

The upper temperature limit of life under high hydrostatic pressure in the deep atmosphere

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Keywords:
i_h h dro tati pre ure
i_h temperature
ie ophile
perthermophile
eep io phere

i_h h dro tati pre ure i a onmon fa tor in the deep ea and pro ide an i_nora le parameter of on ideo ration in all tudie related to deep life i_h temperature en ironment in the deep ea mai rd in ludin h drothermal ent and deep ediment in the u ea oor upport enormous amount of ioma produ ti it and di er it of life an mi ro e li in there are u wall h per thermophilic pie ophile hi h ope ith the dual tre e of hi_h temperature and at the ame time and pla in. A ni ant role in eo hemi al elemental & lin_ no led_e of the upper temperature limit of life under deep ea condition an help u to e timate the oundar of the iosphere and e plore it ha ita ilit on arth and in e traterre trial area ut un ertaintie remain ere e ha e ummari ed the urrent no n kno led_e of ph iolo_i al orrelation et een hi_h temperatur and ell a the effe t of on ell at hi_h temperature The effe t mainl_ ompr i et ota pe t iolo_i al inte_r it and mete oli fea i ilit The former ha een in e ti_ated in man tudie on ariou mi roor ani m from hi h e an dra a _eneral on lu ion that help ell maintain iolo_i al inte_r it under hi_h temperature or the latter e i tin_ tudie ha e pro ided lie u_e tin_ that oth hi_h temperature and halden_e mete oli fea i ilit ut iti till dif ult to dra on lu ion on the additi e effe t on mete oli m due to the la of temati anal i ere e al o propo e a serie of ue tion for further in e titation and alled for more attention on mete oli re pon e to hi_h tem perature and thi ould pro ide a dire t rid_e et een eo hemi tr and e olo_ help u to under tand the mi ro al fun tion in the deep io phere and allo u to e timate the oundarie of life and ha itat

1. Introduction

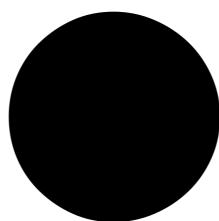
1.1. High hydrostatic pressure (HHP) is a common factor in the deep sea

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1
 reported higher thermophilic piezophile in ludin, other a teria and ar haea perature opt optimal ro th pre ure deep ea h drothermal ent re iation Topt optimal ro th temperature Tma ma imum ro th tem deep ea h drothermal ediment and eothermall heated ea oor no data

Type	strain name	a itat	in situ depth m	Topt °	Tma °	opt a	ran_e a	ner_ meta oli m	eferen e
a teria	<i>Anoxybacter fermentans</i>			–			–	e + redu tion	en_ et al
	<i>Piezobacter thermophilus</i>						–	o idation	Ta ai et al
	<i>Thiopropfundum lithotrophica</i>						–	o idation	Ta ai et al
	<i>Thermosiphon japonicus</i>						–	redu tion	Ta ai and ori o hi
	<i>Marinitoga piezophila</i>						–	redu tion	lain et al
	<i>Pseudothermotoga elfii</i>			–			–	ermentation	ouma_na et al
	<i>DSM9442</i>								a ot et al
	<i>Clostridium paradoxum</i>			–		a a	–	ethano_ene i	i et al oma et al
								ζ	
r haea	<i>Methanocaldococcus jannaschii</i>					a	– a	ethano_ene i	one et al oon aratana orn it
	<i>Methanococcus thermolithotrophicus</i>					a	–	ethano_ene i	u er et al
	<i>Methanopyrus kandleri</i>					a	–	ethano_ene i	urr et al Ta ai et al
	<i>Methanopyrus kandleri</i>					a a		e + redu tion	a he and o le ζ
	<i>Palaeococcus ferrophilus</i>						–	redu tion	Ta ai et al
	<i>Palaeococcus pacificus</i>						–	redu tion	en_ et al
	<i>Pyrococcus abyssi</i>						–	redu tion	rau o et al
	<i>Pyrococcus yayanosii</i>					a a a	–	redu tion	irrien et al ζ
	<i>Thermococcus aggregans</i>					a	–	redu tion	an_anella et al
	<i>T</i>						–	redu tion	
	<i>Thermococcus guaymasensis</i>					–	–	redu tion	an_anella et al
	<i>Thermococcus peptonophilus</i>						–	redu tion	on ále et al
	<i>Thermococcus eurythermalis</i>					a a	–	redu tion	hao et al
	<i>Thermococcus barophilus</i>					a a	–	redu tion	artein on et al
	<i>Thermococcus paralvinellae</i>					a a		redu tion	en le et al ζ
	<i>Thermococcus piezophilus</i>					a	–	redu tion	alma o et al
	<i>Thermococcus barophilus</i>							redu tion	er et al han_ et al
	<i>Thermococcus camini</i>			–			–	redu tion	ourtine et al
	<i>IrI35c</i>								
	<i>Pyrococcus</i> train					a a a		redu tion	olden and aro
	<i>Pyrococcus p</i>					a a a		redu tion	anna h et al
	<i>Desulfurococcus p</i>					a a		redu tion	anna h et al
	<i>Archeoglobus fulgidus</i>	heated ea oor		ζ	ζ	a	– ζ –	sulfate redu tion	li er et al tetter

re iation Topt optimal ro th temperature Tma ma imum ro th temperature opt optimal ro th pre ure deep ea h drothermal ent deep ea h drothermal ediment and eothermall heated ea oor deep oil produ in ell no data

^a or *M. jannaschii* train a orre pond to the hi_he tpre ure te ted ith a hi_her ro th rate ut the optimum pre ure i not no n iller et al

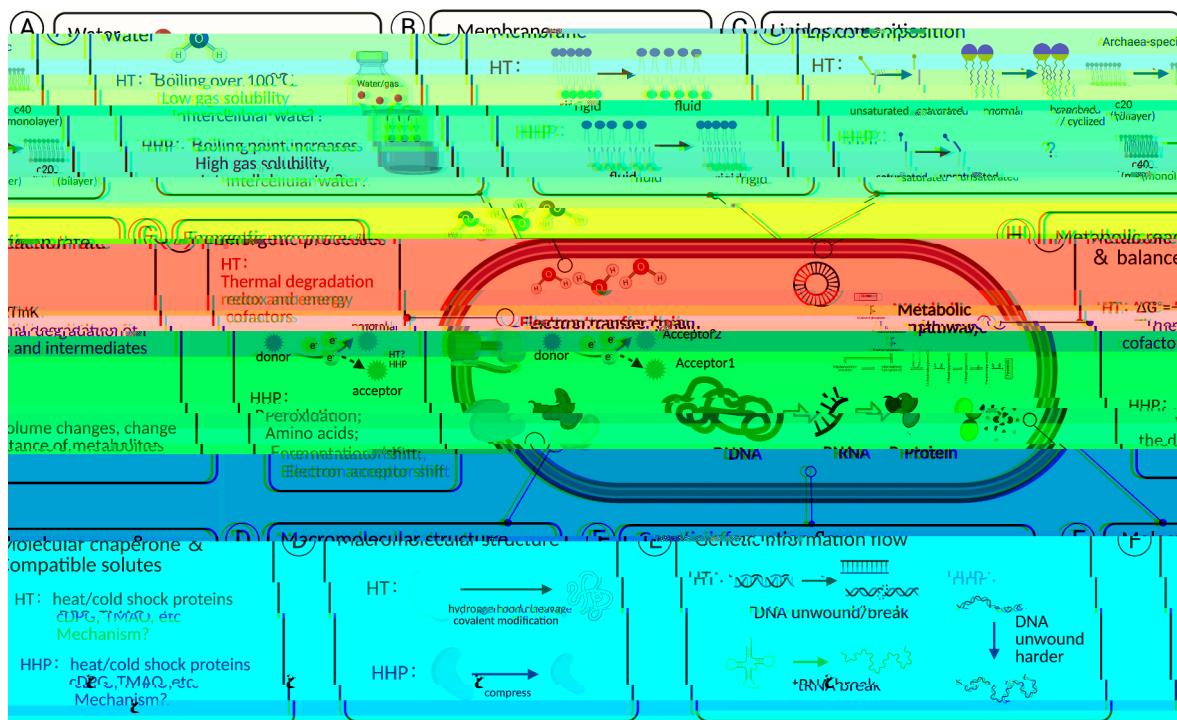
them to maintain and survive. The current record at a pressure of 10 km depth indicates a potential opportunity to investigate the upper temperature limit of life outside of hydrothermal vents.

The optimal growth temperature of *Microarania* has been proposed to be lower than the nucleic acid sequence of the genome (Auer et al., 2015). In total, 11 and machine learning methods (Li et al., 2018) showed that the optimal growth temperature of *Microarania* is 30 °C. These values are an average of predicted optimum growth temperatures from the microbial dipeptide sequence and the optimal temperature of an enzyme predicted for the unique amino acid sequence of the individual enzyme. It was found that the method to predict the optimal growth temperature of *Microarania* ophili is perthermophile mentioned above (Table 1). The values are generally all around 30 °C, except for the experimental optimum growth temperature of 25 °C. On the one hand, the deviation may be due to a larger range of the optimum growth temperature; on the other hand, thermophile with optimum growth temperature over 30 °C may have a peptide thermal stability higher than that of the protein structure of large molecule. There are other mechanisms for adaptation to high temperature environment, such as the training data model, and the microbial optimal growth temperature prediction model of the soft are one and the theoretical model of the soft are to predict the relationship between microbial maximum growth temperature and the growth temperature at the optimum growth temperature and the highest temperature of the *Microarania* do not exhibit a linear relationship (Fig. 2). This means that the highest temperature of *Microarania* are not completely determined by the genome. Therefore, the soft are prediction of the temperature adjustment needed to obtain the highest growth temperature may not be completely accurate.

Training data set at the optimum growth temperature and the highest temperature of the *Microarania* do not exhibit a linear relationship (Fig. 2).

In this section, we review the possible and potential problems arising from studies on the thermal stability of cellular membranes and macromolecules and the synthesis of molecularular

heterogeneity and compatibility under high temperature and in the environment. Some negative impact on cell and cellular component are caused by high temperature.



... hematidram of the effect of high temperature and high hydrostatic pressure on cell and molecular properties of thermophilic pieophiles. T
high temperature high hydrostatic pressure

relation and namic in the hyperthermophilic protein than in a model protein. High pressure induces that the protein undergoes a distorted structure and intermolecular interactions different from hyperthermophilic and mesophilic protein [16]. Tro et al. used ultrafast rational spectroscopy and dielectric relaxation spectroscopy to observe the random orientational motion of water molecule inside the lipid bilayer of three prototypical hyperthermophilic organisms. They found that most of the intra-cellular water is heated by the same random orientational motion as neat water here, a smaller fraction is heated by higher orientational dynamics. This intercellular water interaction proposed to bound primarily to protein and to a lesser extent to other biomolecules and ions [17]. In addition, the high pressure of proteome associated with reduced hydration after a period in hyperthermophilic pieophiles was measured using neutron scattering [18]. High pressure is proposed to have a potential effect on the molecular adaptation of the cell to high hydrostatic pressure [19]. For the detailed physiological effect of temperature and/or pressure on intercellular water in both mesophilic and hyperthermophilic condition remains a matter of interest.

3.2. Cell membrane

The cell membrane is essential for separating the intracellular environment from the outside. The integrity of the membrane enables the cell to maintain a relatively stable internal environment to support biological functions in the face of harsh or extreme external environment. The most part of the cell membrane is at high temperature. With the application of lethal heat treatment, the high mortality of the cell membrane will lead to the appearance of pores and structural damage. High pressure also has a great impact on the addition of lethal heat treatment. On the one hand, it is due to the solubility of membrane protein and the permeability of membrane protein at high temperature. Membrane protein is a membrane protein that is permeable to a certain extent and can maintain the protein's structure to a certain extent to maintain the protein's high temperature resistance. On the other hand, due to the increased accumulation of lipid chains and the increase in protein-membrane interaction [20]. The increase of protein-membrane interaction maintains the stability of the membrane and the normal function of the membrane.

Hyperthermophilic pieophiles have special membrane properties

allowing them to adapt to high temperature or even high-pressure environments. They enhance the thermal stability of membranes in reaction, the proportion of saturated fatty acids in phospholipid molecule in reaction, the length of phospholipid chain and in reaction, the proportion of isomerized and branched-chain fatty acids in hyperthermophilic archaea. A monolayer membrane structure composed of diether lipids and tetraether lipids are more rigid and impermeable than a single-layered or aliphatic membrane made of diether lipids. Their impact on membranes is limited to the high temperature environment [21]. In addition, the interaction between pentane structure and the diether lipid modification of the diether electrons are also a key for hyperthermophilic archaea to maintain the stability of membranes [22]. In addition, the adaptation mechanism enhances the stability through membrane lipid and their composition.

Generally, they help maintain membrane integrity and permeability at the high temperature treatment leading to contraction and relaxation at high temperature [23] and low temperature [24]. Other major cell membrane disorders and compare the position of fatty acid chain order and composition of the membrane is similar to that seen at -10°C under a pressure of 1 kbar [25]. Monotriton et al. found that a decrease in temperature at the same time is determined by the composition of the lipid membrane. The temperature in reaction, the entropy of the membrane and in reaction, the disorder of the high pressure will reduce the entropy. The reaction in pressure reduces the entropy of the membrane to maintain the stability of the membrane within a relatively stable range or membrane protein on the one hand, as mentioned above, the reaction in high pressure is due to the maintenance of the protein's structure to a certain extent to maintain the protein's high temperature resistance. On the other hand, due to the increased accumulation of lipid chains and the increase in protein-membrane interaction [20]. The increase of protein-membrane interaction maintains the stability of the membrane and the normal function of the membrane.

thermophile are subjected to lethal thermal stress and this is one of the areas that can increase the maximum temperature and maintain the integrity of the cell structure.

In most all cases, produce both diether and tetraether lipids. Tourte et al. (1998) and Arai et al. (2000) have demonstrated a strong correlation between the diether/tetraether ratio in lipid composition and adaptation to extreme environment in addition to high temperature. In fact, the diether/tetraether ratio measured the ratio of diether to tetraether under different temperature and pressure. Their results showed the similar trend with increasing pressure and decreasing temperature, both of which led to an increase in the proportion of diether. However, at higher temperatures, a greater proportion of tetraether was observed under low pressure and high temperature conditions. This situation in pressure and temperature also impacted the level of unsaturation of apolar lipid with an irregular polyprenoid chain containing an unsaturated isoprene derivative (Arai et al., 2000).

3.3. Macromolecules

Protein molecules play a major role in maintaining the integrity of the cell under heat stress. The effect of high temperature on biological activity is the most studied threat. Hydrogen bonding and electron interaction and hydrogen bonding in protein help to stabilize protein structure (Teiner, 1998). Protein molecules are irreversibly denatured during the high temperature stress (Harnasch et al., 2000). At the same time, amino acid residues in protein molecules are prone to undergo covalent modification at high temperature, leading to the inactivation and degradation of protein molecules. Such damage includes the deamidation of glutamine residue and a tyrosine residue and the oxidation of threonine residue (Aenige and Terner, 1998). The nucleic acid molecules and are also threatened by high temperature. The double helix of the nucleic acid molecule will open when the temperature exceeds the melting temperature and the structure of the nucleic acid molecule is easily destroyed at high temperature (Aenige and Terner, 1998) and it cannot participate in the functional transition process (Aenige and Terner, 1998).

Thermophiles exhibit physical adaptation mechanisms for adapting to "optimal growth" temperature condition. Such enhancement is not sufficient for example, Hild et al. (1998) improved the stability of the helix structure and reduced hydrophobic effect (Olden and Aniel, 1998) when the thermophile faced nearly lethal thermal stress. The high temperature adaptation mechanism cannot effectively maintain the functional activities of all these scenarios, so help maintain the stability of the biomolecules to some extent. For example, Hild et al. found that high pressure can help maintain the stability of the hydrophobicity of *M. jannaschii*. Based on their results, this effect is proposed to play a role in stabilizing hydrophobic interaction. The author also analyzed how it helped to maintain the internal packing of amino acid groups to prevent the loss of protein hydrophobicity induced by high temperature (Elh et al., 2000). Additionally, high pressure promotes protein compaction and mitigate the damage of the three-dimensional structure caused by the loss of protein hydrophobicity induced by high temperature (Elh et al., 2000). It helps to stabilize the structure of proteins and

deal with the cellular damage associated with lethal heat stress. The accumulation of multiple compatible solutes and their *de novo* synthesis in the pathogenic bacteria has been identified and shown to be associated with survival stress. There is little information on their *in vivo* physiological role and metabolic role in hyperthermophilic microorganisms from the deep sphere.

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hyperthermophilic microorganisms are all hemotrophs. They are mainly in ludin, hemoautotroph and hemoheterotroph. The hemoautotrophs are mainly anaerobic and a few are microaerobic. The arrangement of metabolism through the oxidation of sulfur compounds in different oxidation states and electron transfer agents is shown below.

Hönheit and Höhneit The metabolic pathways of heterotrophic hyperthermophiles mainly include the metabolism of peptide and sulfur compounds. Their energy source is mainly different types of electron donors such as H_2^+ , S^{2-} , sulfur compound in different oxidation states and electrons. Some of them contain more than one electron donor in the electron transfer chain. With the presence of electron acceptors, the electron transfer pathway is determined by the properties of the molecule. For example, if the molecule is a strong oxidant, it will be reduced to other organic products with molecular oxygen. If the molecule is a weak oxidant, it will be reduced to other organic products with molecular oxygen. In their metabolic pathways, they add thermal stress to the system to withstand the temperature difference between the internal and external environment. This is similar to the effect of temperature on the growth of microorganisms mentioned earlier. Therefore, the dual condition of high temperature and high salt concentration is often used in studies on hyperthermophilic bacteria. The reported halophile potential is summarized in Table 1.

Annier et al *et al* observed more amino acid requirement in a hyperthermophilic piezophile under high pressure. It was found that the low energy yield of fermentation of organic polymers to ether with energy constraint imposed high hydrolytic pressure may render *de novo* synthesis of amino acids below all unfavorable conditions. **Annier et al** suggested it is still too early to draw conclusions on how high temperature and pressure impact metabolism and metabolism in relation to the extent of factor availability. It is till insufficient to predict the quantitative analysis needed to highlight how high temperature and pressure impact metabolism individually and how operational

. onc u ion ndr ro ct

Temperature is considered the key limit for life and habitat. The diversity of life in high temperature environment in the deep sea is due to hydrothermal vent and deep sea floor processes. It is tended to be the source of the origin of both life and the habitat environment of the earth with the recognition of the significant importance of the atmosphere in high pressure thermal circulation and metabolism. The problem of life lies in the deep sea and deep atmosphere has also been handled despite the relative dearth and contradiction effect of high temperature and low biological integrity of the biological environment of metabolism. It is reported that the individuality of either high temperature or not to mention the mechanism of adaptation to deeper metabolism mainly provides a direct evidence of the hemispherical and global. The current knowledge approach regarding metabolism limit our understanding of the real microbially functional mediation, the elemental role under *in situ* pole extreme environment in the deep sea and hinder our estimation of where the boundary for both life and habitat are located.

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