## GEPHE SUMMARY

Gephebase=^GH^#gephebase-summary-title) Published	Gephebase Gene Entry Status	GP00002037 Courtier	GephelD Main curator
HENOTYPIC CHANGE			
Morphology (https://www.gephebase.org/search-criteria?/and+Tra Category=^Morphology^#gephebase-summary-title)	Trait Category iit Trait		
Body size (dwarfism) (https://www.gephebase.org/search-criteria?/ (dwarfism)^#gephebase-summary-title)	'and+Trait=^Body size Trait State in Taxon A		
Bos taurus	Trait State in Taxon B		
Bos taurus - miniature Taxon A	Ancestral State		
Domesticated (https://www.gephebase.org/search-criteria?/and+T Status=^Domesticated^#gephebase-summary-title)	Taxonomic Status axonomic		
Taxon A	Latin Name	Taxon B	Latin Name
Bos taurus (https://www.gephebase.org/search-criteria?/and+Taxon and Syno taurus^#gephebase-summary-title)	nyms=^Bos	Bos taurus (https://www.gephebase.org/search-criteria?/and+Taxon an taurus^#gephebase-summary-title)	nd Synonyms=^Bos
cattle	Common Name Synonyms	cattle	Common Name Synonyms
Bos bovis; Bos primigenius taurus; cattle; bovine; cow; dairy cow; do cow; Bos taurus Linnaeus, 1758; Bos Tauurus	omestic cattle; domestic Rank	Bos bovis; Bos primigenius taurus; cattle; bovine; cow; dairy cow; Bos taurus Linnaeus, 1758; Bos Tauurus	cow; domestic cattle; domestic Rank
species	Lineage	species	Lineage
cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eumetazoa Chordata; Craniata; Vertebrata; Gnathostomata; Teleostomi; Eute Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; Theria; Laurasiatheria; Artiodactyla; Ruminantia; Pecora; Bovidae; Bovinae	leostomi; Sarcopterygii; Eutheria; Boreoeutheria;	cellular organisms; Eukaryota; Opisthokonta; Metazoa; Eur Chordata; Craniata; Vertebrata; Gnathostomata; Teleoston Dipnotetrapodomorpha; Tetrapoda; Amniota; Mammalia; T Laurasiatheria; Artiodactyla; Ruminantia; Pecora; Bovidae; E	ni; Euteleostomi; Sarcopterygii; Theria; Eutheria; Boreoeutheria;
Pag (aver gattle) (Pagly gamy)	9903 ) NCBI Taxonomy ID	Bos (oxen, cattle) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.	.cgi?id= 9903 ) NCBI Taxonomy ID
Bos (oxen, cattle) - (Rank: genus) (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=	r tebr raxementy ib		
(https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id= 9913 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=	9913 ) s Taxon A an Infraspecies?	9913 (https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.	.cgi?id= 9913 ) is Taxon B an Infraspecies?

P01241 (http://www.uniprot.org/uniprot/P01241)

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UniProtKB Homo sapiens

GenebankID or UniProtKB

## GENOTYPIC CHANGE

GO:0005148 : prolactin receptor binding

	Generic Gene Name		
GH1	Synonyms		
GH; GHN; GH-N; GHB5; IGHD2; hGH-N; IGHD1A; IGHD1B	Synonyms		
	String		
9606.ENSP00000312673			
(http://string-db.org/newstring_cgi/show_network_section.pl?identifier=			
9606.ENSP00000312673 )			
	Sequence Similarities		
Belongs to the somatotropin/prolactin family.			
(	GO - Molecular Function		
GO:0046872 : metal ion binding (https://www.ebi.ac.uk/QuickGO/	term/GO:0046872)		
GO:0008083 : growth factor activity (https://www.ebi.ac.uk/QuickGO/term/GO:0008083			
GO:0005179 : hormone activity (https://www.ebi.ac.uk/QuickGO/term/GO:0005179)			

(https://www.ebi.ac.uk/QuickGO/term/GO:0005148)	
GO:0005131 : growth hormone receptor binding	
(https://www.ebi.ac.uk/QuickGO/term/GO:0005131)	
GO - Biological Process	
GO:0043406 : positive regulation of MAP kinase activity	
(https://www.ebi.ac.uk/QuickGO/term/GO:0043406)	
GO:0050731 : positive regulation of peptidyl-tyrosine phosphorylation	
(https://www.ebi.ac.uk/QuickGO/term/GO:0050731)	
GO:0043568 : positive regulation of insulin-like growth factor receptor signaling pathway	
(https://www.ebi.ac.uk/QuickGO/term/GO:0043568)	
GO:0032355 : response to estradiol (https://www.ebi.ac.uk/QuickGO/term/GO:0032355)	
GO:0040018 : positive regulation of multicellular organism growth	
(https://www.ebi.ac.uk/QuickGO/term/GO:0040018)	
GO:0010828 : positive regulation of glucose transmembrane transport	
(https://www.ebi.ac.uk/QuickGO/term/GO:0010828)	
GO:0060396 : growth hormone receptor signaling pathway	
(https://www.ebi.ac.uk/QuickGO/term/GO:0060396)	
GO:0060397 : JAK-STAT cascade involved in growth hormone signaling pathway	
(https://www.ebi.ac.uk/QuickGO/term/GO:0060397)	
GO:0046427 : positive regulation of JAK-STAT cascade	
(https://www.ebi.ac.uk/QuickGO/term/GO:0046427)	
GO:0042531 : positive regulation of tyrosine phosphorylation of STAT protein	
(https://www.ebi.ac.uk/QuickGO/term/GO:0042531)	
GO:0014068 : positive regulation of phosphatidylinositol 3-kinase signaling	
(https://www.ebi.ac.uk/QuickGO/term/GO:0014068)	
GO:0048513 : animal organ development	
(https://www.ebi.ac.uk/QuickGO/term/GO:0048513)	
GO:0031667 : response to nutrient levels	
(https://www.ebi.ac.uk/QuickGO/term/GO:0031667)	
GO:0070977 : bone maturation (https://www.ebi.ac.uk/QuickGO/term/GO:0070977)	
GO:0010536 : positive regulation of activation of Janus kinase activity	
(https://www.ebi.ac.uk/QuickGO/term/GO:0010536)	
GO:0045927 : positive regulation of growth	
(https://www.ebi.ac.uk/QuickGO/term/GO:0045927)	
GO - Cellular Component	
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:0070195 : growth hormone receptor complex	
(https://www.ebi.ac.uk/QuickGO/term/GO:0070195)	
GO:0031904 : endosome lumen (https://www.ebi.ac.uk/QuickGO/term/GO:0031904)	
	Presumptive Null
Unknown (https://www.gephebase.org/search-criteria?/and+Presumptive Null=^Unknown^#gephebase-summary-title)	
	Molecular Type
Coding (https://www.gephebase.org/search-criteria?/and+Molecular Type=^Coding^#gephebase-summary-title)	
	Aberration Type
$SNP(https://www.gephebase.org/search-criteria?/and+Aberration Type=SNP^*gephebase-summary-title)$	
	SNP Coding Change
Nonsynonymous	
	Molecular Details of the Mutation
g.48768780C>T c.593C>T p.T198M	
	Experimental Evidence
Candidate Gene (https://www.gephebase.org/search-criteria?/and+Experimental Evidence=^Candidate Gene^#gephebase-summary-title)	

	Taxon A	Taxon B	Position
Codon	ACG	ATG	48768780
Amino-acid	Thr	Met	198

Main Reference

A miniature condition in Brahman cattle is associated with a single nucleotide mutation within the growth hormone gene. (2009) (https://pubmed.ncbi.nlm.nih.gov/19524387)

Authors

McCormack BL; Chase CC; Olson TA; Elsasser TH; Hammond AC; Welsh TH; Jiang H; Randel RD; Okamura CA; Lucy MC

Abstract Miniature Brahman cattle at the USDA ARS Subtropical Agriculture Research Station in Brooksville, FL have normal proportioned growth but are approximately 70% of mature height and weight when compared with Brahman cattle in the same herd. Pedigree analyses suggest that the condition is inherited through a recessive allele. The miniature Brahman cattle in the Brooksville herd have been used for studies of growth and reproduction, but the underlying causative mutation is unknown. Presumably, the miniature condition could arise from a mutation in the GH gene. The objective, therefore, was to clone the GH cDNA from Brooksville miniature Brahman cattle, compare its sequence to normal Brahman cattle, and test the biological activity of the native GH protein. Messenger RNA was isolated from the pituitary, and a cDNA for the protein coding region of the GH gene was amplified by reverse-transcription polymerase chain reaction (PCR) from each of 2 miniature Brahman bulls. The cDNA were cloned into plasmid vectors, and top and bottom strands were sequenced by automated DNA sequencing. The sequence of both cDNA clones derived from miniature cattle differed from Bos indicus GH (GenBank AF034386) at base number 641 because there was a cytosine (C) instead of a thymine (T). The C to T change encoded a mutation (threonine to methionine) at amino acid 200 (T200M mutation). The mutation was confirmed by sequencing of an additional 2 miniature cattle and comparing their sequence to 2 normal cattle. The threonine is located in the fourth alpha helix of GH and is 1 of 8 amino acids that participate in binding of GH to the GH receptor. Twelve miniature Brahman and 9 normal Brahman cattle were tested by using a restriction fragment length polymorphism analysis that employed the BsmBI restriction enzyme (specific for the mutated nucleotide). The 12 miniature Brahman cattle were homozygous for the mutation (-/-). Seven of the normal Brahman cattle were homozygous for the witatity of -/- cattle (n=4) had approximately 60% activ

## **RELATED GEPHE**

Related Genes 5 (aggrecan, LCORL, PLAG1, PRKG2, RNF11) (https://www.gephebase.org/search-criteria?/or+Taxon ID=^9913^/and+Trait=Body size/and+groupHaplotypes=true#gephebasesummary-title)

No matches found.

EXTERNAL LINKS

COMMENTS

https://omia.org/OMIA001821/9913/

Related Haplotypes