

## GEPHE SUMMARY

hemoglobin; HBB ( <a href="https://www.gephebase.org/search-criteria?/and+Gene">https://www.gephebase.org/search-criteria?/and+Gene</a> Gephebase="hemoglobin; HBB" #gephebase-summary-title)	Gephebase Gene	GP00000472	GepheID
Published	Entry Status	Martin	Main curator

## PHENOTYPIC CHANGE

Physiology ( <a href="https://www.gephebase.org/search-criteria?/and+Trait">https://www.gephebase.org/search-criteria?/and+Trait</a> Category="Physiology" #gephebase-summary-title)	Trait Category		
Hypoxia response ( <a href="https://www.gephebase.org/search-criteria?/and+Trait">https://www.gephebase.org/search-criteria?/and+Trait</a> ="Hypoxia response" #gephebase-summary-title)	Trait		
Lophonetta specularioides- low-altitude	Trait State in Taxon A		
Lophonetta specularioides- High-altitude	Trait State in Taxon B		
Taxon A	Ancestral State		
Intraspecific ( <a href="https://www.gephebase.org/search-criteria?/and+Taxonomic">https://www.gephebase.org/search-criteria?/and+Taxonomic</a> Status="Intraspecific" #gephebase-summary-title)	Taxonomic Status		
	Taxon A	Taxon B	
Lophonetta specularioides ( <a href="https://www.gephebase.org/search-criteria?/and+Taxon">https://www.gephebase.org/search-criteria?/and+Taxon</a> and Synonyms="Lophonetta specularioides" #gephebase-summary-title)	Latin Name	Lophonetta specularioides ( <a href="https://www.gephebase.org/search-criteria?/and+Taxon">https://www.gephebase.org/search-criteria?/and+Taxon</a> and Synonyms="Lophonetta specularioides" #gephebase-summary-title)	Latin Name
crested duck	Common Name	crested duck	Common Name
Anas specularioides; Lophonetta specularioides; crested duck; Lophonetta specularioides (King, 1828)	Synonyms	Anas specularioides; Lophonetta specularioides; crested duck; Lophonetta specularioides (King, 1828)	Synonyms
species	Rank	species	Rank
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Sauropsida; Sauria; Archelosauria; Archosauria; Dinosauria; Saurischia; Theropoda; Coelurosauria; Aves; Neognathae; Galloanserae; Anseriformes; Anatidae; Anatinae; Lophonetta	Lineage	cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa; Bilateria; Deuterostomia; Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Euteleostomi; Sarcopterygii; Dipnotetrapodomorpha; Tetrapoda; Amniota; Sauropsida; Sauria; Archelosauria; Archosauria; Dinosauria; Saurischia; Theropoda; Coelurosauria; Aves; Neognathae; Galloanserae; Anseriformes; Anatidae; Anatinae; Lophonetta	Lineage
Lophonetta () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75872">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75872</a> )	Parent	Lophonetta () - (Rank: genus) ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75872">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75872</a> )	Parent
75873 ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75873">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75873</a> )	NCBI Taxonomy ID	75873 ( <a href="https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75873">https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=75873</a> )	NCBI Taxonomy ID
No	is Taxon A an Intraspecies?	No	is Taxon B an Intraspecies?

## GENOTYPIC CHANGE

HBB	Generic Gene Name	P68871 ( <a href="http://www.uniprot.org/uniprot/P68871">http://www.uniprot.org/uniprot/P68871</a> )	UniProtKB Homo sapiens
ECYT6; CD113t-C; beta-globin	Synonyms	()	GenebankID or UniProtKB
9606.ENSP00000333994 ( <a href="http://string-db.org/newstring.cgi/show_network_section.pl?identifier=9606.ENSP00000333994">http://string-db.org/newstring.cgi/show_network_section.pl?identifier=9606.ENSP00000333994</a> )	String		
Belongs to the globin family.	Sequence Similarities		
GO:0046872 : metal ion binding ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0046872">https://www.ebi.ac.uk/QuickGO/term/GO:0046872</a> )	GO - Molecular Function		
GO:0020037 : heme binding ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0020037">https://www.ebi.ac.uk/QuickGO/term/GO:0020037</a> )			
GO:0005344 : oxygen carrier activity ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0005344">https://www.ebi.ac.uk/QuickGO/term/GO:0005344</a> )			
GO:0043177 : organic acid binding ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0043177">https://www.ebi.ac.uk/QuickGO/term/GO:0043177</a> )			
GO:0019825 : oxygen binding ( <a href="https://www.ebi.ac.uk/QuickGO/term/GO:0019825">https://www.ebi.ac.uk/QuickGO/term/GO:0019825</a> )			

GO:0031721 : hemoglobin alpha binding  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0031721)  
 GO:0030492 : hemoglobin binding (https://www.ebi.ac.uk/QuickGO/term/GO:0030492)  
 GO - Biological Process

GO:0006898 : receptor-mediated endocytosis  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0006898)  
 GO:0007596 : blood coagulation (https://www.ebi.ac.uk/QuickGO/term/GO:0007596)  
 GO:0008217 : regulation of blood pressure  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0008217)  
 GO:0042542 : response to hydrogen peroxide  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0042542)  
 GO:0043312 : neutrophil degranulation  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0043312)  
 GO:0015701 : bicarbonate transport (https://www.ebi.ac.uk/QuickGO/term/GO:0015701)  
 GO:0098869 : cellular oxidant detoxification  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0098869)  
 GO:0042744 : hydrogen peroxide catabolic process  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0042744)  
 GO:0015671 : oxygen transport (https://www.ebi.ac.uk/QuickGO/term/GO:0015671)  
 GO:0010942 : positive regulation of cell death  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0010942)  
 GO:0051291 : protein heterooligomerization  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0051291)  
 GO:0030185 : nitric oxide transport (https://www.ebi.ac.uk/QuickGO/term/GO:0030185)  
 GO:0070527 : platelet aggregation (https://www.ebi.ac.uk/QuickGO/term/GO:0070527)  
 GO:0045429 : positive regulation of nitric oxide biosynthetic process  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0045429)  
 GO:0050880 : regulation of blood vessel size  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0050880)  
 GO:0070293 : renal absorption (https://www.ebi.ac.uk/QuickGO/term/GO:0070293)

GO - Cellular Component

GO:0005829 : cytosol (https://www.ebi.ac.uk/QuickGO/term/GO:0005829)  
 GO:0070062 : extracellular exosome (https://www.ebi.ac.uk/QuickGO/term/GO:0070062)  
 GO:0005576 : extracellular region (https://www.ebi.ac.uk/QuickGO/term/GO:0005576)  
 GO:0005615 : extracellular space (https://www.ebi.ac.uk/QuickGO/term/GO:0005615)  
 GO:0072562 : blood microparticle (https://www.ebi.ac.uk/QuickGO/term/GO:0072562)  
 GO:0071682 : endocytic vesicle lumen  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0071682)  
 GO:0031838 : haptoglobin-hemoglobin complex  
 (https://www.ebi.ac.uk/QuickGO/term/GO:0031838)  
 GO:0005833 : hemoglobin complex (https://www.ebi.ac.uk/QuickGO/term/GO:0005833)  
 GO:1904813 : ficolin-1-rich granule lumen  
 (https://www.ebi.ac.uk/QuickGO/term/GO:1904813)  
 GO:1904724 : tertiary granule lumen (https://www.ebi.ac.uk/QuickGO/term/GO:1904724)

No (https://www.gephebase.org/search-criteria?/and+Presumptive Null="No"#gephebase-summary-title) Presumptive Null  
 Coding (https://www.gephebase.org/search-criteria?/and+Molecular Type="Coding"#gephebase-summary-title) Molecular Type  
 SNP (https://www.gephebase.org/search-criteria?/and+Aberration Type="SNP"#gephebase-summary-title) Aberration Type  
 Nonsynonymous SNP Coding Change  
 Asp94Glu Molecular Details of the Mutation  
 Candidate Gene (https://www.gephebase.org/search-criteria?/and+Experimental Evidence="Candidate Gene"#gephebase-summary-title) Experimental Evidence

	Taxon A	Taxon B	Position
Codon	-	-	-
Amino-acid	-	-	-

Convergent Evolution of Hemoglobin Function in High-Altitude Andean Waterfowl Involves Limited Parallelism at the Molecular Sequence Level. (2015)  
 (https://pubmed.ncbi.nlm.nih.gov/26637114) Main Reference

Natarajan C; Projecto-Garcia J; Moriyama H; Weber RE; Muñoz-Fuentes V; Green AJ; Kopuchian C; Tubaro PL; Alza L; Bulgarella M; Smith MM; Wilson RE; Fago A; McCracken KG; Storz JF Authors

Abstract  
 A fundamental question in evolutionary genetics concerns the extent to which adaptive phenotypic convergence is attributable to convergent or parallel changes at the molecular sequence level. Here we report a comparative analysis of hemoglobin (Hb) function in eight phylogenetically replicated pairs of high- and low-altitude waterfowl taxa to test for convergence in the oxygenation properties of Hb, and to assess the extent to which convergence in biochemical phenotype is attributable to repeated amino acid replacements. Functional experiments on native Hb variants and protein engineering experiments based on site-directed mutagenesis revealed the phenotypic effects of specific amino acid replacements that were responsible for convergent increases in Hb-O<sub>2</sub> affinity in multiple high-altitude taxa. In six of the eight taxon pairs, high-altitude taxa evolved derived increases in Hb-O<sub>2</sub> affinity that were caused by a combination of unique replacements, parallel replacements (involving identical-by-state variants with independent mutational origins in different lineages), and collateral replacements (involving shared, identical-by-descent variants derived via introgressive hybridization). In genome scans of nucleotide differentiation involving high- and low-altitude populations of three separate species,

function-altering amino acid polymorphisms in the globin genes emerged as highly significant outliers, providing independent evidence for adaptive divergence in Hb function. The experimental results demonstrate that convergent changes in protein function can occur through multiple historical paths, and can involve multiple possible mutations. Most cases of convergence in Hb function did not involve parallel substitutions and most parallel substitutions did not affect Hb-O<sub>2</sub> affinity, indicating that the repeatability of phenotypic evolution does not require parallelism at the molecular level.

Additional References

## RELATED GEPHE

Related Genes

1 (hemoglobin; HBA2) (<https://www.gephebase.org/search-criteria?/or+Taxon ID=^75873^/and+Trait=Hypoxia response/and+groupHaplotypes=true#gephebase-summary-title>)

Related Haplotypes

1 (<https://www.gephebase.org/search-criteria?/or+Gene Gephebase=^hemoglobin; HBB^/and+Taxon ID=^75873^/or+Gene Gephebase=^hemoglobin; HBB^/and+Taxon ID=^75873^#gephebase-summary-title>)

## EXTERNAL LINKS

## COMMENTS

Needs curation