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Hemoglobin alpha in the blood vessel wall

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Abstract

Hemoglobin has been studied and well haracterized in red blood cells for over one hundred years. However, new work has indicated that the hemoglobin alpha subunit (Hba) is also found within the blood vessel wall, where it appears to localize at the myoendothelial junction (MEJ) and plays a role in regulating nitric oxide (NO) signaling between endothelium and smooth muscle. This discovery has created a new paradigm for control of endothelial nitric oxide synthase activity, nitric oxide diffusion, and ultimately, control of vascular tone and blood pressure. This review will discuss the current knowledge of hemoglobin's properties as a gas exchange molecule in the blood stream, and extrapolate the properties of Hba biology to the MEJ signaling domain. Specifically, we propose that Hba is present at the MEJ to regulate NO release and diffusion in a restricted physical space, which would have powerful implications for the regulation of blood flow in peripheral resistance arteries.

Keywords

hemoglobin; hemoglobin alpha; alpha thalassemia; myoendothelial junction; Cytochrome B5 reductase 3; nitric oxide

Introduction

For over one hundred years¹, hemoglobin has been known as the gas exchange molecule found within red blood cells (RBC) that is responsible for delivering oxygen to tissues and subsequently removing carbon dioxide². The structure of normal functional adult hemoglobin (HbA; Figure 1A) is composed of twin alpha and beta globin subunits ($\alpha_2\beta_2$), each of which contains a heme, or iron ion, within a heterocyclic ring of four pyrrole

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 (C_4H_4NH) molecules known as a porphyrin. The assembled tetramer can interact with the aforementioned gases, as well as carbon monoxide and NO. Today, this paradigm in RBC remains unchanged. However we³, and others^{4,5}, have demonstrated that Hba is expressed in the blood vessel wall but in contrast to other globins in the blood vessel wall^{3,6,7}, Hba localization and physiological effects are concentrated at myoendothelial junctions (MEJs) in the endothelial cells (EC) lining the lumen of blood vessels.

Although the discovery of Hb α in the blood vessel wall is unique, there have been clues from human diseases of Hba deletion, possibly independent of RBC function, that have already indicated its possible presence. The Hba gene is located on chromosome 16 and has two identical but duplicated coding sequences, HBA-1 and HBA-2. Genetic deletion or lossof-function mutations to the Hba alleles are known as alpha-thalassemia, a condition which is defined by the severity of changes to red blood cell indices^{8,9}. Patients with alpha⁺ thalassemia (silent carriers) will have deletion or loss of function mutations to one of the HBA-1 or HBA-2 alleles. However, compensation enables them to remain asymptomatic and many go undiagnosed. Deletion of two of the HBA-1 and HBA-2 alleles causes alpha⁰ thalassemia, which presents with moderate or severe RBC indices, depending on the deletion type or mutation⁹. In terms of the vascular phenotype, this frequently includes moderate hypotension. Deletion of three of the HBA-1 and HBA-2 alleles is known as HbH disease and characterized by hemolytic anemia, hepatosplenomegaly, and the formation of a tetramer of β chains *in vivo*. Due to severity of RBC indices, it remains unclear if the vascular phenotype remains consistent with alpha⁰ thalassemia or is hidden by development of other pathologies. Hydrops fetalis syndrome (HP Bart Syndrome) occurs with deletion of all of the Hb α alleles and results in death in utero¹⁰. The vascular phenotype of dilated cerebral arteries is observed via ultrasound., and also used to assist in the diagnosis of the disease although difficult to delineate from the accompanied severe anemia (Table 1) 11,15 . Regardless, this would correlate with observations from patients with different degrees of Hga deletion presenting with differences in capillary diameter.¹⁶ In each of these cases, a role for Hba in RBC is challenging to fully explain the vascular phenotype presented. However, reduction or deletion of Hb α in EC of the resistance arteries, where it is hypothesized to regulate NO delivery to vascular smooth muscle cells (VSMC), could be a reason. Future research in this area could lead to important new insight into understanding the pathology as well as diagnosis related to these diseases.

Hemoglobin alpha, certae sedis

The first evidence of localization and function for Hba in the blood vessel wall was seen in MEJ from resistance arteries in the systemic vasculature³. Resistance arteries are the small arteries that contribute the greatest amount to peripheral resistance and thus overall blood pressure regulation^{17,18}. In resistance vasculature heterocellular communication is critical to the maintenance of vascular tone and blood pressure (for review, see ^{19,20}). Crucial to heterocellular communication in resistance arteries are the presence of MEJs. The MEJ is the physical link between EC and the VSMC, characterized as a small protrusion of mostly EC (approximately 0.5 m wide and long) through the internal elastic lamina, linking with VSMC through gap junctions.¹². Myoendothelial junctions are found throughout the vasculature; however, a gradient is observed such that the junctions are more prevalent as

the diameter of the vascular tree decreases²¹. Thus, MEJs represent the closest physical location of EC to VSMC in the small resistance arteries. The MEJ is acknowledged as an important signaling microdomain (for review, see (Billaud et al., 2014)²²) not only because of its spatially limited structure, but also because the MEJ serves as the gateway between EC signal transduction to the VSMC²². For instance, S-nitrosylation/denitrosylation of connexin 43 is controlled by specific proteins localized to the MEJ, including S-nitrosoglutatione reductase (GSNOR), endothelial nitric oxide synthase (eNOS), and IP₃ receptor type 1 (IP3RI)²³. Together, these proteins participate in localized signaling that promotes negative feedback on vasoconstriction induced by α_1 -adrenergic receptor activation.

Importantly, while eNOS is localized to intracellular regions including plasma membrane caveolae and the Golgi complex, it is also localized to the MEJ^{24,25}. Endothelial cells exhibit a strict and varied control over the production of NO (reviewed in (Shaul, 2002)²⁶) and the predominant enzyme responsible for the generation of NO in these cells is eNOS. This localization of eNOS at the MEJ presumably occurs because NO is a highly reactive gaseous free radical, enabling targeted release of NO at proximal sites to the overlying VSMCs, where it functions as potent vasodilator. Subsequent work demonstrated that eNOS and Hba reside in close proximity to each other and potentially form a macromolecular complex at the MEJ³, where they provide a regulatory mechanism for NO-mediated vasodilation (Figure 2).

Three general outcomes await NO once it is produced in EC: it can 1) diffuse into the blood stream where it is rapidly scavenged, first by cell free hemoglobin in the plasma and even further by hemoglobin in RBC²⁷, 2) diffuse to neighboring VSMCs and cause vasorelaxation, 3) be scavenged by any number of molecules, such as reactive oxygen species (ROS) (for review, see Martinez et al., 2009²⁸). In the blood stream, most NO diffuses across its concentration gradient into RBC, where it is either scavenged or stored. NO scavenging proceeds via oxidation by oxyhemoglobin, which produces methemoglobin and nitrate. This is the favored reaction; however, NO can also bind directly to deoxygenated hemoglobin in a simple addition reaction to form iron-nitrosyl-hemoglobin (Hb-NO)²⁹. Furthermore, it has been proposed that nitric oxide reacts in an oxygen-sensitive mechanism with the conserved Cys93 residue of the hemoglobin β -chain³⁰; however, these findings have been intensely debated^{31–33}. Finally, data has been suggested that eNOS is present in RBC³⁴, which possibly contributes to vasoprotectivity^{35–37}. Once in the VSMC, NO activates soluble guanylyl cyclase (sGC) by binding to its heme moiety, allowing a several hundred-fold increase in the catalysation of GTP to cGMP³⁸ and causing vasodilation through several mechanisms³⁹. Scavenging of NO occurs in pathologies that result in, produce, or maintain elevated oxidant stress, especially when accompanied with an inability to be counter balanced by antioxidants. Increased ROS can scavenge NO directly or alter pathways that mediate NO production. The multiple fates of NO in the vasculature is further complicated by its very short half-life (<5 seconds)⁴⁰. Conceivably, in the EC at the MEJ, any one of the above mechanisms may be in $play^{41}$.

An important regulator in the binding of NO (and oxygen) to hemoglobin is the oxidation state of the heme iron: ferrous (Fe²⁺), or ferric (Fe³⁺). The ferric state of hemoglobin is also frequently known as methemoglobin (MHb)^{42,43}. However, the HbA in the bloodstream

contains less than 1% MHb due to the fact that it possesses a higher affinity to oxygen and excess MHb can result in tissue hypoxia and even death⁴⁴. The oxidation state of iron also controls the sensitivity of Hba to NO; the Fe²⁺ state rapidly scavenges NO and the resulting production of nitrate and methemoglobin decreases NO bioavailability, whereas the heme moiety in the Fe³⁺ state reacts slowly and transiently with NO, allowing for increased diffusion of NO into smooth muscle^{45–47}. This reaction would presumably hold true for the Hbg found in the MEJ. In RBC, a methemoglobin reductase, cvtochrome b5 reductase 3 (CytB5R3, also known as diaphorase 1), is present to recycle the Fe³⁺ state and prevent the accumulation of MHb. When CytB5R3 is absent due to genetic defect or ingestion of oxidizing toxins the resulting methemoglobinemia⁴⁸(famously characterized as a bluish tint to the skin) causes decreased tissue oxygenation, hypoxia, and cyanosis⁴⁹. Not surprisingly, the enzyme CytB5R3 found in RBC is also found in EC and at the MEJ [although not as strictly localized as Hba). However, in contrast to the RBC where the Fe²⁺ oxidation state predominates, the MEJ heme moieties in resistance arteries exist in both states (Fe³⁺ approximately 58%). Knockdown of CytB5R3 reveals altered reactivity in ex vivo vessels to adrenergic and endothelial dependent NO stimulus, revealing a permissive effect of CytB5R3 on NO bioavailability³. Interestingly, the CytB5R3 inhibitor and anti-thyroid drug propylthiouracil (PTU) also presents with a reduction in blood pressure in rats⁵⁰. This could be explained by PTU preventing the CytB5R3 enzyme from reducing the Fe³⁺ heme moieties in Hb α the MEJ, thus maintaining the slower NO scavenging reaction and allowing for increased NO diffusion into the VSMC. The evidence increasingly indicates that Hba could play an important role in regulating NO bioavailability through control of the heme oxidation state.

Lastly, it's possible that Hb α is found in resistance arteries to chelate excessive NO. This rationale is based on the fact that conduit arteries have been previously characterized as largely dependent on NO for vasodilator activity, whereas smaller resistance vessels rely on the action of endothelium-derived hyperpolarizing factor (EDHF)^{22,51}. However, there is no observable difference in the total protein expression of eNOS in carotid and third order mesenteric arteries (i.e., resistance arteries; Figure 3). Thus, Hba at the MEJ in resistance arteries may be chelating the NO generated by eNOS, allowing for EDHF to remain dominant for increased endothelial dependent vasodilation, and maintaining strict control over NO for other cellular functions (e.g., negative feedback after vasoconstriction). It remains unclear how Hb α scavenges the excessive NO, and whether this action results in a permanent loss of NO or simply a reservoir of NO that can be released upon appropriate stimulus. It is also possible that the amount of eNOS between resistance arteries and conduit arteries is inconsequential because the eNOS that is present in resistance arteries is uncoupled as compared to conduit arteries, negating eNOS generation (e.g., ^{52,53}). It is unclear why such a potent enzyme would still be present if this were the case, however the possibility exists and it is clear more work needs to be done in this regard.

The importance of NO scavenging by Hba at the MEJ is predominantly derived from its location. Nitric oxide scavenging by RBC hemoglobin requires diffusion through the EC monolayer, the glycocalyx, the plasma, and then interacts with cell free hemoglobin or must cross the RBC membrane before it reaches HbA. Based on these assumptions, it is tempting

to speculate that the likelihood of locally produced NO at the MEJ interacting with Hba is greatly increased due to it's proximity to the NO source. However, NO produced on the luminal side of the endothelium would favor scavenging by RBC hemoglobin. This implies that although the relative abundance of Hb in RBC may be greater, it's importance for the regulation of diffusion of NO into VSMC may be reduced, although this may change with deviation from basal conditions or pathological states.

Why hemoglobin a in the endothelium?

One of the interesting questions that arises from the discovery of Hba at MEJs of EC is, why Hba, and not Hb β ? Wouldn't both subunits be better than one for microcirculation where MEJs are prevalent? As of yet there is no clear answer to this question.

Expression of both hemoglobin subunits has beef reported in many cell types beyond erythrocytes^{54–59}, which has sparked hypotheses of functions in addition to oxygen transport and delivery. The oxygen transport by Hb is specific to vertebrates with globins in other walks of life having a variety of functions such as NO dioxygenase and peroxidase activity⁶⁰. Individually, the Hb subunits have damaging effects; thus, the assembly of HbA is regulated. Only Hba has a chaperone, alpha hemoglobin stabilizing protein [AHSP], which maintains Hba solubility and reduces reactive oxygen species production by stabilizing the Fe³⁺ state⁶¹. AHSP binds to Hb α in proximity to the Hb β binding site [Fig. 1B] and is displaced during the assembly of Hb. Therefore, Hba subunits may be considered the less toxic of the two subunits for additional physiological roles due to the added protection of AHSP. The Hb α and Hb β subunits have very similar structures [Figure 1C–D] yet very different amino acid sequences (43% identity]. There are some structural rearrangements of Hba upon binding AHSP; however, the overall structure is very similar to both Hb α and Hb β in the heterotetramer. Both subunits similarly bind haem; thus, in terms of oxygen binding and reactivity the subunits are similar. However, the sequence differences between the subunits could certainly provide differences in signaling. For instance, Hba molecular interactions that regulate NO signaling at the MEJ³ may be mediated by distinct amino acid regions of Hb α that are different than Hb β such that Hb β would not elicit the same signaling cascade. More work on this fundamental question will certainly be required.

Vasculature effects of genetic hemoglobin deletion

In mice, similar to humans, deletion of all of the Hb α alleles is lethal, making genetic studies on animals *in vivo* difficult. However, there are other animal models found in nature where Hb α and β have been reported to be deleted, yet the animals still prove viable, with very interesting effects on the vasculature^{62,63}. These animals, Icefishes (family Channichthyidae), dominate the fish fauna of the Southern Ocean surrounding Antarctica⁶⁴. Icefishes are exceptional in that they are the only known vertebrate animals to completely lack the oxygen-carrier HbA in their blood⁶⁵. Loss of this key respiratory protein has ensued a phenotypic pale, translucent white blood and has earned these animals the common names of 'white-blooded fishes' and 'icefishes' (Figure 4)⁶⁶. Absent a gaseous transporter protein, oxygen is carried unassisted in solution of icefish blood and results in an oxygen-carrying capacity that is <10% of that exhibited by red-blooded notothenioid fishes⁶⁷. Given this

consequence and the fact that the HbA-null condition proves lethal in all other examples, both laboratory-based and natural, how has this condition persisted in icefishes then? Although more complex and beyond the scope of what we can present here (for an excellent review, see⁶⁸), there are several basic considerations that help to answer this question. First, icefishes possess cardiovascular systems with unusually enhanced features as compared to their red-blooded notothenioid relatives; large hearts, large diameter capillaries, and large blood volumes collectively enable icefishes to maintain a high-throughput circulatory design without excessive pressure development 69,70 . Combined with the abundantly high oxygen content of Antarctic waters and their relatively low metabolic rates, these cardiovascular traits permit icefishes to sufficiently oxygenate their tissues and support their aerobic mode of metabolism⁷¹. As one might expect however, the loss of expression of HbA has implications for the metabolism of NO. It was proposed that the loss of NO-oxygenase activity with genetic deletion of HbA may have led to subsequent elevation of NO levels that could explain many, if not all, of the unique cardiovascular and physiological traits that evolved in icefishes⁶⁸. Substantiating at least part of that hypothesis, Beers et al. (2010) established that NO concentrations in blood plasma appear to be greater in icefishes than in HbA-expressing species⁷². They also reported that the high NO levels in icefishes were not the result of greater synthesis but, rather, appeared to be due primarily to the loss of the degradative pathway for NO⁷².

Building upon the above work, Borley and collaborators subsequently conducted a study in which they induced severe anemia in *Notothenia coriiceps*, an HbA-expressing notothenioid with a normal hematocrit of 35–40%⁷³. Surgically implanted osmotic pumps were used to treat individuals with a powerful hemolytic agent that resulted in a drastic reduction in hematocrit (>90%) and HbA concentration (>70%). Levels of NO were significantly higher in anemic animals compared to the full HbA-expressing controls and were similar to the levels of NO reported for white-blooded icefishes^{72,73}. Although MEJs have not been identified in these fish, it would be surprising if these anatomical structures weren't present. The regulation of NO in the arteries of icefish could be one potential avenue for understanding how Hbα regulates NO delivery in arteries with an *in vivo* model capable of withstanding severe hemoglobin depletion.

Summary

This review describes a new paradigm of localized NO regulation by Hb α in MEJs of resistance arteries. This unique microdomain has the potential for pharmacological targeting and serves as an explanation for several different pathologies associated with Hb α deletion. Although some work has been done on this observation, there is still much to do and we look forward to extending this review in the future.

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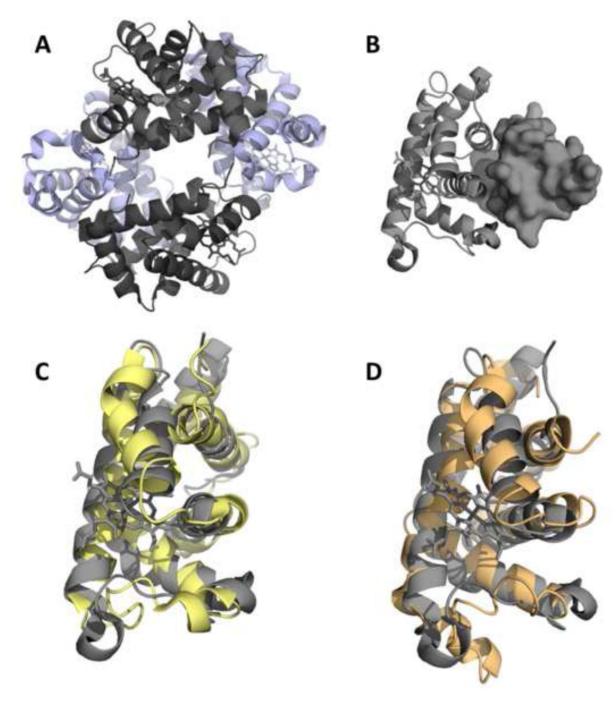


Figure 1. Structure of Hba

(A) HbA is a heterotetramer with two a subunits (gray) and two β subunits (purple). (B) Monomeric Hba (rendered in cartoon) is stabilized by AHSP (surface representation). The overall backbone fold of Hba bound to AHSP (C and D, gray) is similar to Hba (C, yellow) and Hb β (D, orange) of HbA.

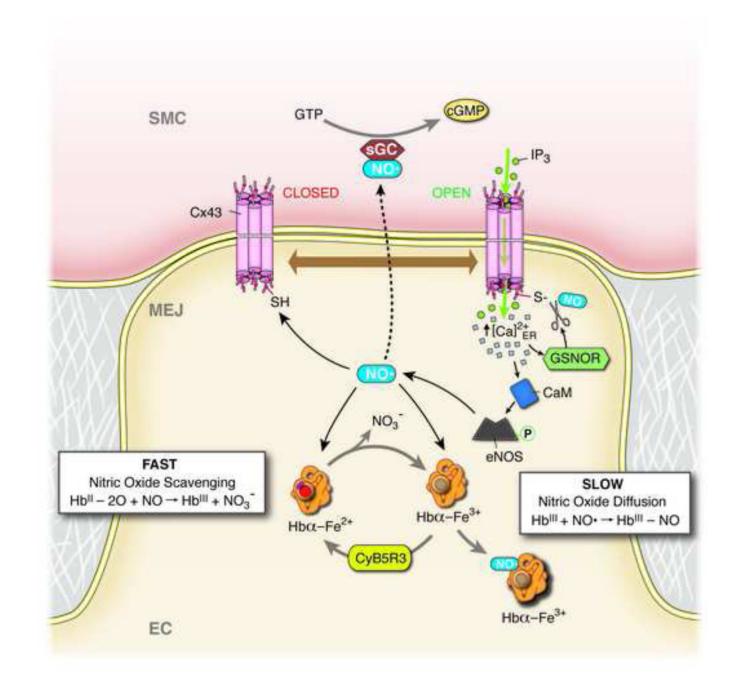


Figure 2. Schematic of NO regulation at the MEJ

Negative feedback following α_1 -adrenergic receptor induced vasoconstriction at the myoendothelial junction (MEJ) of an endothelial cell (EC). α_1 -adrenergic receptor agonists cause an increase in smooth muscle cell (VSMC) cytosolic inositol 1,4,5-trisphosphate (IP₃), which travels down its concentration gradient through open (i.e. S-nitrosylated) connexin 43 (Cx43) channels. IP₃ binds to localized IPs-receptor 1 on the endoplasmic reticulum (not shown), which causes localized calcium (Ca²⁺) release and an increase in cytosolic Ca²⁺ concentration at the MEJ. This increase in cytosolic Ca²⁺ facilitates activation of S-nitrosoglutathione reductase (GSNOR), which denitrosylates and closes Cx43 channels,

making them impermeable to IP₃. The increase in cytosolic Ca²⁺ also activates calmodulin (CaM) through binding-induced conformational changes. The Ca²⁺/CaM complex binds to endothelial nitric oxide synthase (eNOS), facilitating its phosphorylation, activation, and production of nitric oxide (NO). Some of this newly-produced NO will diffuse into VSMC, where it binds to and activates soluble guanylyl cyclase (sGC), facilitating the conversion of GTP to cGMP, which ultimately leads to a reduction in constriction. The newly-produced NO may also facilitate opening of Cx43 channels through S-nitrosylation. Finally, excess NO may be scavenged by free hemoglobin alpha (Hb α) at the MEJ through reaction with oxyhemoglobin alpha (fast), or NO may be chelated through reaction with methemoglobin alpha (slow). Cytochrome B5 reductase (CyB5R3) converts methemoglobin alpha to Hb α , which readily binds oxygen^{3,6,21–23}.

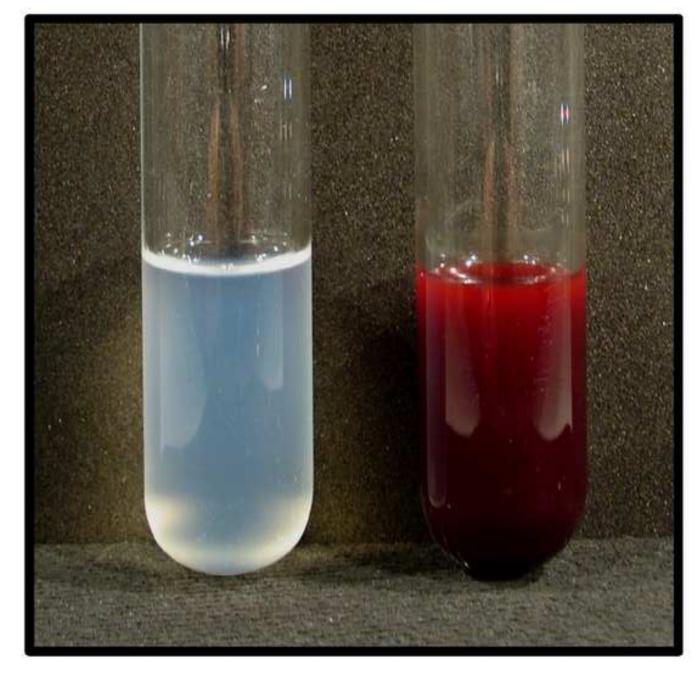
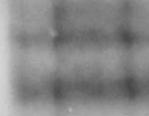


Figure 3. Expression of eNOS between resistance and conduit arteries is equal Representative western blot analysis of endothelial nitric oxide synthase (eNOS) protein level in third-order mesenteric arteries compared to carotid. Samples were equalized according to the amount of VE-cadherin, an endothelial cell marker.

mes carotid



eNOS

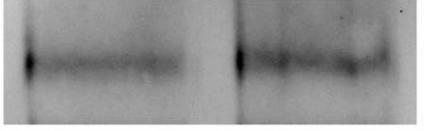




Figure 4. Blood from HbA and HbA null fish

Blood samples from two Antarctic notothenioid fishes illustrate a striking contrast in level of hemoglobin expression. The test tube on the right contains blood from an HbA-expressing species, *Notothenia coriiceps*, while the tube on the left depicts the completely HbA-null phenotype of the 'crocodile' icefish, Chaenocephalus aceratus.

Table 1

Chromosome 16 contains two identical coding sequences for hemoglobin alpha, hemoglobin alpha 2 and hemoglobin alpha 1 (shown here in reference to the zetaglobin gene, which is also located on chromosome 16). Deletions/non-functional mutations are depicted by the absence of a corresponding box. Genotype, red blood cell (RBC) indices, vascular phenotype with relation to blood pressure, and medical classification of the various alpha thalassemia's are also listed^{8,10,44,74}.

| Chromosome 16 (both copies shown) | | | Genotype | RBC Indices | Vascular Phenotype | Classification |
|-----------------------------------|--|-------|-------------|---|--|--|
| HBZ | m | HBA-1 | | | | |
| ====== a ====== a ====== a ====== | | aa/aa | Normal | Normal | Normal | |
| ====== a ====== a ====== a ====== | | | | | | |
| | ====== a = ======= a = | α | a-/aa | Normal | Moderate hypotension with potential compensation | Alpha thalassemia minima, also known as heterozygo sity fo alpha (+) thalassemia, silent carrier of alpha thalassemia, and alpha thalassemia-2 trait |
| ====== z or ======= z | ====== α = ======= α = ========= α = | | a-/a- or/aa | Minimal anemia, decreased MCV and MCH | Moderate hypotension | Alpha thalassemia minor, also known a alpha thalassemia-1 trait (Due to homozygosity for alpha (+) thalassem (α -/ α -) or heterozyg sity for alpha (0) thalassemia (/ $\alpha\alpha$) |
| 1 | 1-1 | | a-/ | Hemolytic anemia with the formation of P-chain tetramers | None documented | Hemoglobin H (HbH) disease |
| | | | -/ | Severe anemia due to formation of gamma-4 tetramers (hemoglobin Bart's) | Hydrops fetalis with increased cerebral blood flow | Hydrops fetalis syndrome with hemoglobin Bart's |

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