

Hsp90aa1 Cas9-KO Strategy

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Project Overview



Project Name Hsp90aa1

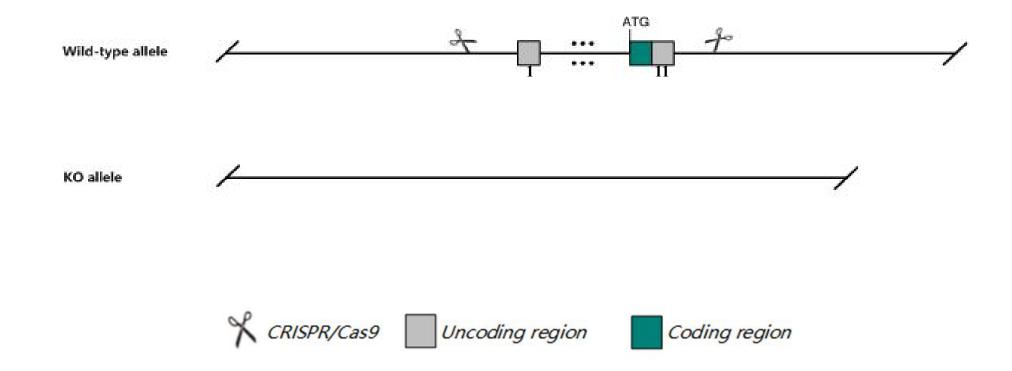
Project type Cas9-KO

Strain background C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Hsp90aa1* gene. The schematic diagram is as follows:



Technical routes



The *Hsp90aa1* gene has 8 transcripts. According to the structure of *Hsp90aa1* gene, exon1-exon11 of *Hsp90aa1-202* (ENSMUST00000094361.10) 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 *Hsp90aa1* gene. The brief process is as follows: CRISPR/Cas9 sys

Notice



According to the existing MGI data, Mice homozygous for a gene trap allele exhibit male sterility associated with arrested male meiosis and male germ cell apoptosis. Mice homozygous for a transgenic gene disruption exhibit male sterility and small testis.

The *Hsp90aa1* gene is located on the Chr12. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information NCBI



Hsp90aa1 heat shock protein 90, alpha (cytosolic), class A member 1 [Mus musculus (house mouse)]

Gene ID: 15519, updated on 7-Apr-2019

Summary



Official Symbol Hsp90aa1 provided by MGI

Official Full Name heat shock protein 90, alpha (cytosolic), class A member 1 provided by MGI

Primary source MGI:MGI:96250

See related Ensembl:ENSMUSG00000021270

Gene type protein coding
RefSeq status PROVISIONAL
Organism Mus musculus

Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;

Muroidea; Muridae; Murinae; Mus; Mus

Also known as 86kDa, 89kDa, AL024080, AL024147, Hsp86-1, Hsp89, Hsp90, Hspca, hsp4

Expression Broad expression in placenta adult (RPKM 235.9), CNS E11.5 (RPKM 213.3) and 19 other tissuesSee more

Orthologs <u>human</u> all

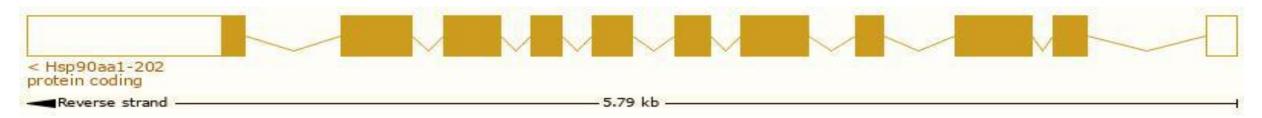
Transcript information Ensembl



The gene has 8 transcripts, all transcripts are shown below:

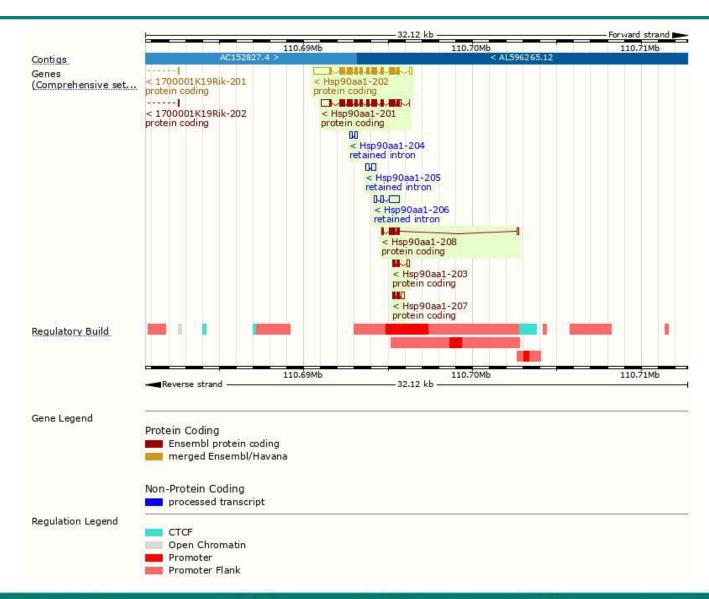
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Hsp90aa1-202	ENSMUST00000094361.10	3281	733aa	Protein coding	CCDS26172	P07901 Q80Y52	TSL:1 GENCODE basic APPRIS P1
Hsp90aa1-201	ENSMUST00000021698.12	2852	<u>733aa</u>	Protein coding	CCDS26172	P07901 Q80Y52	TSL:5 GENCODE basic APPRIS P1
Hsp90aa1-208	ENSMUST00000155242.7	711	201aa	Protein coding	20	B7ZC50	CDS 3' incomplete TSL:3
Hsp90aa1-207	ENSMUST00000149189.1	481	98aa	Protein coding	29	A2A6A2	CDS 3' incomplete TSL:2
Hsp90aa1-203	ENSMUST00000124156.7	427	103aa	Protein coding	-	B7ZC49	CDS 3' incomplete TSL:2
Hsp90aa1-206	ENSMUST00000145255.1	924	No protein	Retained intron	-	#3	TSL:1
Hsp90aa1-205	ENSMUST00000134967.1	487	No protein	Retained intron	20	49	TSL:2
Hsp90aa1-204	ENSMUST00000129005.1	325	No protein	Retained intron	29	20	TSL:1

The strategy is based on the design of *Hsp90aa1-202* transcript, The transcription is shown below



Genomic location distribution





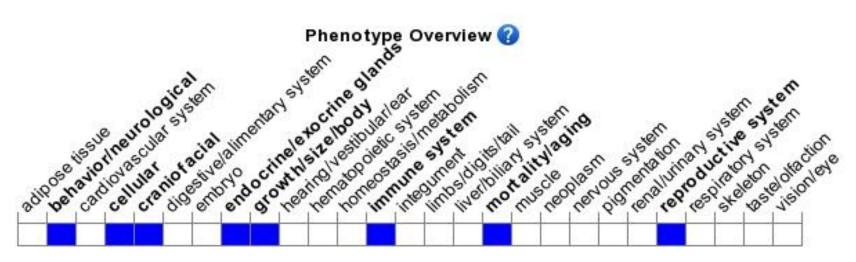
Protein domain





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, Mice homozygous for a gene trap allele exhibit male sterility associated with arrested male meiosis and male germ cell apoptosis. Mice homozygous for a transgenic gene disruption exhibit male sterility and small testis.



If you have any questions, you are welcome to inquire. Tel: 400-9660890





