

Adrb3 Cas9-KO Strategy

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Design Date: 2019-8-5

Project Overview



Project Name

Adrb3

Project type

Cas9-KO

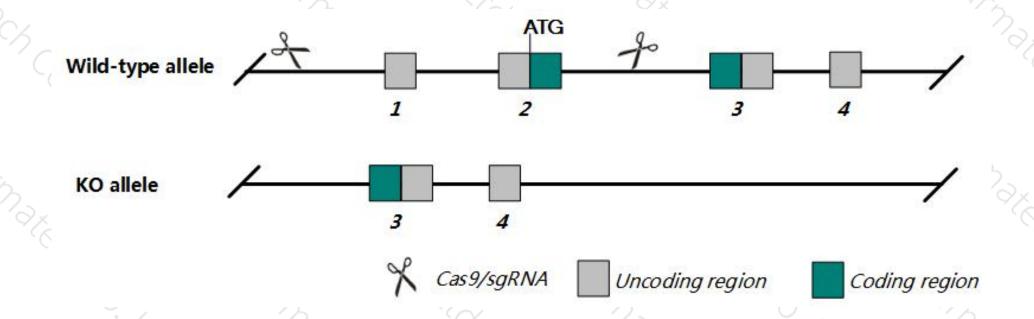
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Adrb3* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Adrb3* gene has 5 transcripts. According to the structure of *Adrb3* gene, exon1-exon2 of *Adrb3-201*(ENSMUST00000081438.9) 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 *Adrb3* gene. The brief process is as follows: gRNA was transcribed in vitro.Cas9 and gRNA 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.

Notice



- ➤ According to the existing MGI data, Homozygotes for targeted null mutations develop greater adiposity, especially on a high-fat diet, and are unresponsive to the beta3-adrenergic receptor agonist, CL316,243.
- \triangleright Partial sequence of Gm45470 gene will be deleted together in this strategy.
- The floxed region is near to the N-terminal of *Got111* gene, this strategy may influence the regulatory function of the N-terminal of *Got111* gene.
- The *Adrb3* gene is located on the Chr8. 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)



Adrb3 adrenergic receptor, beta 3 [Mus musculus (house mouse)]

Gene ID: 11556, updated on 25-Jun-2019

Summary

△ ?

Official Symbol Adrb3 provided by MGI

Official Full Name adrenergic receptor, beta 3 provided by MGI

Primary source MGI:MGI:87939

See related Ensembl: ENSMUSG00000031489

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 Adrb-3; beta 3-AR

Expression Biased expression in mammary gland adult (RPKM 215.3), subcutaneous fat pad adult (RPKM 163.4) and 4 other tissues See more

Orthologs human all

Genomic context

☆ ?

Location: 8 A2; 8 15.94 cM

See Adrb3 in Genome Data Viewer

Exon count: 6

Annotation release	Status	Assembly	Chr	Location	
106	current	GRCm38.p4 (GCF_000001635.24)	8	NC_000074.6 (2722577627230807, complement)	
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	8	NC_000074.5 (2833624828340060, complement)	

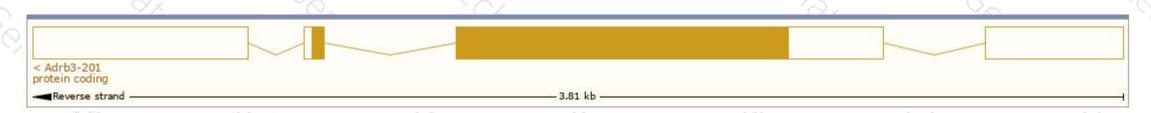
Transcript information (Ensembl)



The gene has 5 transcripts, all the transcript are shown below:

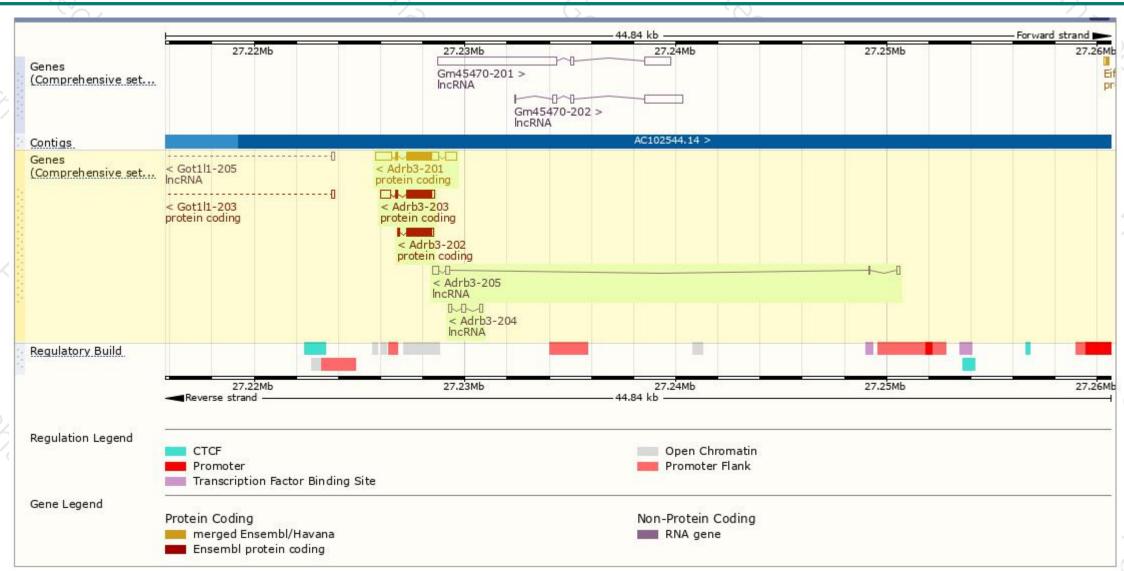
Name 🌲	Transcript ID A	bp 🌲	Protein #	Biotype 🍦	CCDS .	UniProt 🍦	Flags	4
Adrb3-201	ENSMUST00000081438.9	2795	400aa	Protein coding	CCDS22213@	P25962@ Q3UP63@	TSL:1 GENCODE ba	sic APPRIS P2
Adrb3-202	ENSMUST00000117565.1	1336	404aa	Protein coding		P25962⊌	TSL:1 GENCODE bas	ic APPRIS ALT2
Adrb3-203	ENSMUST00000121838.1	1864	400aa	Protein coding	CCDS22213 ₽	P25962@Q3UP63@	TSL:1 GENCODE ba	asic APPRIS P2
Adrb3-204	ENSMUST00000209447.1	450	No protein	IncRNA	i i	-	TSL:3	
Adrb3-205	ENSMUST00000211346.1	669	No protein	IncRNA	:=	-	TSL:3	

The strategy is based on the design of Adrb3-201 transcript, The transcription is shown below



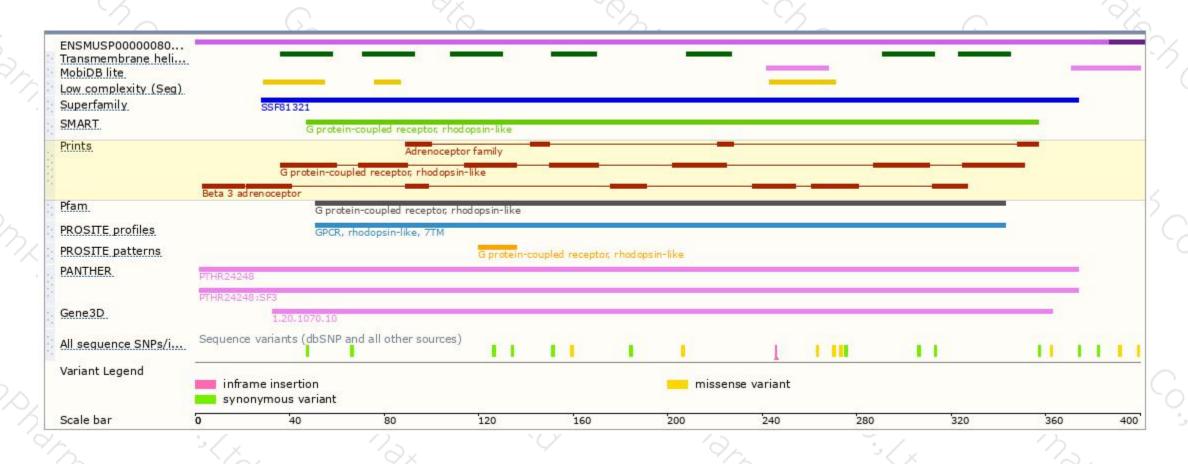
Genomic location distribution





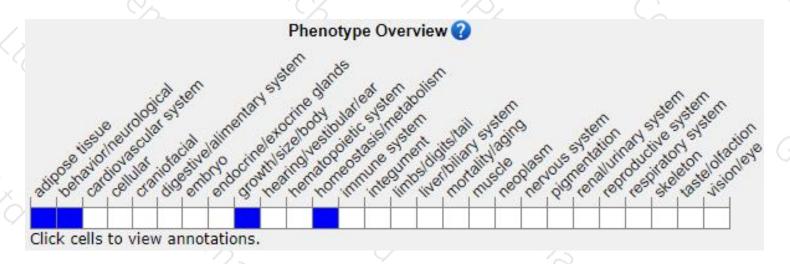
Protein domain





Mouse phenotype description(MGI)





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

Homozygotes for targeted null mutations develop greater adiposity, especially on a high-fat diet, and are unresponsive to the beta3-adrenergic receptor agonist, CL316,243.



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





