

Ghsr Cas9-CKO Strategy

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Project Overview

Project Name

Ghsr

Project type

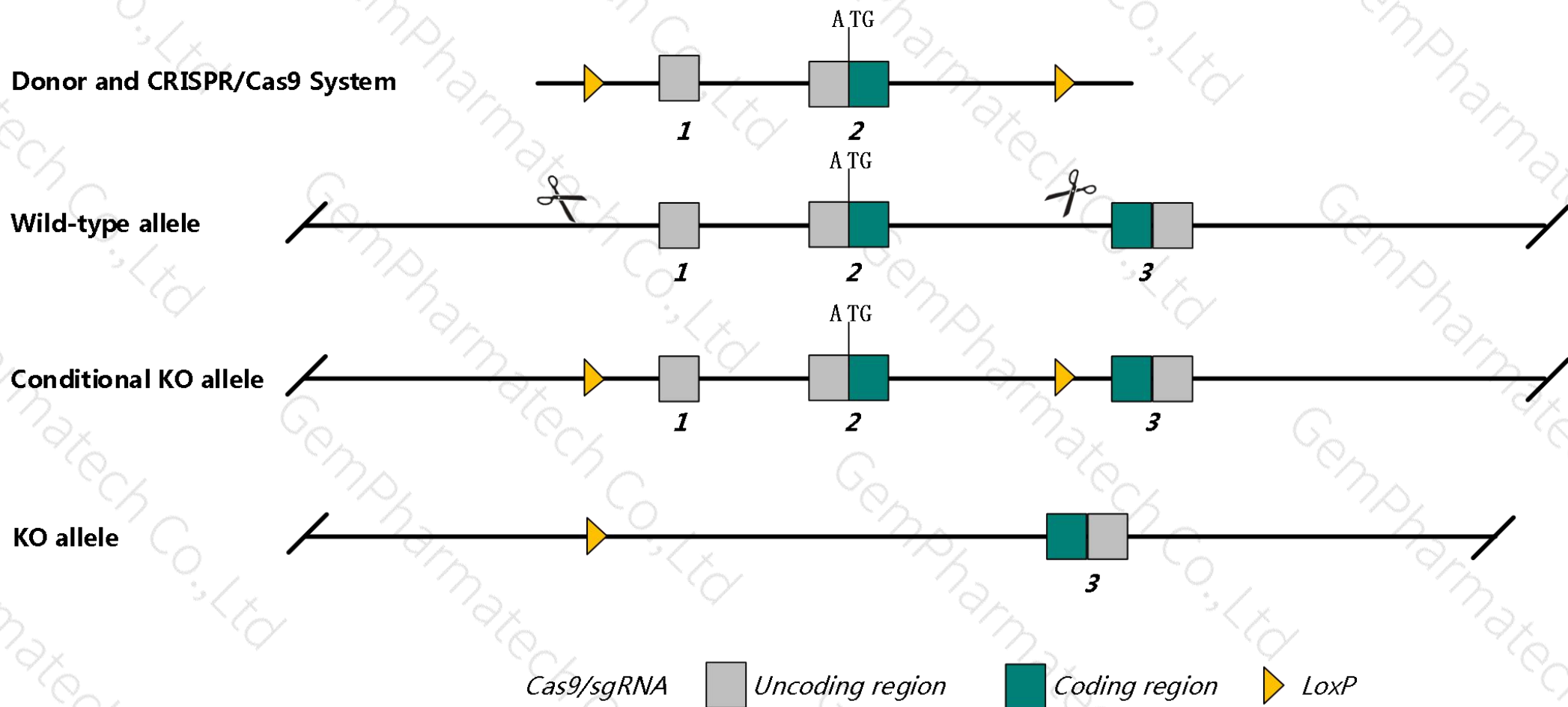
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Ghsr* gene. The schematic diagram is as follows:



- The *Ghsr* gene has 1 transcript. According to the structure of *Ghsr* gene, exon1-exon2 of *Ghsr-201* (ENSMUST00000057186.1) transcript is recommended as the knockout region. The region contains most of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ghsr* gene. The brief process is as follows: CRISPR/Cas9 system and Donor were microinjected into the fertilized eggs of C57BL/6JGpt mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6JGpt mice.
- The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

- According to the existing MGI data, Homeostasis is disrupted by inactivation of this gene, namely growth hormone release and appetite stimulation.
- The *Ghsr* gene is located on the Chr3. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

Gene information (NCBI)

Ghsr growth hormone secretagogue receptor [Mus musculus (house mouse)]

Gene ID: 208188, updated on 12-Mar-2019

Summary



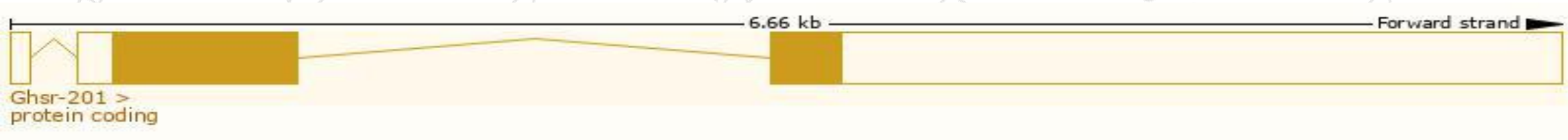
Official Symbol	Ghsr provided by MGI
Official Full Name	growth hormone secretagogue receptor provided by MGI
Primary source	MGI:MGI:2441906
See related	Ensembl:ENSMUSG00000051136
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	C530020I22Rik, GHRP, GHS-R, Ghsr1a
Expression	Low expression observed in reference dataset See more
Orthologs	human all

Transcript information (Ensembl)

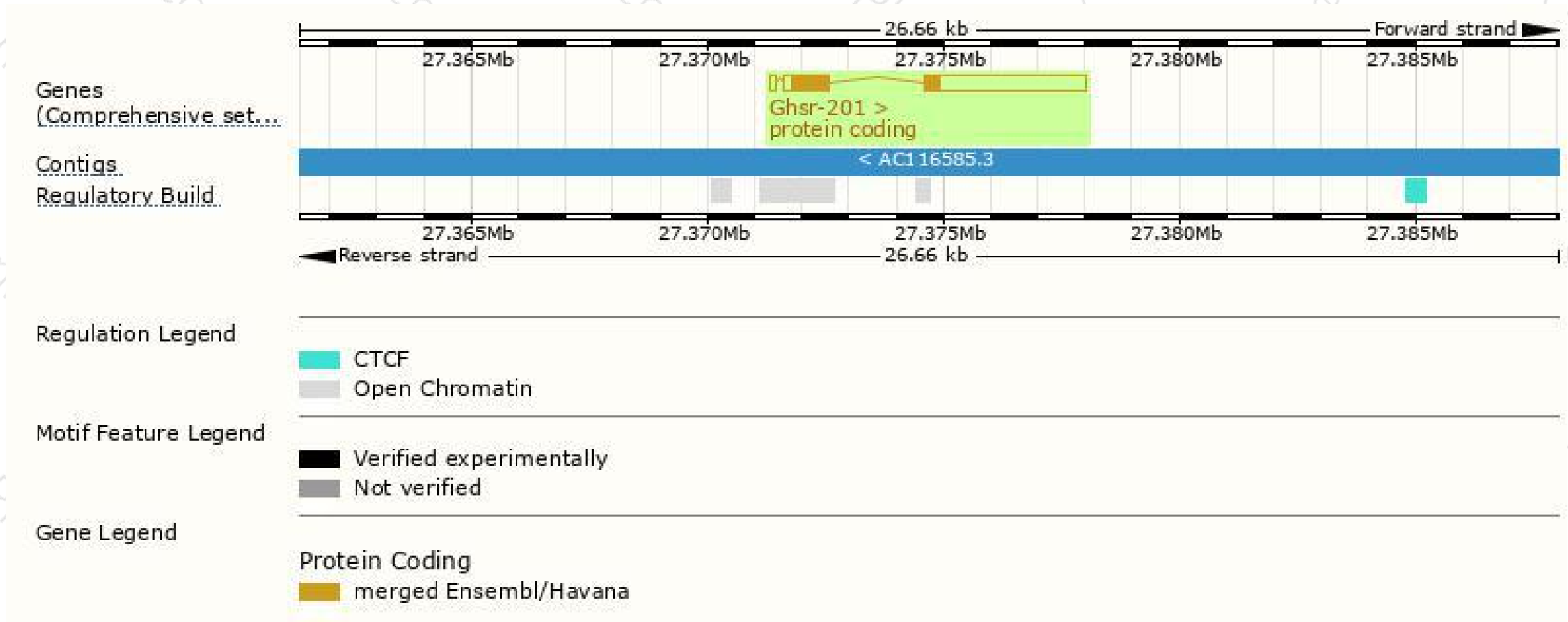
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ghsr-201	ENSMUST00000057186.1	4433	364aa	Protein coding	CCDS17273	Q0VBE5 Q99P50	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Ghsr-201* transcript, The transcription is shown below



Genomic location distribution



Protein domain

ENSMUSP00000061...

Transmembrane heli...

Low complexity (Seg)

Conserved Domains

hmmpanther

PTHR24243

Growth hormone secretagogue receptor

Superfamily domains

SSF81321

Prints domain

Growth hormone secretagogue receptor/motilin receptor

G protein-coupled receptor, rhodopsin-like

Pfam domain

G protein-coupled receptor, rhodopsin-like

PROSITE profiles

GPCR, rhodopsin-like, 7TM

PROSITE patterns

G protein-coupled receptor, rhodopsin-like

Gene3D

1.20.1070.10

All sequence SNPs/i...

Sequence variants (dbSNP and all other sources)

Variant Legend

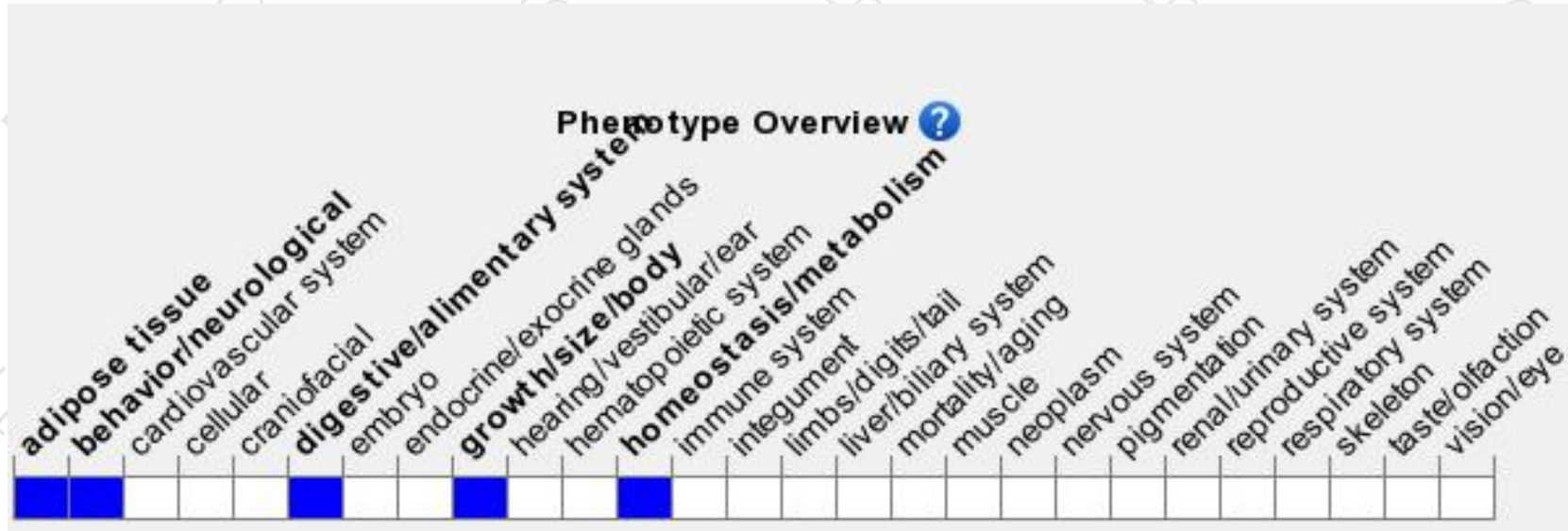
splice region variant

synonymous variant

Scale bar

0 40 80 120 160 200 240 280 320 364

Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

According to the existing MGI data, Homeostasis is disrupted by inactivation of this gene, namely growth hormone release and appetite stimulation.

If you have any questions, you are welcome to inquire.

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