

# Systemic Effects of Energy Metabolism Regulator Amber-Antitox during Vibration-Induced Dysregulation

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Energy metabolism regulator Amber-antitox had a systemic effect under experimental conditions of vibration-induced dysregulation. Whole-body vibration was accompanied by nonlinear changes in energy metabolism in the heart, liver, kidneys, and lymphocytes and impairment of intrasystemic interorgan relationships between mitochondria. Amber-antitox prevented vibration-induced deenergization of mitochondria and contributed to the preservation of multidimensional relationships of energy metabolism in vital internal organs.

**Key Words:** *energy metabolism; Amber-antitox; vibration-induced dysregulation; factor analysis*

The introduction of medical methods with energy metabolism regulators (EMR) suggests systemic evaluation [3] of their efficiency by means of multivariate analysis [5,9]. Preclinical studies of EMR are performed on models for the *in vitro* response of the organism to exogenous stimulation. The cycle of mitochondrial function in various tissues (rest—activity—rest) is initiated by addition of substrates and ADP. An incomplete rest—activity cycle is observed after treatment with substrates and uncoupler [4,6,11]. The regulatory role of mitochondria in homeostasis is evaluated by recording of various parameters, which hinders the interpretation of data. Summarization of data involves the analysis of interactions between input factors, whose effects are modified by the mitochondrial response in organs and tissues.

Whole-body vibration is a promising experimental model to study the systemic multilevel effect of EMR. Long-term vibration under experimental conditions or exposure to the adverse industrial or environmental factor over many years is followed by

systemic dysregulation of homeostasis, hypoxia, and energy deficiency [2,10]. Systemic approach should be used to develop new schemes for the prevention and therapy of vibration-induced changes (multiple visceropathies and angiopathies [10]) from the positions of bioenergetic pharmacology [12,13].

Here we studied the effect of an EMR Amber-antitox (AA) on vibration-induced dysregulation in energy-producing systems of the myocardium, liver, kidneys, and lymphocytes.

## MATERIALS AND METHODS

Experiments were performed on 60 male Chinchilla rabbits weighing 2.5-3.0 kg and obtained from Podgornyi nursery (Kirov region). The study was conducted according to the rules of the European Convention for the Protection of Vertebrate Animals Used for Experimental and Other Scientific Purposes. Vertical whole-body vibration (amplitude 0.5 mm, frequency 44 Hz) was induced on an UV 70/200 industrial device. The animals were exposed to 60-min vibration in the morning time (9.00-11.00) over 21 and 56 days. Aqueous suspension of AA was administered intragastrically through a tube 60 min before the start of vibration. Single dose of AA (8.4 mg/kg) was estimated from the

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coefficient of bioequivalent doses for various mammalian species and humans taking into account the relations between the weight and relative area of the body surface [1]. Control animals received AA, but were not exposed to vibration.

Blood samples were taken from the marginal vein 60 min before vibration. The state of regulatory systems of energy homeostasis was evaluated from the variability of population and cellular characteristics of lymphocyte SDH activity. The cytochemical method was used to estimate the number and distribution of formazan granules in 50 cells [8]. The energy state of lymphocytes was described by mean enzyme activity and coefficients of variation, asymmetry, and excess [7,8]. Oxidative processes in vibration-sensitive organs (heart, liver, and kidneys) [10] were studied *in vitro* by the method of polarography. The rate of mitochondrial respiration in tissue homogenates was estimated on the model of incomplete rest—activity cycle [4]. The metabolic state of mitochondria under rest conditions was modeled by addition of a FAD-dependent substrate (FAD-DS) succinate (1 mM) or a mixture of NAD-dependent substrates (NAD-DS) glutamate and malate (3 mM). ATPase activity was reproduced by oxidation of FAD-DS and NAD-DS with 2,4-dinitrophenol (20  $\mu$ M). Partial oxidase activity for endogenous substrates was estimated by the sensitivity of endogenous respiration ( $V_E$ ) to inhibitors of NAD-dependent and FAD-dependent respiration (2 mM amytal and 2 mM malonate, respectively). This index was calculated by the equation:  $1 - V_{INH}/V_E$ , where  $V_{INH}$  is endogenous respiration after addition of amytal ( $V_A$ ) or malonate ( $V_M$ ) [6].

The significance of intergroup differences was evaluated by parametric (multiple LSD test) and nonparametric methods (Mann—Whitney  $U$  test) [9]. Interference of the effects of vibration and AA on mitochondria was estimated by factor analysis and principle component analysis [5]. The data were analyzed by means of ANOVA/MANOVA and Principle components & Classification Analysis (Statistica 6.0 software) [11].

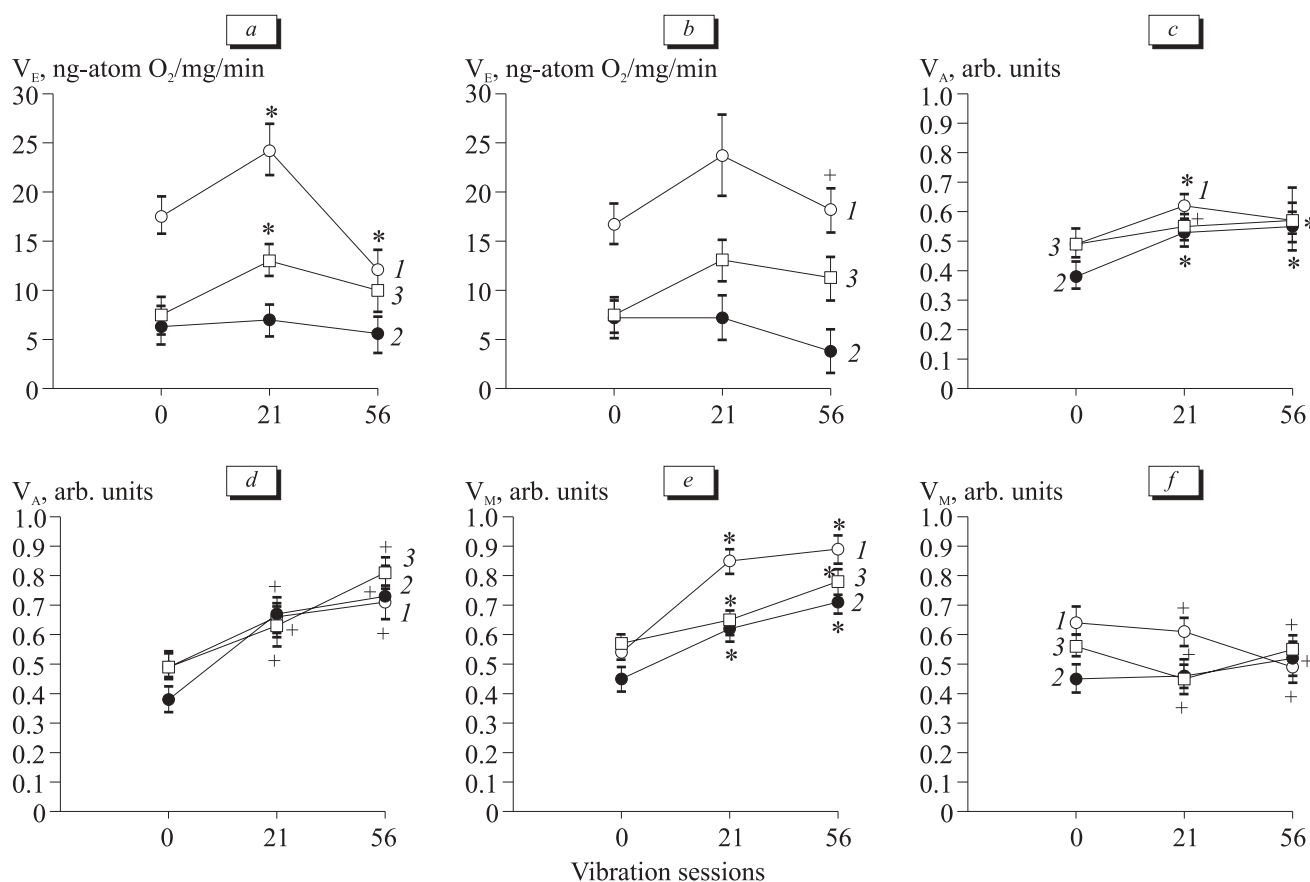
## RESULTS

The study of population and cellular characteristics of SDH activity showed that long-term whole-body vibration impairs the energy state of lymphocytes. It was manifested in several stereotypic signs (Table 1). Mean enzyme activity decreased by 20%. According to the variation coefficient, heterogeneity of cells increased by 51% ( $p < 0.01$ ). Transition of asymmetry to the region of negative values reflected predominance of cells with decreased activity ( $p < 0.001$ ). A negative shift of the excess indicated that lymphocyte pools are unmatched. Low-activity cells dominated under these conditions. Administration of AA before vibration preserved energy status of lymphocytes. SDH activity in treated rabbits was 46% higher than in animals without energy protection ( $p < 0.001$ ). Population variability in treated rabbits decreased by 28% ( $p < 0.001$ ). According to the decrease in asymmetry, pools of cells with high and low SDH activity were more balanced compared to the control. The negative shift of the excess reflected cell transition from the pool with normal activity to the pool with high activity. Evaluation of the energy state of lymphocytes showed that AA has a consistent and complete effect. Clinical and physiological observations showed that these cytochemical parameters of lymphocyte SDH activity reflect a positive effect of EMR [7,8]. Vibration and energy protection with AA produced phasic changes in mitochondria of the heart, liver, and kidneys, which nonlinearly depended on the duration of exposure. We observed asynchronous changes in several parameters, while other parameters underwent stereotypic changes. Variations in  $V_E$  differed in the range and directionality (Fig 1, a).  $V_E$  decreased in myocardial mitochondria, but returned to the control level in mitochondria of the liver and kidneys. FAD- and NAD-dependent oxidases had similar activity in utilization of endogenous substrates. The slight increase in  $V_A$  reflected insignificant acceleration of NAD-dependent reactions in the respiratory chain (Fig. 1, c).  $V_M$

**TABLE 1.** Effect of AA and Long-Term Whole-Body Vibration on Energy State of Rabbit Lymphocytes

Parameter	Control (n=12)	Whole-body vibration (n=10)	Whole-body vibration+AA (n=10)
Mean enzyme activity	10.5 (9.8-11.1)	8.4 (7.5-9.3)*	12.3 (11.1-13.7)*
Variation coefficient	54 (51-58)	82 (75-89)*	59 (55-63)*
Distribution asymmetry coefficient	0.60 (0.56-0.75)	-0.67 (-0.86 - -0.50)*	0.23 (0.08-0.39)*
Excess coefficient	-0.41(-0.67-0.17)	-0.56 (-0.87 - -0.27)	-0.65 (-0.94 - -0.38)

**Note.** Mean values; 95% confidence intervals are shown in parentheses.  $p < 0.05$ : \*compared to the control; \*compared to animals of the vibration group.



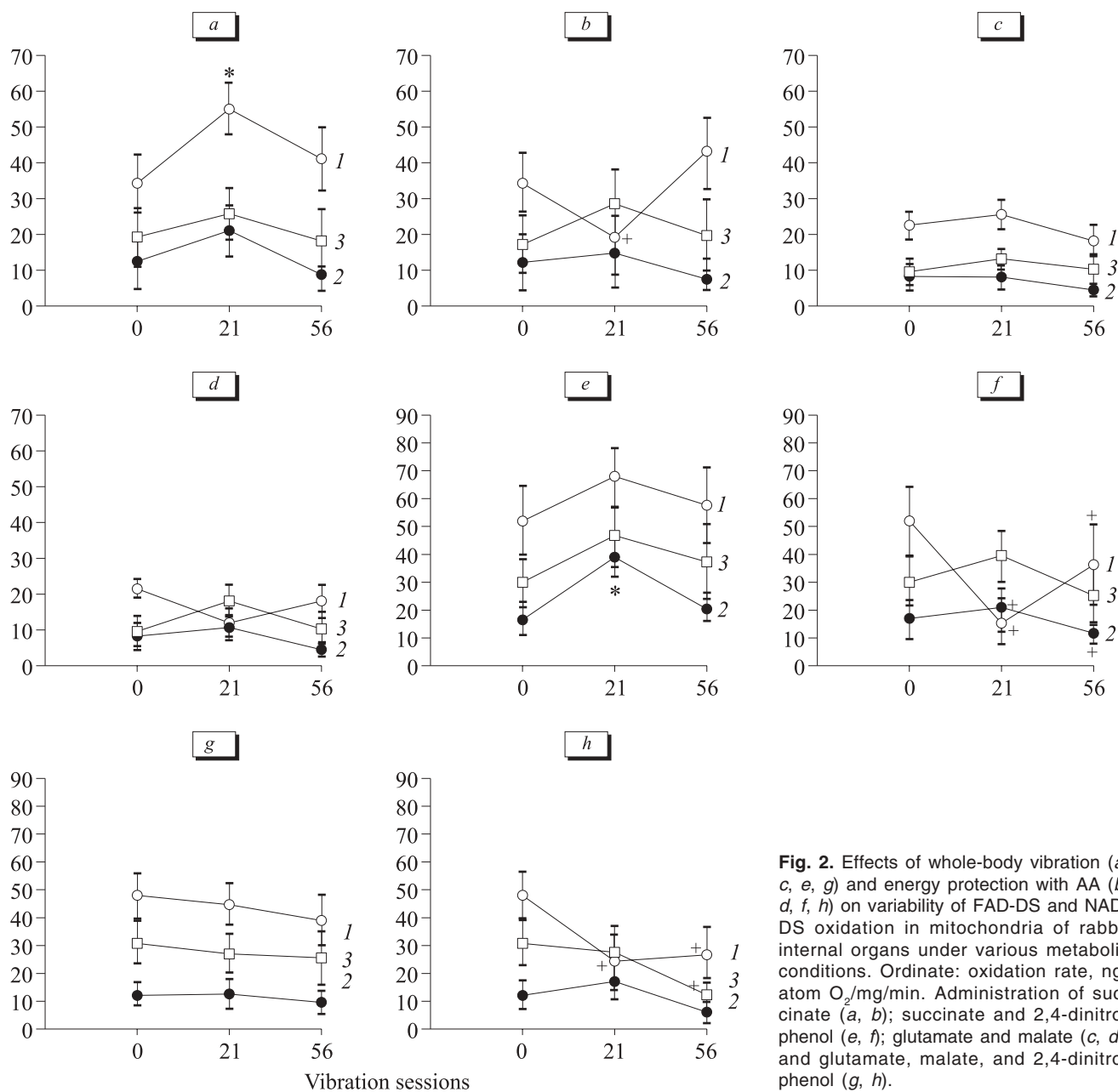
**Fig. 1.** Effects of whole-body vibration (a, c, e) and AA (b, d, f) on oxidation of endogenous substrates in rabbit myocardium (1), liver (2), and kidneys (3) and sensitivity of  $V_E$  to inhibitors of NAD- (b) and FAD-dependent fractions of the mitochondrial respiratory chain (c). Here and in Fig. 2:  $p < 0.05$ : \*compared to pre-vibration parameters; \*compared to animals with no pharmacological protection.

underwent a monotonous increase (Fig. 1, e) and prevailed over  $V_A$ . Therefore, dysregulation is accompanied by synchronous mobilization of the succinate-dependent bioenergetic system in internal organs.

After combined exposure to vibration and AA, myocardial  $V_E$  practically did not differ from the control (Fig. 1, b). However, this parameter in treated rabbits was higher compared animals of the vibration group.  $V_E$  remained nearly unchanged in the kidneys, but decreased in the liver. Inhibitory analysis showed that variations in the contribution of  $V_A$  to the monotonous increase of  $V_E$  in all organs (Fig. 1, d) are in inverse proportion to small changes in  $V_M$  (Fig. 1, f). These data indicate that AA decreases tissue demands for O<sub>2</sub> and improves mitochondrial function. AA prevents hyperactivation of the FAD-dependent  $V_E$  fraction, but increases the contribution of NAD-DS. Oxidation of FAD-DS and NAD-DS in the rest—activity cycle nonlinearly depended on the duration of exposure to input factors (Fig. 2). Oxidation of exogenous succinate corresponded to variations in  $V_E$  (Fig. 2, a, e). However, oxidation of NAD-DS decreased in the myocardium or remained practically unchanged

in mitochondria of the liver and kidneys (Fig. 2, c, g). The regulatory effect of AA was most significant in myocardial mitochondria. The myocardium is one of the organs with high-intensity production and consumption of energy (Fig. 2, b, d, f, h). The extremely high rate of respiration decreased, while the respiratory rate corresponding to the range of control values underwent only small changes. Similar results were obtained in experiments with the liver and kidneys. However, the rate of substrate oxidation in these organs was lower. Plasticity of the reactions in rapid metabolic cluster [6,11-13] was higher compared to the NAD-dependent system of energy production (Figs. 1 and 2). The system for production and utilization of succinate probably plays a key role in accumulation of the effects of adverse factor and formation of the response to AA. It is necessary to take into account complex interorgan relationships of mitochondria. They vary under conditions of dysregulation and pharmacological correction, but are unobvious during the data comparison on a plane.

Factor analysis allowed us to decrease the size of information on the interaction of vibration, AA,



**Fig. 2.** Effects of whole-body vibration (a, c, e, g) and energy protection with AA (b, d, f, h) on variability of FAD-DS and NAD-DS oxidation in mitochondria of rabbit internal organs under various metabolic conditions. Ordinate: oxidation rate, ng-atom O<sub>2</sub>/mg/min. Administration of succinate (a, b); succinate and 2,4-dinitrophenol (e, f); glutamate and malate (c, d); and glutamate, malate, and 2,4-dinitrophenol (g, h).

and mitochondria in various organs to 2 independent variables (principle components, Table 2). Vibration and energy production modified the factor structure of the correlation matrix, which reflected the comparability of the strength of their interaction. In animals of the control and vibration groups, the degree of variance absorption by principle components ( $F_1$ ) was 2.2-2.5 times higher compared to auxiliary components ( $F_2$ ).  $F_1$  and  $F_2$  were balanced after AA administration. The regulatory effect of EMR counteracted the influence of a dysregulating factor, which contributed to the maintenance of interorgan relationships in mitochondria at the control level. Vibration decreased the ratio of the

total variance absorbed by both components (decrease in regulation). The ratio of principle components to the total variance remained unchanged after AA administration (preservation of regulation).

In control animals, strong negative multiple correlations were revealed between kinetic parameters and  $F_1$  (factor loadings  $>0.8$ ). The contribution of partial reactions was low. However,  $F_1$  mainly depended on  $V_M$ .  $F_2$  had a moderate effect only on  $V_E$ . The remaining interrelations were insignificant. The factor structure of the matrix significantly changed during dysregulation. Multiple correlations between parameters produced a strong effect on  $F_1$  (factor loadings  $\geq 0.8$ ), but had a posi-

tive sign. Therefore, the regulatory interactions were diminished under these conditions. As distinct from  $V_A$ ,  $V_M$  had little effect on  $F_1$ . The value of  $F_2$  absorbing a small part of the total variance distributed the loading on parameters for metabolic activity of mitochondria during administration of succinate, succinate+2,4-dinitrophenol, and glutamate+malate (Table 2). After AA administration, most correlations of test parameters with  $F_1$  had a negative sign. However, significant changes were revealed in absolute parameters of factor weights for individual parameters and their relation to  $F_2$ . Factor loadings for  $V_E$ ,  $V_M$ , and administration of glutamate+malate+2,4-dinitrophenol and signs of their influence on  $F_1$  were preserved. Loadings for administration of succinate, succinate+2,4-dinitrophenol, and glutamate+malate on  $F_2$  were redistributed and had a negative sign. These changes show that the system of rapid metabolic cluster is involved in reorganization of tissue bioenergetics. Intersystem regulatory interactions were preserved and partially transformed due to energy production during vibration and accumulation of the effects at the level of mitochondria. Similar tendencies were found in generalized covariations of grouping signs with  $F_1$  and  $F_2$ . The influence of vibration and EMR on intersystem interactions of mitochondria (objectively present and implicit regularity) was manifested during formalized presentation of test objects in  $F_1/F_2$  coordinates (Fig. 3). In control animals, the points

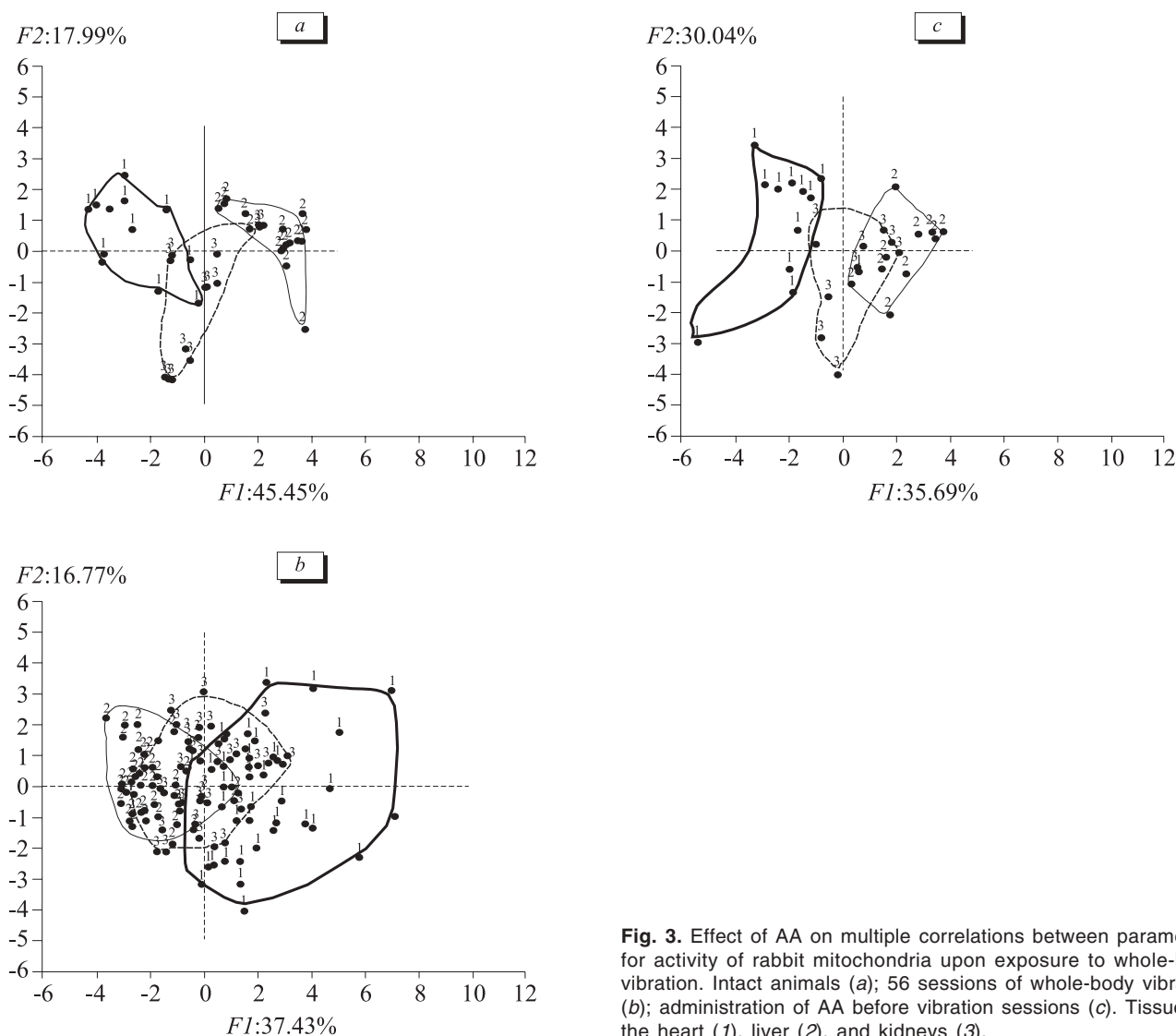
corresponding to mitochondria of internal organs formed the separate and compact clusters (Fig. 3, *a*). During vibration these clusters became diffuse, superposed, and disoriented and changed the position (Fig. 3, *b*). Administration of AA decreased the degree of dysregulation changes and contributed to the preservation of compactness, separateness, and orientation of point clusters, which did not differ from the control (Fig. 3, *c*).

The strength and directionality of correlations between various parameters reflect the rate and effectiveness of system transition from one state to another state [3]. Our results indicate that EMR not only has a regulatory effect on activity of rapid metabolic cluster in mitochondria of each organ (increase or decrease) [6,11-13], but also optimizes interorgan relationships between parameters of mitochondrial bioenergetics that are impaired under the influence of an adverse factor. This phenomenon can be considered as a sign for systemic integral function of mitochondria [13], which is activated by AA. AA is the exogenous factor that preserves partial energy systems in tissues and organs and maintains stable energy homeostasis during adverse exposure to vibration. Tissue mitochondria summarize the damaging effects, which contributes to the appearance of a material substrate for adaptation stages associated with various levels of energy deficiency [11,12]. The protective effects of EMR are also summarized at the level of tissue mitochondria.

**TABLE 2.** Effect of Long-term Whole-Body Vibration and AA on Factor Structure of Intertissue Interactions of Rabbit Mitochondria

Parameter	Control		Whole-body vibration		Whole-body vibration+AA	
	$F_1$	$F_2$	$F_1$	$F_2$	$F_1$	$F_2$
Ratio of fraction absorbed by each component, %	45.5	17.9	37.4	16.8	35.7	30.0
Dispersion of the amount of basic data, %	63.4	54.2	65.7			
Factor loadings of parameters for mitochondrial function, arb. U						
$V_E$	-0.84	0.51	0.82	0.26	-0.90	0.26
$V_A$	-0.40	-0.01	0.10	0.00	0.19	0.00
$V_M$	-0.55	-0.13	0.32	0.31	-0.52	0.31
after addition of						
succinate	-0.91	0.25	0.79	-0.53	-0.69	-0.53
succinate and 2,4-dinitrophenol	-0.96	0.06	0.89	-0.84	-0.68	-0.84
glutamate and malate	-0.87	0.38	0.87	-0.66	-0.49	-0.66
glutamate, malate, and 2,4-dinitrophenol	-0.95	-0.13	0.91	-0.37	-0.89	-0.37
Covariations of components and grouping signs						
tissue	-0.99	0.12	0.45	0.05	-0.25	-1.85
vibration	0.0	0.0	0.51	-0.21	0.42	1.12





**Fig. 3.** Effect of AA on multiple correlations between parameters for activity of rabbit mitochondria upon exposure to whole-body vibration. Intact animals (a); 56 sessions of whole-body vibration (b); administration of AA before vibration sessions (c). Tissues of the heart (1), liver (2), and kidneys (3).

dria. EMR contribute to the preservation and stabilization of system interactions between mitochondria in cells and organs upon exposure to an adverse factor. They increase resistance of the organism and have similar pharmacodynamic characteristics under various pathological conditions.

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