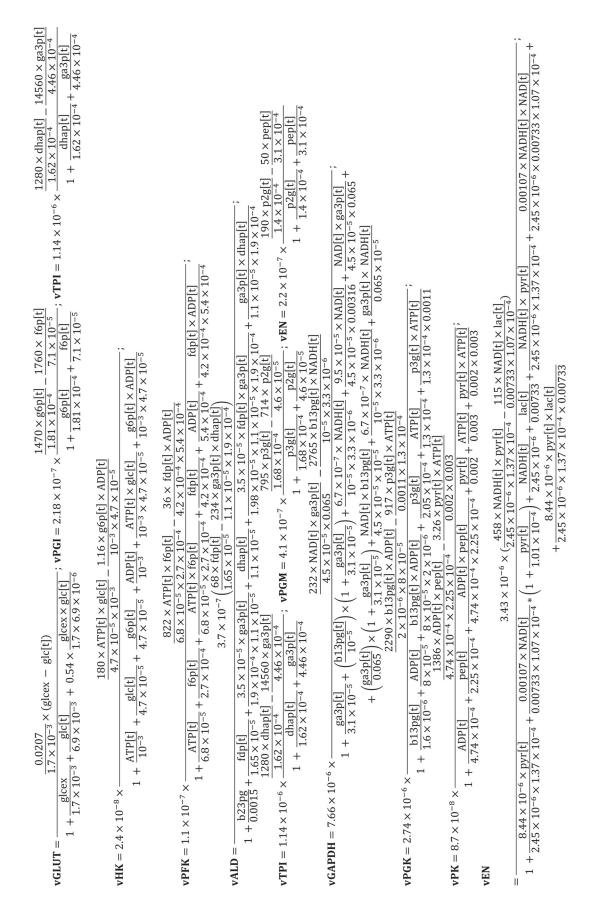
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# APPENDIX

The system of differential equations used in this study is presented below. In this form, the system was presented in the Wolfram Mathematica software.

\_

glc'[t] = vGLUT - vHK	$glc[0] = 5 \times 10^{-3}$
g6p'[t] = vHK - vPGI - vG6PDH	$g6p[0] = 6 \times 10^{-5}$
f6p'[t] = vPGI - vPFK + vTA + vTK2	$f6p[0] = 1.9 \times 10^{-5}$
fdp'[t] = vPFK - vALD	$fdp[0] = 5.6 \times 10^{-6}$
dhap'[t] = vALD - vTPI	dhap $[0] = 1.5 \times 10^{-5}$
ga3p'[t] = vALD + vTPI - vGAPDH - vTA + vTK1	$ga3p[0] = 3.6 \times 10^{-6}$
+ vTK2	
b13pg'[t] = vGAPDH - vPGK	$b13pg[0] = 2.3 \times 10^{-7}$
p3g'[t] = vPGK - vPGM	$p3g[0] = 4.8 \times 10^{-5}$
p2g'[t] = vPGM - vEN	$p2g[0] = 1.4 \times 10^{-5}$
pep'[t] = vEN - vPK	$pep[0] = 8.1 \times 10^{-6}$
pyr'[t] = vPK - vLDH - vpyrtr	$pyr[0] = 5.2 \times 10^{-5}$
lac'[t] = vLDH - vlactr	$lac[0] = 1.3 \times 10^{-3}$
ATP'[t] = -vHK - vPFK + vPGK + vPK	$ATP[0] = 1.4 \times 10^{-3}$
ADP'[t] = vHK + vPFK - vPGK - vPK	$ADP[0] = 1.5 \times 10^{-4}$
gl6p'[t] = vG6PDH - vPGLase	$gl6p[0] = 3.6 \times 10^{-4}$
go6p'[t] = vPGLase - v6PGODH	$go6p[0] = 4.5 \times 10^{-5}$
ru5p'[t] = v6PGODH - vX5PI - vR5PI	$ru5p[0] = 4.9 \times 10^{-6}$
x5p'[t] = vX5PI - vTK1 - vTK2	$x5p[0] = 9 \times 10^{-6}$
r5p'[t] = vR5PI - vTK1	$r5p[0] = 5.8 \times 10^{-6}$
s7p'[t] = vTK1 - vTA	$s7p[0] = 2.1 \times 10^{-5}$
e4p'[t] = vTA - vTK2	$e4p[0] = 4.5 \times 10^{-7}$
Cat1'[t] = -v3 + v4	$Cat1[0] = 5.5 \times 10^{-6}$
$\operatorname{Cat2'}[t] = v3 - v4$	$Cat2[0] = 5.5 \times 10^{-6}$
H202ex'[t] = -v1	$H202ex[0] = (1 \dots 250) \times 10^{-6}$
H202'[t] = v2 - v3 - v4 - v5 - v6 - v7	H2O2[0] = 0
cytb5r'[t] = -v8 + v9	$cytb5r[0] = 8.1 \times 10^{-6}$
cytb5ox'[t] = v8 - v9	cytb5ox[0] = 0
metHb'[t] = v6 - v7 - v8 - v13 + v10 + v14	metHb[0] = 0
ferHb'[t] = v7 - v10 - v15 + v16	ferHb[0] = 0
metHbcomplex' $[t] = v13 - v14$	metHbcomplex[0] = 0
ferHbcomplex' $[t] = v15 - v16$	ferHbcomplex[0] = 0
B3'[t] = -v13 + v14 - v15 + v16	$B3[0] = 2.6 \times 10^{-5}$
Hb'[t] = -v6 + v8	Hb[0] = 0.01
Asc'[t] = -v10 + v11	$Asc[0] = 7.5 \times 10^{-5}$
DHA'[t] = v10 - v11	DHA[0] = 0
$GSH'[t] = 2 \times (-v5 - v11 + v12)$	$GSH[0] = 1.5 \times 10^{-3}$
GSSG'[t] = v5 + v11 - v12	GSSG[0] = 0
NADP'[t] = -vG6PDH - v6PGODH + v12	$NADP[0] = 0.2 \times 10^{-6}$
NADPH'[t] = vG6PDH + v6PGODH - v12	$NADPH[0] = 5 \times 10^{-5}$
NAD'[t] = -vGAPDH + vLDH - v9	$NAD[0] = 8.9 \times 10^{-5}$
NADH'[t] = vGAPDH - vLDH + v9	$NADH[0] = 1.5 \times 10^{-7}$



$\mathbf{vG6PDH} = 6.4 \times 10^{-5} \times \frac{NADP[t] \times g6p[t]}{1 + \left(\frac{NADP[t]}{2.67 \times 10^{-6}}\right) \times \left(1 + \frac{g6p[t]}{6.7 \times 10^{-5}}\right) + \frac{NADPH[t]}{2.72 \times 10^{-5}}; \ \mathbf{v6PGLase} = \frac{6.26 \times 10^{-4} \times g16p[t]}{7.99 \times 10^{-5} + g16p[t]};$
$k1 = 240000; k2 = 410; k3 = 2 \times 10^9; k4 = 26000; k5 = 48; k6 = 30; k7 = 630; k8 = 36000; k9 = 800; k10 = 225000; k11 = 300; k12 = 4950000; N1 = k1k3k5k7k9; N2 = k2k4k6k8k10; D1 = k2k9(k4k6 + k5k6 + k5k7); D2 = k1k9(k4k6 + k5k6 + k5k7); D3 = k3k5k7k9; D4 := k2k4k6k8; D5 = k2k10(k4k6 + k5k7); D6 = k1k3(k5k7 + k5k9 + k6k9 + k7k9); D7 = k1k4k6k8; D8 = k3k5k7k10; D9 = k2k10(k2k4 + k2k5 + k2k6 + k4k6); D10 = k1k3(k5 + k6); D11 = k3k8k10(k5 + k6); D3 = k3k5k7k10; D9$
v6PGODH 2.1 × 10 <sup>-6</sup> × (N1 × NADP[t] × go6p[t] − N2 × ru5p[t] × NADPH[t])
$= \frac{1}{D1 + D2 \times \text{NADP}[t] + D3 \times \text{go6p}[t] + D4 \times \text{ru5p}[t] + D5 \times \text{NADPH}[t] \times \text{go6p}[t] + D7 \times \text{NADP}[t] \times \text{ru5p}[t] + D8 \times \text{go6p}[t] \times \text{NADPH}[t] + D9 \times \text{ru5p}[t] \times \text{NADPH}[t] + D1 \times \text{NADPH}[t] \times \text{ru5p}[t] + D1 \times \text{NADPH}[t] \times \text{ru5p}[t] \times ru5p$
$\mathbf{vRSPI} = 1.42 \times 10^{-5} \times \frac{21600}{1 + 33.3 + 14.2} \times \mathbf{vSPI} = 4.22 \times 10^{-6} \times \frac{31490000}{1 + 438 + 305} \times \frac{1490000}{216000} \times \frac{1}{1 + 428 + 305} \times \frac{1490000}{233.3 + 14.2} \times \frac{1}{133.3 + 14.2} \times \frac{1}{233.3 + 1$
21600; k2 = 45.3; k3 = 16.3; k4 = 30000; k5 = 490000; k6 = 60; k7 = k5k7(k2 + k3); T13 = k2k4(k6 + k7); T14 = k6k8(k2 + = k1k4(k6 + k7);
VIA M11 × s7p[t] × ga3p[t] – M12 × e4p[t] × f6p[t]
$= 6.9 \times 10^{-7} \times 111 \times 57p[t] + T12 \times ga3p[t] + T13 \times e4p[t] + T14 \times f6p[t] + T15 \times s7p[t] \times ga3p[t] + T16 \times e4p[t] \times f6p[t] + T17 \times ga3p[t] \times f6p[t] + T12 \times ga3p[t] \times f6p[t] + T17 \times ga3p[t] \times f6p[t] + T17 \times ga3p[t] \times f6p[t] + T17 \times ga3p[t] \times f6p[t] + T10 \times ga3p[t] \times f6p[t] \times f6p[$
$M21 \times x5p[t] \times r5p[t] - M22 \times ga3p[t] \times s7p[t]$
$\times$ 10 <sup>-7</sup> T <sub>2</sub> 1 × x5p[t] + T <sub>2</sub> 2 × r5p[t] + T <sub>2</sub> 3 × ga3p[t] + T <sub>2</sub> 4 × s 216000; k2 = 38; k3 = 34; k4 = 156000; k5 = 2240000; k6 = k5k7(k2 + k3); T <sub>3</sub> 3 = k2k4(k6 + k7); T <sub>3</sub> 4 = 1 = k1k4(k6 + k7);
vTK2 $M31 \times x5p[t] \times e4p[t] - M32 \times ga3p[t] \times f6p[t]$
= 3.3 × 10 × T31 × x5p[t] + T32 × e4p[t] + T33 × ga3p[t] + T34 × f6p[t] + T35 × x5p[t] × e4p[t] + T36 × ga3p[t] × f6p[t] + T37 × e4p[t] × f6p[t] + T38 × x5p[t] × ga3p[t]

SPELL: 1. OK