

Extracellular hemoglobin and environmental stress tolerance in *Chironomus* larvae

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ABSTRACT

Hemoglobin (Hb) is one of the most common conserved molecules found in organisms belonging to all major kingdoms of life. Chironomid midge larvae are unique among the invertebrates being the only free-living group of organisms possessing extracellular hemoglobins (Hbs) in monomeric and dimeric forms floating in their hemolymph. Remarkable ability of individual species of chironomid midges to tolerate wide range of environmental stressors prevailing in their respective ecosystems has enabled chironomids to sustain as one of the most widely distributed insects in the world. Ability of different *Chironomus* spp to thrive under extreme hypoxic conditions as well as in the presence of chemical pollution made chironomid midges an efficient model system to assess the effect of environmental stress in different freshwater ecosystems. The modulation of Hb concentration has been found to be a function of different type of abiotic stressors and thus Hb of chironomid midges (Ch-Hb) has attracted the attention as a potential biomarker for environmental biomonitoring. Large body of literature on Ch-Hb accumulated mainly during sixties to eighties of twentieth century has enriched our understanding of its physiological, developmental and biochemical features. Empirical as well as *in-silico* studies carried out in recent years provided insights into many structure-function relationships of Ch-Hb. However, for the last few decades, majority of the studies were carried out to project and validate Ch-Hb as potential bioindicator for evaluating toxicants, chemical pollutants and environmental gradients of dissolved oxygen concentration. In this review, historical development of the subject has been compiled with notes on future implications of contemporary studies on Ch-Hb.

Key words: Chironomus; chironomid midge; extracellular hemoglobin; stress tolerance; hypoxia; biomarker; biomonitoring freshwater ecosystem.

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INTRODUCTION

Hemoglobin (Hb) happens to be the most studied of all proteins. Occurrence of hemoglobins (Hbs) across living kingdoms, its diversity amongst various taxa, and its complex developmental regulation makes Hbs a fascinating family of proteins and genes (Hardison, 1998; Vinogradov *et al.*, 2005). Hbs are ubiquitous among all the vertebrates known till date with little variation in the quaternary structure. On the other hand, extracellular Hbs of invertebrates are sporadic and whenever present, the Hbs exhibit large scale structural heterogeneity (Vinogradov, 1985; Weber and Vinogradov, 2001) making these globins interesting and intriguing from the physiological, ecological and evolutionary context. Comparative information on invertebrate globins is very less compared to wealth of literature available for vertebrate globins. In the past, red body fluid in the invertebrates was described as cruorine, erythrocrurion and also as hemerythrin or hemoglobin depending on the time of publication and organisms (Terwilliger, 1980). The body fluid of many invertebrates is characterized by

presence of respiratory pigments other than Hb, which are reviewed by Weber and Vinogradov (2001). Among insects, Hbs have been fully characterized till date only from *Chironomus thummi thummi*, *Drosophila melanogaster* and the larvae of the parasite botfly *Gasterophilus intestinalis*. Amongst these, *Chironomus* Hbs are extracellular, freely circulating in the hemolymph, while the Hbs of *Drosophila* and *Gasterophilus* are intracellular (Osmulski and Leyko, 1986; Pesce *et al.*, 2005; Burmester *et al.*, 2006).

Chironomid non-biting midges comprise one of the largest insect families (Insecta: Order Diptera) inhabiting diverse freshwater ecosystem across the world. Lifecycle of the non-biting midges consists of aquatic egg, larval, pupal and terrestrial adult stages. Aquatic larval stage is the longest and these larvae are conspicuous in any freshwater ecosystem due to the presence of high concentration of Hb which renders them a bright coloration and hence the genesis of its popular name as 'blood worms' (Oliver 1971; Pinder 1986; Armitage *et al.*, 1995; Ferrington, 2008). Chironomid midge hemoglobin (Ch-Hb) attracted attention of molecular evolutionary biologists with Antoine and Niessing's publication (1984)

of intron-less globin genes in *C. thummi thummi*. The lack of intron could not be explained by convergent evolution of vertebrate and *Chironomus* globins. One plausible hypothesis could be existence of an ancestral globin gene with three (or more) introns and during the divergence of the Kingdoms Plantae and Animalia, the plants retained three introns (in Leghemoglobins) and later during the divergence of vertebrates and invertebrates, all vertebrate animals retained few introns while amongst invertebrates, the chironomid midges lost all introns (Hardison, 1998). This postulation became little complicated when Hankeln *et al.* (1997) reported presence as well as absence of central introns in many European and Australian species of *Chironomus*. The speculated insertion by gene conversion like mechanism indicated transposition type events leading to Hb gene polymorphism found in extant *Chironomus* spp. (Gruhl *et al.* 2000). However, phylogenetic analyses based on amino acid sequences of Hb variants as well as nucleotide sequences of corresponding Hb genes indicated that evolution of chironomid Hb (Ch-Hb) genes arose through series of successive gene duplication followed by diversification (Hankeln *et al.*, 1998). It is now a widely accepted view that the divergence of Ch-Hb genes occurred from a monomeric globin lineage 270-300 million years ago (Goodman *et al.*, 1988). Number of globin genes coding for extracellular hemoglobins in *C. thummi thummi* was estimated to be 27, although other studies in *C. tentans*, the number exceeded over 40 which is till date the largest Hb gene family reported in any organism (Hankeln *et al.*, 1998 and references therein).

In the past, mode of larval life and ecological parameters were linked with properties of Hb in different species of *Chironomus*. Among these, intrinsic factors like hormones and extrinsic factors like hypoxia and presence of toxicants were the major causative factors implicated with the modulation of Hb concentration and corresponding gene expression (Osmulski and Leyko, 1986). Environmental biologists have been using different stages of *Chironomus* as freshwater biomonitoring agent for the last three decades or so (Choi, 2004). Morphological deformities have been used as endpoints to assess the impact of pollutants/ toxicants including radionuclide, heavy metals and other stressors (Ingersoll and Nelson, 1990; Clarke, 1993; Armitage *et al.*, 1995; Hamalainen, 1999; Dias *et al.*, 2008). Salivary gland polytene chromosome of chironomid larval midges served as an excellent cytogenetic tool as indicators of metal-genotoxicity and other stressors (Nath and Lakhotia, 1989; Martin, 1996; Bernabo *et al.*, 2011; Michailova *et al.*, 2011, 2012; Michailova and Petrova, 2015; Lencioni *et al.*, 2016). Additionally, stress proteins like Hsp70 (Nath and Lakhotia, 1989; Karouna-Renier and Zehr, 1999; Morales *et al.*, 2011; Datkhile *et al.*, 2011) and

stress-responsive metabolites (Thorat *et al.*, 2017) of chironomid larvae were also used as indicator biomolecules for assessing environmental stress. However, in this review, I shall focus only chironomid Hb, as and when attempts were made to establish Hb as bioindicator of environmental stress.

HEMOGLOBIN IN CHIRONOMID MIDGES

Developmental features

Heterogeneity and ontogenic difference of Hbs among different larval stages were first reported by Manwell (1966). Thompson and English (1966) found abundance of larval hemoglobin towards later stages of larval development and later on, an elaborate account of ontogenic changes in Hb synthesis in *C. tentans* was published by English (1969). Appearance of Hb could be first noticed during the second instar larvae and the Hb content reached up to 90% of the total hemolymph protein by the time larvae moved to fourth instar of development. Bergtrom *et al.* (1976) demonstrated larval fat body as the major site of Hb synthesis and secretion. Radiolabelling studies also confirmed that the biosynthesis of both heme and globin could take place in the larval fat body. The seminal study of Bergtrom and his group was the first demonstration of Hb synthesis using organ culture in any invertebrate. Tissue specificity of Hb synthesis in the larval sub-epidermal fat body of *C. thummi* was further confirmed by Vafopoulou-Mandalos and Laufer (1983) using autoradiographic technique. Hbs are exported to hemolymph from the fat body cells by means of N-terminal signal peptides. By the time the metamorphosis concludes at the adult stage, more than 95% of the Hb can be found as degraded. Using meticulously designed experiments with radioactive tracers (⁵⁹Fe-labelled Hb) Schin *et al.* (1974) demonstrated gradual decline of Hb concentration post-fourth instar larval period till the pharate adult stage in *C. pallidivittatus*. Hbs are degraded in the Malpighian tubules (Jarial 1988). The mature adults were hardly found to possess any Hb which underwent degradation represented by increment in bile pigments, identified as bililatrene (Schin *et al.*, 1974). Detailed studies, especially during 80's decade of the last century, established Ch-Hbs as potent allergens (Trewitt and Bergtrom, 1981; Cranston, 1988; Baur *et al.*, 1986). Other studies implicated degraded products of Hb are responsible for the high prevalence of sensitization producing conjunctivitis, rhinitis and asthma by inhalation. IgE mediated hypersensitivity was reported among handlers of fish food in their workplace because chironomid midges are popularly used as nutrients in fish aquarium (Ballesteros *et al.*, 2006).

Molting in holometabolous insects are regulated by

hormones like ecdysones and juvenile hormones. On the other hand, a strong correlation of molting and appearance of new Hbs in a stage-specific manner suggested hormonal role and its control over HBb synthesis (Trewitt and Bergtrom, 1981). Subsequently, using radio-isotope based tracing techniques, Vofopoulou-Mandalos and Laufer (1982, 1984) demonstrated how ecdysterone and Juvenile hormone regulate Hb synthesis during larval development of *C. thummi*. They also suggested that both these hormones could play a role in modulating the expression of Hb gene.

Cytogenetic features

Although there have been plenty of cytogenetic studies on various aspects using *Chironomus* larval salivary gland polytene chromosomes, literature on any cytogenetic studies relevant to chironomid Hb is almost nil except two seminal studies by Hankeln *et al.* (1988) and Schmidt *et al.* (1988). Globin gene cluster from the subspecies *C. thummi piger* was localized to chromosome II (the chromosomal band position F2b2) by *in situ* hybridization (Hankeln *et al.* 1988). The same laboratory embarked on an exploration of distribution of Hb genes on polytene chromosomal regions of different *Chironomus* species. Schmidt *et al.* (1988) investigated 13 *Chironomus* species and their robust data demonstrated localization of Hb genes coding for monomeric and dimeric Hb proteins on chromosome arms E and D respectively. These cytogenetic data on the distribution of Hb genes on polytene chromosomes agreed with the cytological data of Keyl (1962), often used to establish evolutionary relationships among the different *Chironomus* species.

There has been a long gap in the literature concerning any cytogenetic studies linked with chironomid Hb subsequent to Hankeln *et al.* (1988) and Schmidt *et al.* (1988). Much later, my laboratory carried out cytogenetic study in the oriental non-biting midge *C. ramosus* which has relevance to larval Hb content as a function of dietary iron content. I used two variant larval populations of *C. ramosus* generated by Babrekar (2009) under differential simulated conditions of providing one and half fold more iron (Fe) containing diet *versus* low-Fe diet relative to the standard food recipe prescribed by Nath and Godbole (1998). The resultant larval population exhibited concomitant higher and lower Hb level in larval hemolymph (Babrekar 2009). Interestingly, the two variant group of larval population showed heterochrony with respect to pupal emergence calendar (Tab. 1). One of the cytogenetic hallmarks of prepupal developmental stage of *C. ramosus* is the appearance of Balbiani Ring (BR) at 18C locus of chromosome IV (Fig. 1). I studied the appearance and extent of puffing of 18C locus in the two developmentally variant larval populations with respect to their Fe-rich *versus* low-Fe dietary rearing conditions concomitant with high and low Hb level (Tab. 1). The findings showed a cytogenetic correspondence of heterochrony in the developmental timing of metamorphic event as indicated by the cytogenetic marker (onset of BR manifestation). These data warrant further studies to explore involvement of hormonal and molecular events signaling critical levels of Hb in the larval hemolymph prior to metamorphosis.

Tab. 1. Heterochrony in onset and extent of manifestation of Balbiani Ring (BR) activity of 18C locus of chromosome IV (Reference map: Nath and Godbole, 1997) in two groups of larvae of larvae of *Chironomus ramosus*, grown either in Iron (Fe) rich or low diet (Babrekar, 2009) possessing high (0.8 ± 0.09 gm Hb/gm body weight) and low (0.5 ± 0.04 gm Hb/gm body weight) hemoglobin content. Puffing index is estimated from the ratio (mean \pm SD) of chromosomal locus 18C (which exhibits developmental stage specific expression of BR) width to that of 18B locus (which never shows any puffing).

Larval age (days post hatching)	18C BR puffing index	
	Larvae with low Hb level grown with low-Fe diet	Larvae with high Hb level grown with Fe-rich diet
15	1.00 \pm 0.01	2.00 \pm 0.02
16	1.00 \pm 0.02	2.30 \pm 0.01
17	1.00 \pm 0.01	3.00 \pm 0.02
18	1.75 \pm 0.02	3.75 \pm 0.03
19	2.00 \pm 0.02	4.25 \pm 0.03
20	2.00 \pm 0.02	4.00 \pm 0.03
21	3.00 \pm 0.02	Onset of pupation
22	3.25 \pm 0.03	
23	3.00 \pm 0.01	
24	3.00 \pm 0.02	
25	Onset of pupation	

Biochemical features

Among all the chironomid midges known till date, hemoglobin (also referred to as erythrocrucorin in literature available 4-5 decades earlier) from *Chironomus thummi thummi* has remained as the most studied molecule. Different components of hemoglobin from *C. thummi thummi* (CTT-Hb) revealed many unique features which could be correlated with the structure, function and adaptation (Buse *et al.*, 1979). Pioneering work by G. Braunitzer and his collaborators characterized twelve different isoforms of Hbs from the hemolymph of *C. thummi thummi*, abbreviated as CTT as prefix followed by roman numerical representing chromatographic fractions. Amongst these, CTT-III was the first high resolution structure of any invertebrate Hb, which turned out to be the subject of many biochemical, biophysical and molecular biological studies (Green *et al.*, 1998 and references therein). Other studies of *Chironomus* species of plumosus group revealed presence of sixteen different Hb fractions (Shobanov 2004).

Hbs of *C. thummi* exist in the hemolymph in a pH dependent equilibrium between monomers and dimers (Braun *et al.*, 1968; Sick *et al.*, 1972). Thomson *et al.* (1968) demonstrated larval Hbs of *C. tentans* as exclusively monomeric with a molecular weight of ~15,900. There has been contradicting views and findings regarding molecular weight of Hbs of different *Chironomus* species, including the monomeric *versus* dimeric forms of Hbs. It started from the initial observation of Svedberg (1934) which was re-estimated by others (Osmulski and Leyko, 1986). Series of findings in *C. thummi thummi* confirmed subsequently

that isoforms of electrophoretically different Hbs exist either as monomeric or as dimeric form and maintain a pH dependent equilibrium (Braun *et al.*, 1968; Pfletschinger *et al.*, 1980). Chironomid Hbs can bind oxygen with high affinity (P_{50} =0.6-1.5 Torr at pH 7 and 20°C) and in a non-cooperative manner. Under physiological conditions, O₂ binding can get modulated by changes in pH of milieu which is known as Bohr effect. Chironomid Hbs are exceptional for showing absence of cooperativity and an alkaline Bohr effect except for CTT-I (Wollmer *et al.*, 1972; Steffens *et al.*, 1977; Burmester and Hankeln, 2007).

Under physiological pH condition of around six monomeric and homodimeric Hb components correspond to molecular weight of about 16,000-17,000 and 32,000-34,000, respectively. Notwithstanding the fact that almost all Hb related biochemical and biophysical characterization was carried out on different European chironomid midges, the first report of Hb of any oriental chironomid midge species was reported by Das and Handique (1996). Eleven Hb components from Indian midge species *C. ramosus* were described which were found to comprise three monomers, seven dimers and one on both monomeric and dimeric forms. This is in contrast with ten and twelve isoforms of *C. plumosus* and *C. thummi thummi* respectively. Many workers in this field developed and refined biochemical procedures of preparation of Ch-Hb for experimental and analytical studies. Ruf *et al.* (1994) published comprehensive methods for preparation and characterization of CTT-Hbs which was standardized by many scientists for quantitative assay of Ch-Hbs, often with modification (Ronad and Nath, 2017). Besides conventional biochemical studies, an electrophoretic method to obtain Hb profile using denaturing polyacrylamide gel was shown to be useful for chironomid species identification (Oh *et al.*, 2014). This is a first report of an interdisciplinary approach for a potential application of *Chironomus* Hb in taxonomy.

In the literature, three-dimensional structure of deoxy hemoglobin (component III) from *Chironomus thummi thummi* has remained the only source for a reference structure to carry out biochemical and bioinformatics studies (Gaikwad *et al.* 2016). From the evolutionary point of view, it is important to note that amino acid in heme binding pockets of Hb molecules underwent drastic evolutionary diversifications resulting in complete absence of amino acids like isoleucine and glutamine in human Hb during the course of evolution since the speciation of non-biting chironomid midges (Tab. 2). Such biochemical combined with bioinformatics analyses hold promise for future revelation of many unknown facts of evolutionary transitions of invertebrate Hb to vertebrate Hb.

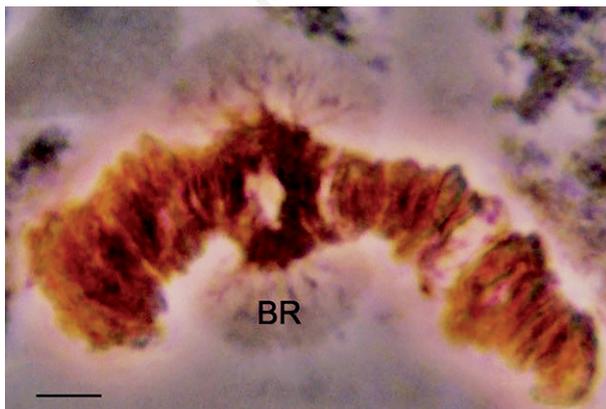


Fig. 1. Representative photomicrograph of chromosome IV from the salivary gland polytene chromosome complements of *Chironomus ramosus* (Nath and Godbole, 1997) showing a conspicuous Balbiani Ring (BR) at 18C locus. Scale bar: 10 μ m.

CHIRONOMUS HEMOGLOBIN AND STRESS TOLERANCE

Hypoxia

One of the most remarkable ability of chironomid midge larvae has been their tolerance to extreme hypoxic conditions. Hb and the bright red color of chironomid midge larvae were linked as early as in the 19th century (Rollett, 1861; Lankester, 1872). Miall and Hammond (1900) and Pause (1919) speculated Hb's role in oxygen storage by chironomid larvae. Fox (1955) observed an elevated level of Hb synthesis in chironomid larvae in poorly aerated water. Ch-Hb facilitates absorption of oxygen from the surrounding water at partial oxygen pressures ranging from 10-50 Torr (Redecker and Zebe, 1988; Zebe, 1991). There have been other studies demonstrating how larvae could carry out aerobic metabolism under hypoxic conditions. Concentration of Hb as a function of increasing oxygen depletion was established in *C. riparius* by Grazioli *et al.* (2016) who provided metabolic explanation of Hb's role during anaerobiosis and how Hb might play a role in short-term oxygen storage.

Kawai *et al.* (2004) evaluated respiration rates and survival rates of Japanese *Chironomus* spp. under anoxic conditions and found high Hb content in chironomids living in lentic environments in contrast to those living in lotic waters. Moreover, survival duration was found longer with increasing respiration rates and Hb contents. Varying level of Hb has been reported to depend on the ambient concentration of dissolved oxygen in the laboratory reared tropical Indian species *C. ramosus* using a fabricated hypoxia-inducing device customized for experimentation with chironomid larvae (Ronad and Nath, 2017). They reported that simulated hypoxic conditions of 0.5ppm of dissolved oxygen could induce a multifold increase in Hb gene expression as a function of

time of exposure. While finding an evolutionary logistics of chironomid midge's remarkable ability to tolerate extreme hypoxic environment, one needs to find out an explanation of proposition of Burmester *et al.*, (2007) that extracellular Hbs of chironomid midges originated as an evolutionary derivative of intracellular Hbs in response to selection pressure of hypoxic environment.

Chemical pollutants/toxicants

Larval discoloration due to decreased red pigmentation typically associated with Hb was used as one of the early indicators of heavy metal (*e.g.*, copper) toxicity (Majumdar and Gupta, 2012) in *C. ramosus*. Gradual loss of red color was also observed by MacDonald *et al.*, (2004) due to perfluorooctane sulfonic acid (a fluorinated surfactant) toxicity. Larvae of *C. riparius* exposed to cadmium showed reduction in total Hb content (Choi and Ha, 2009). Other studies on natural population of *Glyptotendipes* spp and *Endochironomus* spp. exposed to heavy metal contamination (cd, Pb, Hg, Cr, Cu, Zn) revealed significant reduction in genetic diversity and variable level of Hb protein diversity (Jacobs and Marie, 2006). Amongst environmental pollutants/ toxicants many chemical substances are redox-active. Studies carried out by Choi and Roche (2004) in *C. riparius* suggested auto-oxidation of Hb in response to such redox-active chemicals. In one of the interesting findings in *C. riparius* adapted to acid environment suggested remarkable buffering capacity of Hb in pH as low as 3.5. This Canadian strain of *C. riparius* possessed twice the content of other strains of same species inhabiting non-acidic environment (Jernelöv *et al.*, 1981).

Chemical-specific Hb gene expression in *C. tentans* was reported by Lee *et al.*, (2006) while evaluating consequences of larval exposure to pollutants. Interestingly, alkyl phenolic compounds elevated the expression of Hb gene while pesticides decreased the expression. Extension of similar studies by the same research group revealed that *C. riparius* globins could exist in both inducible and constitutively expressed forms. Presence of multiple globin isoforms facilitated the organism to adapt to pollutants (Ha and Choi, 2008). They observed downregulation of oxyhemoglobin on exposure to bisphenol A and chlorpyrifos suggesting its elevated autoxidation to methemoglobin. Similar to hypoxia, oxygen deficiency could be induced chemically by herbicide treatment (*e.g.* Atrazine) and *C. tentans* larvae were found to modulate Hb level by regulating its gene expression at the level of transcription (Anderson *et al.*, 2007). Quite often, due to chemical specific response of Ch-Hb, many studies remained inconclusive for validating appropriateness of Hb as bioindicator of all types of pollutants/ toxicants *per se* (Darville and Whilm, 1984).

Tab. 2. Bioinformatics analysis of presence (+) and absence (–) of amino acid residues in the heme pocket lining in hemoglobins of *Chironomus thummi thummi* (CTT-Hb) [PDB id: 1ECA] and human (Hu-Hb) [PDB id: 2DN2] (data from Dr. Sangeeta Sawant, Bioinformatics Center, SPPU, Pune, India, *personal communication*).

Amino acid in heme-binding pockets of Hb molecule	CTT-Hb	Hu Hb
Isoleucine (Ile)	+	–
Tyrosine (Tyr)	–	+
Alanine (Ala)	–	+
Glutamine (Gln)	+	–
Asparagine (Asn)	–	+

Radiation

Among earliest reports on radiation tolerance of chironomid midges, cytogenetic findings of Blaylock (1966) seems to be the most noteworthy followed by other studies from Michailova's (Michailova and Petrova, 1994, 2015) and Nath's research group (Datkhile *et al.*, 2009a and cross references citing previous works on radiation induced deformities in *Chironomus* spp.). Detailed studies on the cellular and molecular basis of gamma radiation tolerance in Indian species of midge *C. ramosus* were initiated by Nath and his collaborators (Rita Mukhopadhyaya, BARC, Mumbai and Sangeeta Sawant, S.P. Pune University, Pune, India) who established chironomid midges as one of the insect groups tolerant to high doses of gamma radiation (Datkhile *et al.*, 2009a, 2009b, 2011, 2015). Recently the focus of this research group shifted to Hb of *C. ramosus* larvae subjected to high doses of gamma radiation and a few interesting facts got revealed (Kumar *et al.*, 2017, Gaikwad *et al.*, 2016) which are worth mentioning in this review.

Sensitivity of Ch-Hb and human hemoglobin (Hu-Hb) to gamma radiation (2400 Gy with dose rate of 52.6 Gy/min) was evaluated and a remarkable stability as well as integrity of Ch-Hb were observed when compared to Hu-Hb. Irradiated Ch-Hb exhibited a slight decrease in the intensity in UV-visible absorption spectrum without any alteration in Soret peak position at 415nm, characteristic for Hbs. On the other hand, Hu-Hb showed molecular damage and a shift in Soret band when subjected to similar dose of gamma radiation. Unlike Ch-Hb, additional biophysical analyses indicated relatively higher sensitivity of Hu-hb with reference to heme pocket of Hb molecules (Gaikwad *et al.*, 2016). This was interesting to note from the evolutionary point of view because Hu-Hb represents vertebrate Hbs which underwent little changes since the time of divergence of vertebrates and invertebrates. A sequence and structure-based bioinformatic analysis (Gaikwad *et al.*, 2016 and unpublished data of Sawant and Nath) revealed higher hydrophobicity around heme pockets of Ch-Hb when compared to Hu-Hb. The remarkable resistivity of chironomid midges to adverse environmental conditions in general could be attributed to hydrophobicity of heme pockets of Ch-Hb and molecular evolutionary history of this group of insects.

CONCLUSIONS AND FUTURE PERSPECTIVE

Unlike the vertebrates, Hbs among the invertebrate animals are not ubiquitous. Despite rapid evolutionary divergence, vertebrate Hb retained structural similarity to greater extent when compared to repertoire of invertebrate Hbs. Much remains to be understood regarding extensive interspecific polymorphism and the presence of large

number of isoforms of Hb which are developmentally regulated during larval phase in chironomid midges. We do not know much about the selective advantage for multiplicity of Hb loci in *Chironomus* spp. Perhaps, temporal pattern of developmental stage specific expression of Hb loci in the larvae provides adaptive fitness in aquatic environments with very low oxygen tensions as well as in the presence of pollutants and toxicants in the ambient water.

Although Ch-Hb has been advocated by many as a candidate biomarker for environmental monitoring purpose, one should ensure whether Ch-Hb always qualifies to be a 'unique indicator' for any particular stressor. For that matter, can Ch-Hb either qualitatively or quantitatively distinguish hypoxia-anoxia related stress response *versus* consequences due to heavy metal pollution in the same ecosystem? Nevertheless, one needs to be cautious before euphorically proposing Ch-Hb as bioindicator of a particular stressor in natural habitats where other stress-inducing variables also prevail. Much remains to be done to monitor synergistic as well as antagonistic impact on the hemoglobin of *Chironomus* spp. in freshwater ecosystem where the population is subjected to more than one environmental stressors. Presence of large number of Hb variants in any given species of chironomid midge is also intriguing. Perhaps, this multiplicity of Hbs would be explained in future studies at the functional level with reference to tolerance towards multitude of stressors in the natural habitats of different midge species.

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