

GO:0042542 : response to hydrogen peroxide
GO:0015701 : bicarbonate transport
GO:0098869 : cellular oxidant detoxification
GO:0042744 : hydrogen peroxide catabolic process
GO:0015671 : oxygen transport
GO:0010942 : positive regulation of cell death
GO:0051291 : protein heterooligomerization

GO - Cellular Component

GO:0005829 : cytosol
GO:0016020 : membrane
GO:0070062 : extracellular exosome
GO:0005576 : extracellular region
GO:0005615 : extracellular space
GO:0072562 : blood microparticle
GO:0071682 : endocytic vesicle lumen
GO:0022627 : cytosolic small ribosomal subunit
GO:0031838 : haptoglobin-hemoglobin complex
GO:0005833 : hemoglobin complex

Presumptive Null

No

Molecular Type

Coding

Aberration Type

SNP

SNP Coding Change

Nonsynonymous

Molecular Details of the Mutation

No more than 12 amino acid substitutions required for providing crocodile-like properties in engineered human Hb - effect of single amino acid changes not tested

Experimental Evidence

[Candidate Gene](#)

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

Main Reference

[Transplanting a unique allosteric effect from crocodile into human haemoglobin. \(1995\)](#)

Authors

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Abstract

Crocodiles are able to remain under water for more than one hour without surfacing to breathe and often kill their prey by drowning it. How do crocodiles stay under water for a long time? When they hold their breath, bicarbonate ions, the final product of respiration, accumulate and drastically reduce the oxygen affinity of haemoglobin, releasing a large fraction of haemoglobin-bound oxygen into the tissues. We have now located the bicarbonate-ion-binding site at the alpha 1 beta 2-subunit interface by making various human-crocodile chimaeric haemoglobins. Furthermore, we have been able to transplant the bicarbonate effect into human haemoglobin by replacing only a few residues, even though the amino-acid sequence identity between crocodile (*Crocodylus niloticus*) and human haemoglobins is only 68% for the alpha- and 51% for the beta-subunit. These results indicate that an entirely new function which enables species to adapt to a new environment could evolve in a protein by a relatively small number of amino-acid substitutions in key positions.

Additional References

RELATED GEPHE

Related Genes

4 (BHLHE41, EGLN1, EPAS1, PPAR-alpha)

Related Haplotypes

No matches found.

COMMENTS

