Influence of Low Temperature on the Evolution of Amino Acid Pools Adaptive Modifications in Poikilothermal Animals

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Abstract

Interpretation of phylogenesis in the light of comparative biochemistry is currently still limited. Adaptive changes in the concentration of the FAA in response to the decrease in temperature are one of the essential features of poikilothermic animals living in areas with large temperature difference. It is purpose of this review to reveal the hierarchy of certain biochemical sign, namely the FAA that underlie of differences in invertebrate and vertebrate poikilotherms and in the process of evolution begin to discover themselves only at low temperatures. The review summarizes literature and author information about evolutionary development of adaptive FAA responses of poikilothermic animals, at different levels of phylogenesis, as a result of the seasonal drop in temperature to negative or zero values. It is concluded that the non-specific accumulation of proteinogenic amino acids, characteristic for many invertebrates, is replaced in the vertebrates animals by the accumulation of new participants in the mechanisms of the low temperatures adaptation, which is a nonproteinogenic sulphoamino acid, taurine, and in the brain it is PEA and phosphoamine acid, PS.

Keywords Evolution, Adaptation, Low Temperatures, Poikilothermal Animals, Brain, Phosphoethanolamine, Phosphoserine, Taurine.

1. Introduction

It is presumed that global climate change is one of the main driving forces behind the evolution of living organisms [1]. Mutations of genes seem to occur more frequently during periods of cooling, giving a powerful impetus to the evolution of organisms and resulting intensive formation of new types [2, 3]. Traces of a progressive evolutionary change, their stages and "witty" finds can be seen when studying the effect of low temperatures on the biochemical mechanisms in different types of ectothermic animals. The expression of many genes that encode proteins involved in signal transductions and chaperones may be changed when exposed to cold shock or cold acclimation [4–6]. The synthesis of cold-induced proteins and modification of existing proteins, conversion of lipid composition of cell membranes, activation of isoenzymes, change of ion, oxygen and metabolite transport are required for adaptation of poikilothermic animals to rapid decreases in environmental temperatures [7, 8]. In addition, there are the syntheses or accumulation of substances that perform the cryoprotective function (alcohols, sugar, antifreeze proteins, ice-nucleating agent) [9–12]. The FAA and PEA, that is an ethanolamine derivative, also contribute their mite in biochemical strategy of low-temperature adaptation.

2. Seasonal Changes of the FAA Pool in the Invertebrates

The content of certain FAA is notably increased in the blood, hemolymph and in bodies of poikilothermic animals, both invertebrates and vertebrates when exposed to low temperatures as a result of the change of seasons or the sudden cold stress. The FAA which perform functions of membrane protectant, osmolytes or antioxidants [13–15], differ depending on the stage of the macroevolution or habitats [9–11]. Accumulation of the FAA in invertebrates (except for intertidal shellfish [16, 17]) is the less than in vertebrates and not accumulated FAA are specific for low-temperature adaptation [18–24]. In addition, a common classifies sign of "protective" the FAA for most invertebrate is that all they are proteinogenic ("protein-building") amino acids, that is, they combine into peptide chains to form the building blocks of proteins. Most of the accumulated amino acids are glucoplastic (e.g., alanine, glutamate, glycine, proline, serine, valine, histidine) [18–24]. Many of them are active participants in the metabolism.

Taking into account the huge variety of the animal world, you should not expect that all types and classes of
poikilothermic animals must be examined in the context of the response FAA to cold temperatures impact, but even scattered evidence of comparative biochemistry can give an idea about the typical features of pool FAA adaptive modifications of the poikilotherms. Now, thanks to sufficient works on changes of amino acid pools of cryotolerant and cryoresistant invertebrates when exposed to cold, we know about representative features of their adaptive FAA responses [16–24]. Thanks to the work performed on the intertidal molluscs and mussels, surviving temperatures down to -10°/-15°C, it is known about the specific FAA pattern of these species of marine invertebrates [16, 17]. For example, the pool of taurine in leg muscle of the Littorinalitioria is 77% of the total FAA pool – both at freezing to -4°C and in control at 5°C [17]. In gill, mantle, and muscle of the marine mussel Geukensiademissus, taurine together with glycine amount to 80–85% of the total pool at the 2 or 12h freezing to -6°C [16].

Ecological factors such as average annual temperature, water salinity significantly affect the biochemical features even related of poikilothermic animals. It is not surprising that the biochemical properties of adaptation in freshwater molluscs, wintering at a temperature of about 0°C, different from the above-mentioned properties for intertidal molluscs [22, 23]. Unfortunately, the study of freshwater hydrobionts in this direction is virtually non-existent, and the author can mention only data on fresh water mollusc Lymnaea Stagnalis [22, 23]. According to our data, the great pond snail L. stagnalis, hibernating in winter deep in pond clay until spring, contains taurine in summer and before winter hibernation only in trace quantities (Table 1, Fig. 1) [23]. At a decrease of the temperature to 0°C at the autumn period, concentration of free alanine in body fluids of L. stagnal, is increased in 2,7 times (Table 1). Pool of alanine as a percentage of the total FAA pool, which grows also, increased in 1,4 times and amounts to 25.8% (Fig. 1). An increase of histidine, glutamate, glycine, and serine concentrations was less pronounced.

Increasing the concentration of free alanine during adaptation of L. stagnalis, correlated with increasing levels of serine (Table 1), is typical as well for insects, as for example the southwestern corn borer and the European corn borer in relation to their diapause [9, 18]. Some types of insects during the period of seasonal adaptations increase the level of free alanine and proline [9]. Increased alanine, histidine, glycine and lysine pools, which is typical of freshwater mollusk L. stagnalis, happens in the hemolymph of oak silkworm Antheraea pernyi along with increased proline, arginine and lysine [19]. Increase of alanine, lysine, glycine and phenylalanine is also shown for larvae of the wax moth Galleria mellonella [20]. Alanine, glutamate, valine and lysine pools among some species of beetles are also increased as a result of adaptation to the low temperature [21].

Table 1. Amino acid pool changes in tissue fluids of mollusk L. stagnalis before the winter hibernation (µmol/100 ml-1) [22].

| Aminoacid       | June         | October       |
|-----------------|--------------|---------------|
| Taurine         | tr           | tr            |
| Threonine       | 25.2 ± 1.7   | 20.4 ± 0.4    |
| Serine          | 21.9 ± 1.9   | 37.0 ± 2.3    |
| Glutamic        | 55.1 ± 2.1   | 69.3 ± 4.4    |
| Proline         | 24.0 ± 7.7   | –             |
| Glycine         | 38.5 ± 2.1   | 59.5 ± 2.9    |
| Alanine         | 47.1 ± 2.1   | 128.4 ± 7.5   |
| Cysteine/Cystine| –            | 29.7 ± 15.3   |
| Valine          | 11.5 ± 1.2   | 22.6 ± 0.9    |
| Methionine      | 3.0 ± 0.8    | 7.0 ± 1.3     |
| Isoleuzine      | 10.5 ± 1.2   | 16.2 ± 0.7    |
| Leuzine         | 10.2 ± 1.4   | 20.5 ± 0.9    |
| Tyrosine        | tr           | 10.6 ± 1.4    |
| Phenylalanine   | tr           | 14.7 ± 1.5    |
| Histidine       | 8.1 ± 1.2    | 27.1 ± 1.7    |
| Lysine          | tr           | 20.5 ± 1.2    |
| Total           | 264.4 ± 21.2 | 497.0 ± 42.1  |

Values are means ±σ (n = 3); tr – substance was discovered in trace quantities. The animals were captured in late October.

On the basis of such fragmentary data, it would be reasonable to suppose that in the world of invertebrates at some stages of phylogenesis there is no specificity of adaptive responses of the FAA at low temperatures, and in the process of compensatory increasing of total FAA pool involved many proteinogenic amino acids, which are mostly the same for different types of organisms.

The appearance at any stage of phylogensis non-proteinogenic amino acids, accumulated during the period of seasonal temperatures fall, their specific reaction to cold are a marker of the transition to a higher level of evolution. For example, our data show that ornithine pool in hemolymph of freshwater higher crustaceans Gammarus lacustris, living in an area with large difference of summer and winter temperatures, in autumn is 16.5% of the total FAA pool, taking second after alanine, which is 23.3% (table 2, Fig 1) [24]. However, this non-proteinogenic amino acid is not detected in the freshwater amphipod Gammarus pseudolimnaeus, living in a warmer climate zone [25].
phylogenesis, lacking in the literature, however the accumulation of taurine on earlier, than teleosts, stages of membrane stabilizator, cellular protectant. Data on associated not with metabolism, and serves as an antioxidant, involved in it. Taurine in the circumstances, is probably torpor at zero temperature and proteinogenic FAA are indicates that metabolism is carried out even in a state of accounting for 11,9% of the total FAA pool [26]. This fact after the release of the winter torpor, turns sharply, during this period virtually unchanged, but in the spring, three-month hibernation of P. glehni in ice, level of taurine proteinogenic amino acid that alter their pools during the total pool (table 4, Fig 2) [26].In contrast to the December to 7,7 times in comparison with July and is 45% of the level of taurine in the blood rises in 4,3 times, making up 52,6% of the total FAA pool (table 3, Fig 2) [26]. The level of taurine in the muscles increases over the summer period to 4,40 ± 0,08, making up 52,6% of the total FAA pool (table 3, Fig 2) [26]. The most impressive data are obtained for Nonproteinogenic Amino Acid in the Vertebrates Animals at Low Temperatures

The most impressive data are obtained for the phenomenon of adaptive accumulation of taurine (2-amino-ethanesulfonic acid) is found first in our works in examining the impact of seasonal temperature drops or cold shock effect on the pattern of FAA in the eurythermal freshwater fish P. glehni [26–27].

In winter, before beginning hibernation, the contents of taurine in the muscles increases over the summer period to 4,3 times, making up 52,6% of the total FAA pool (table 3, Fig 2) [26]. The level of taurine in the blood rises in December to 7,7 times in comparison with July and is 45% of the total pool (table 4, Fig 2) [26]. In contrast to the proteinogenic amino acid that alter their pools during the three-month hibernation of P. glehni in ice, level of taurine during this period virtually unchanged, but in the spring, after the release of the winter torpor, turns sharply, accounting for 11,9% of the total FAA pool [26]. This fact indicates that metabolism is carried out even in a state of torpor at zero temperature and proteinogenic FAA are involved in it. Taurine in the circumstances, is probably associated not with metabolism, and serves as an antioxidant, membrane stabilizator, cellular protectant. Data on accumulation of taurine on earlier, than teleosts, stages of phylogeny, lacking in the literature, however the

exceptions are the above-mentioned intertidal clams and mussels with taurine at negative temperatures is very high [16,17]. High levels of taurine in the invertebrate can result in misleading in determining phylogenetic relationships, because this fact is probably an example of biochemical convergence. Taurine in hemolymph of freshwater molluscs L. stagnalis found at any time of the year only in trace amounts, and in insects taurine is the minimum, and in amphipods G. lacustris is 2% of the total FAA pool in the fall [22–24].

Table 2. Amino acid pools of amphipod G. lacustris run-up to the winter hibernation [24].

| Aminoacid   | μmol/ml of homogenate | % of total pool |
|-------------|-----------------------|-----------------|
| Taurine     | 2,08 ± 0,15           | 2,2             |
| Aspartic    | 1,12 ± 0,12           | 1,2             |
| Threonine   | 3,35 ± 0,68           | 3,5             |
| Serine      | 5,28 ± 0,86           | 5,4             |
| Glutamic    | 5,76 ± 0,59           | 5,9             |
| Proline     | 4,5 ± 0,31            | 4,6             |
| Glycine     | 3,9 ± 0,42            | 4,0             |
| Alanine     | 22,6 ± 1,39           | 23,3            |
| Valine      | 5,03 ± 0,23           | 5,2             |
| Methionine  | 2,04 ± 0,22           | 2,1             |
| Isoleuzine  | 3,82 ± 0,21           | 3,9             |
| Leuzine     | 6,32 ± 0,44           | 6,5             |
| Tyrosine    | 2,01 ± 0,11           | 2,1             |
| Phenylalanine Histidine | 1,94 ± 0,23 | 2,0          |
| Lysine      | 8,39 ± 0,55           | 8,7             |
| Ornithine   | 17,0 ± 1,4            | 17,6            |
| Arginine    | tr                    | tr              |
| Total       | 96,8 ± 8,2            | 100             |

Values are means ±σ(n = 3). The collection of animals was conducted in mid-September.

3. The Uprise of Nonproteinogenic Amino Acid in the Vertebrates Animals at Low Temperatures

Table 3. Influence of seasonal temperature changes on amino acids pools in the blood plasma of freshwater fish P.glehni before beginning hibernation (μmol/ml) [26].

| Amino acid | July     | December |
|------------|----------|----------|
| PS         | –        | –        |
| Taurine    | 0,24 ± 0,03 | 1,85 ± 0,10 |
| PEA        | –        | –        |
| Aspartic   | 0,08 ± 0,01 | 0,06 ± 0,01 |
| Threonine  | 0,25 ± 0,06 | 0,17 ± 0,01 |
| Serine     | 0,36 ± 0,03 | 0,28 ± 0,01 |
| Glutamic   | 0,34 ± 0,04 | 0,27 ± 0,02 |
| Proline    | 0,33 ± 0,04 | –        |
| Glycine    | 0,43 ± 0,04 | 0,35 ± 0,01 |
| Alanine    | 0,50 ± 0,02 | 0,52 ± 0,02 |
| Valine     | 0,21 ± 0,01 | 0,09 ± 0,01 |
| Cystathionine | –        | 0,14 ± 0,03 |
| Methionine | 0,06 ± 0,01 | 0,14 ± 0,01 |
| Isoleuzine | 0,15 ± 0,01 | –        |
| Leuzine    | 0,25 ± 0,01 | –        |
| GABA       | –        | 0,36 ± 0,04 |
| Tyrosine   | 0,17 ± 0,01 | –        |
| Phenylalanine | 0,39 ± 0,04 | –        |
| Histidine  | 0,24 ± 0,53 | 0,07 ± 0,01 |
| Lysine     | 0,19 ± 0,01 | 0,10 ± 0,01 |
| Arginine   | 0,20 ± 0,01 | –        |
| Total      | 4,40 ± 0,08 | 4,11 ± 0,16 |

Values are means ±σ(n = 3); P < 0.05. Catching the animals on December 26.

Figure 2. Increase the pool of taurine in muscles and in blood plasma of eurythermal freshwater fish P.glehni before beginning hibernation. S – summer (July), W – winter (December).
Having successfully carried out its tasks with a low-temperature adaptation of teleosts, taurine moves to a higher level of evolution, including reptiles and mammals [29–34]. Data on low temperature modifications the FAA pools for amphibians could not be found in the literature, however, the work carried out on reptilian, reported high content of taurine in plasma and organs of various species of snakes and turtles at moderate temperatures and about a significant increase in the pool of taurine in winter [29, 30]. The total FAA pool in turtle blood C. pictamarginata after a three-day exposition at temperatures from –6 to –8°C increases at 2,25 times, and 52% of increase is due to the taurine [29]. Increased more than 2 times of taurine concentration in freezing up to –2,5° Coccurs in the muscles of the garter snake T. sirtalisparietalis [30]. Elevation of taurine level in 1,5–3°C of taurine concentration in freezing up to –2,5° Coccurs in the muscles of the garter snake T. sirtalisparietalis [30]. Elevation of taurine level in 1,5–3°C of taurine concentration in freezing up to –2,5°

In summer [34]. And now, taurine is a ubiquitous amino acids during winter hibernation and artificial hypothermia in the human body of a wide spectrum of effects. These include stabilization of plasma membrane, modulation of intracellular calcium homeostasis, prevention of cell apoptosis, osmoregulation, reactive-radical scavenging, oxidative stress, neuromodulation and neurotransmission, protein phosphorylation modulation etc. [35–38].

4. PEA and Taurine in the Brain of Vertebrates at Low Temperatures

Unexpected facts were found by us when examining low-temperature responses of the FAA in the brain of eurythermal fresh water fish P. glehni[27, 28]. It turned out that seasonal decrease in temperature, as well as the effect of cold shock, stimulates the emergence in the fish brain of two new members, PEA and PS, which are linked to the structure of membrane phospholipids [39–41]; PEA is a derivative of ethanolamine, and PS is nonproteinogenic amino acid. The appearance of PEA and PS is specific to the brain of fish because in plasma and in the muscle these amino acids are not detected either in summer or in winter [27]. PEA in the brain of P. glehni accumulates in much greater numbers than PS. Pool of PEA, consisting in the middle of summer only 0,3% of the total FAA pool, at the beginning of winter rising to 33,6% (table 5, Fig 3). PS, which not found in the summer in a free form, at the beginning of winter is 3,75% of the total pool (table 5) [28]. It is also interesting that accumulation of the PEA has a negative correlation with the content of taurine [27, 28]. In contrast to the usual increase of taurine level in the blood and organs at low temperatures, in the brain of fishtaurine is declining against the background of increasing levels of PEA and PS (table 5, Fig 3).

Table 4. Seasonal changes of FAA pool in the muscle of freshwater fish P. glehni (µmol/g wet weight) [26].

| Aminoacid | July       | December   |
|-----------|------------|------------|
| PS        | 3.26 ± 0.40| 14.06 ± 0.99* |
| Taurine   | 0.07 ± 0.01| –          |
| Aspartic  | 1.59 ± 0.13| 0.96 ± 0.07** |
| Serine    | 0.9 ± 0.1  | 0.46 ± 0.04** |
| Glutamic  | 0.97 ± 0.08| 1.50 ± 0.11** |
| Proline   | 1.24 ± 0.12| –          |
| Glycine   | 2.0 ± 0.15 | 1.47 ± 0.06** |
| Alanine   | 1.5 ± 0.13 | 2.92 ± 0.33** |
| Valine    | 0.53 ± 0.05| 0.29 ± 0.03** |
| Cystathionine | 0.98 ± 0.09  |
| Methionine| 0.04 ± 0.01| 0.18 ± 0.01* |
| Isoleuzine| 0.15 ± 0.01| –          |
| Leuzine   | 0.31 ± 0.02| tr         |
| GABA      | –          | 2.40 ± 0.13 |
| Tyrosine  | 1.08 ± 0.09| 0.94 ± 0.11 |
| Phenylalanine | 1.14 ± 0.06 | 0.19 ± 0.01* |
| Histidine | 2.06 ± 0.12| 0.43 ± 0.04* |
| Lysine    | 0.87 ± 0.09| 0.38 ± 0.04** |
| Arginine  | 0.49 ± 0.04| –          |
| Total     | 17.88 ± 1.02| 26.71 ± 0.90** |

Values are means ± s.e.m (n = 4); * – P < 0.001; ** – P < 0.05. Catching the animals on December 26.

Table 5. Seasonal changes of FAA and PEA pool in the brain of freshwater fish P. glehni (µmol/g wet weight) [28].

| Aminoacid | June       | December   |
|-----------|------------|------------|
| PS        | 0          | 410 ± 43   |
| Taurine   | 3897 ± 166 | 905 ± 66*  |
| PEA       | 39 ± 15    | 3663 ± 137*|
| Aspartic  | 82 ± 7     | 0          |
| Threonine | 0          | 14± 4      |
| Serine    | 0          | 614 ± 37*  |
| Glutamic  | 2079 ± 140 | 344 ± 22*  |
| Glycine   | 947 ± 53   | 669 ± 40** |
| Alanine   | 896 ± 39   | 717 ± 66** |
| Valine    | 203 ± 17   | 76 ± 13*   |
| Isoleuzine| 25 ± 3,5   | 0          |
| Leuzine   | 33 ± 8     | 0          |
| GABA      | 1105 ± 84  | 1137 ± 70  |
| Tyrosine  | 77 ± 9     | 0          |
| Histidine | 3290 ± 283 | 2167 ± 94* |
| Lysine    | 602 ± 51   | 214 ± 20** |
| Total     | 13275 ± 481| 10931 ± 198** |

Values are means ± s.e.m (n = 4); * – P < 0.001; ** – P < 0.05.

Figure 3. Influence of seasonal temperature changes on PEA and taurine pools in the brain of eurythermal freshwater fish P. glehni. S – summer (July), W – winter (December).
The absence of serine in the summer, but high level in winter and substantial increase at impact of cold shock are also characteristic of low temperature adaptation in the brain of *P. glehni* [27, 28]. Metabolism of serine is associated with PEA via ethanolamine (serine $\rightarrow$ ethanolamine $\rightarrow$ PEA). This hydroxyamino acid is one of the key figures of the phospholipid metabolism in brain, turning into phosphatidylserine by the reaction with phosphatidylcholine or PtdE, as well as participating in the synthesis of sphingolipid molecules [39]. Increasing the concentration of PEA and serine are likely reflects changes in the phospholipid component of cell membranes at low temperatures. It is appropriate to add that negative correlation of serine and taurine pools in the brain slices of chicken, where it was shown that perfusion of L-serine reduced the concentration of taurine [42].

Despite wide expansion of PtdE in the animal world, literary data on the presence in the brain poikilotherms of free PEA are missing, although there are sometimes messages about its detection in whole body of invertebrates [21, 43]. Discrete, dependent on temperature, the presence of PEA and PS in the brain of teleosts presumes the start of such change in evolution of the animal world, in which a special role will be given to this substrate, which related, probably, to adaptive changes of membrane sphyngomyelin or phospholipid metabolism [44–46]. For example, accumulation of PEA, which is an intermediate substrate in the synthesis of PtdE and phosphatidylcholine[40, 41, 45, 46], could be explained by removal of PEA from PtdE by phospholipase C.

Indeed, an eukaryotic cell has phospholipase C that cleavage link between glycerine and the phosphate group in most types of phospholipids, however phospholipase C of an eukaryotes does not disconnect this link in PtdE[47, 48]. The enzyme, cleaving this link precisely in PtdE, was found only in pathogenic bacteria *Clostridium perfringens*, synthesizing «aggressive enzymes» such as proteinase, lecithinase, collagenase, hyaluronidase and α-toxin, which is a phospholipase C [49]. In eukaryotic cells such enzyme is still not detected for PtdE, so the reliability of the first assumptions is unlikely. It can be presumed that the accumulation of the PEA is due to a breach of PtdE synthesis at the stage of PEA and CTP interaction with the participation of CTP: phosphoethanolamine cytidylyltransferase. The content of PtdE in the membranes of the brain in this case should be decreased, and the PEA should be increased. However, according to Chang and Roots [50, 51], amounts of PtdE in microsomes and inner mitochondrial membranes from brain of goldfish *Carassiusauratus* increases significantly at low temperatures (5°С). The content of PtdE increases in the winter not only in the brain (41%), but also in edible flesh of rainbow trout and perhaps in other organs[52].

Mention should be made of other interesting data. So, it is shown that the increase PtdE synthesis in isolated rat hepatocytes was accompanied by a considerable increase in the pool size of PEA, whereas the amount of cytidine diphosphate ethanolamine remained constant [53, 54]. Therefore, it can be assumed that the expression of a gene occurs in the brain of eurythermal fishes at low temperatures, inducing the synthesis of isoenzyme of ethanolamine phosphokinase, and ATP phosphorylates ethanolamine (forming PEA) in the presence of this isoenzyme with greater speed than at normal temperatures. It is possible PEA in an excessive concentration is the activator of the PtdE synthesis at low temperatures.

Accumulation of free PEA in brain of teleosts when exposed to low temperatures presumes its specific role in evolution. It is unknown about existence of PEA in the brain jawless and cartilaginous fishes standing at the lower stages of phylogenesis. High level PEA in the brain of mammals [55] and its possible start in the brain of fish suggest that PEA gradually increases its presence in the brain in the process of evolution.

5. The Algorithm of Evolution

Further, taking into account the algorithm of evolution, it can be assumed that PEA and PS are absent in ganglia of invertebrate even at low temperatures. The results of our new work performed in autumn on the brain of previously mentioned the freshwater mollusk *L. stagnalis*, confirmed this assumption, but they added again the fact of participation in adaptation to cold the other nonproteinogenic amino acid, namely β-alanine (unpublished data). It can be presumed that discussed PEA and PS should be kept (and possibly even the enhance) its presence in a brain of superior vertebrates (amphibians, reptiles, birds). This question, in the absence of relevant studies, is still open. It is known that PEA affect the synthesis of acetylcholine in cholinergic neurons [56], initiates apoptosis [57], affects the activity of the Ca^{2+}-dependent mammalian channels [58]. The level of PEA and PS are changed in Alzheimer's disease and Huntington's disease [59-61].

6. Conclusion

Striking features of biochemical evolution are suddenly detected when studying the effect of low temperatures upon biochemical adaptation of different classes of poikilothermic animals. Synthesis and accumulation of nonproteinogenic FAA(taurine, PS) and also PEA, which is specific to low temperature, carried out in extreme survival situations only starting from a certain stage of phylogensis. PEA and PS is a little inherent or absolutely not typical in poikilotherms in normal temperature conditions, in summer, but are becoming gradually norm in the course of evolution the even for the homeothermic animals. The aforementioned substances reflect progressive complication of biochemical organization of animals and indicates that low temperatures caused the evolutionary changes in the central nervous system.
poikilotherms. This review is an attempt to identify one of the branches of the hierarchy, which was a gap in comparative biochemistry.

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