

Ube3a Cas9-KO Strategy

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Project Overview



Project Name

Ube3a

Project type

Cas9-KO

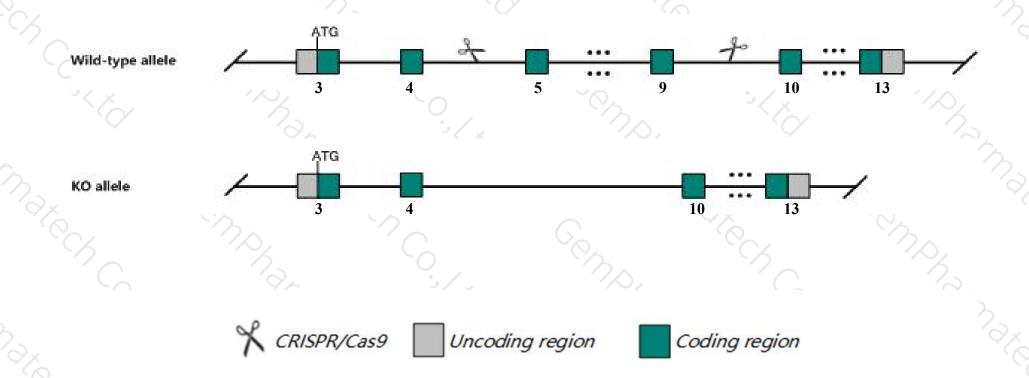
Strain background

C57BL/6JGpt

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Ube3a* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Ube3a* gene has 18 transcripts. According to the structure of *Ube3a* gene, exon5-exon9 of *Ube3a-203* (ENSMUST00000200758.3) transcript is recommended as the knockout region. The region contains 2053bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Ube3a* gene. The brief process is as follows: CRISPR/Cas9 system

Notice



- ➤ According to the existing MGI data, Mice with maternally inherited targeted null mutations exhibit reduced brain weight, impaired motor function, inducible seizures, learning deficits, abnormal hippocampal electroencephalographic recordings, and severely impaired long-term potentiation.
- > The *Ube3a* gene is located on the Chr7. 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)



Ube3a ubiquitin protein ligase E3A [Mus musculus (house mouse)]

Gene ID: 22215, updated on 9-Apr-2019

Summary

☆ ?

Official Symbol Ube3a provided by MGI

Official Full Name ubiquitin protein ligase E3A provided by MGI

Primary source MGI:MGI:105098

See related Ensembl: ENSMUSG00000025326

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 4732496B02, 5830462N02Rik, A130086L21Rik, Hpve6a

Expression Ubiquitous expression in adrenal adult (RPKM 21.0), ovary adult (RPKM 10.5) and 28 other tissuesSee more

Orthologs <u>human</u> all

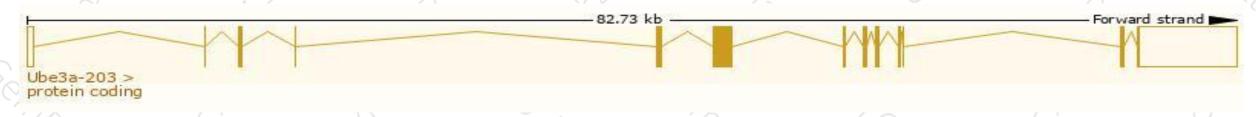
Transcript information (Ensembl)



The gene has 18 transcripts, all transcripts are shown below:

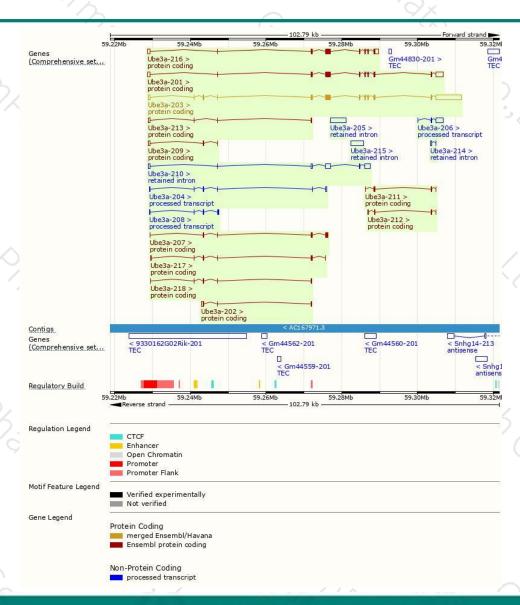
Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ube3a-203	ENSMUST00000200758.3	9856	870aa	Protein coding	CCDS39973@	008759₽	TSL:1 GENCODE basic APPRIS P3
Ube3a-201	ENSMUST00000107537.4	4911	849aa	Protein coding	CCDS80733@	<u>O08759</u> ₽	TSL:1 GENCODE basic APPRIS ALT2
Ube3a-216	ENSMUST00000202945.3	3889	762aa	Protein coding	=======================================	<u>O08759</u> ₽	TSL:1 GENCODE basic
Ube3a-207	ENSMUST00000201409.3	1119	313aa	Protein coding	8-	A0A0J9YUK0€	CDS 3' incomplete TSL:3
Ube3a-217	ENSMUST00000207686.1	797	<u>126aa</u>	Protein coding	2	A0A140LHS2@	CDS 3' incomplete TSL:5
Ube3a-213	ENSMUST00000202440.3	729	<u>49aa</u>	Protein coding	85	A0A0J9YTX6₽	CDS 3' incomplete TSL:5
Ube3a-202	ENSMUST00000200709.1	664	<u>40aa</u>	Protein coding	2	A0A0J9YUF2@	CDS 3' incomplete TSL:3
Jbe3a-209	ENSMUST00000201794.3	616	<u>15aa</u>	Protein coding	89	A0A0J9YV26 €	CDS 3' incomplete TSL:3
Ube3a-211	ENSMUST00000202247.3	579	<u>192aa</u>	Protein coding	2	A0A0J9YUY4€	CDS 5' incomplete TSL:3
Ube3a-212	ENSMUST00000202288.1	442	143aa	Protein coding	89	A0A0J9YVG1@	CDS 5' incomplete TSL:5
Ube3a-218	ENSMUST00000208313.1	441	78aa	Protein coding	22	A0A140LI50@	CDS 3' incomplete TSL:5
Ube3a-205	ENSMUST00000200949.1	4254	No protein	Retained intron	89	2-1	TSL:NA
Ube3a-210	ENSMUST00000202207.3	3586	No protein	Retained intron	2	120	TSL:1
Ube3a-215	ENSMUST00000202776.1	3348	No protein	Retained intron	19	5-1	TSL:NA
Ube3a-214	ENSMUST00000202685.1	436	No protein	Retained intron	2	120	TSL:2
Jbe3a-206	ENSMUST00000201162.1	2129	No protein	IncRNA	19	5-1	TSL:1
Ube3a-204	ENSMUST00000200781.3	742	No protein	IncRNA	2	120	TSL:5
Ube3a-208	ENSMUST00000201480.1	549	No protein	IncRNA		-	TSL:5

The strategy is based on the design of *Ube3a-203* transcript, The transcription is shown below



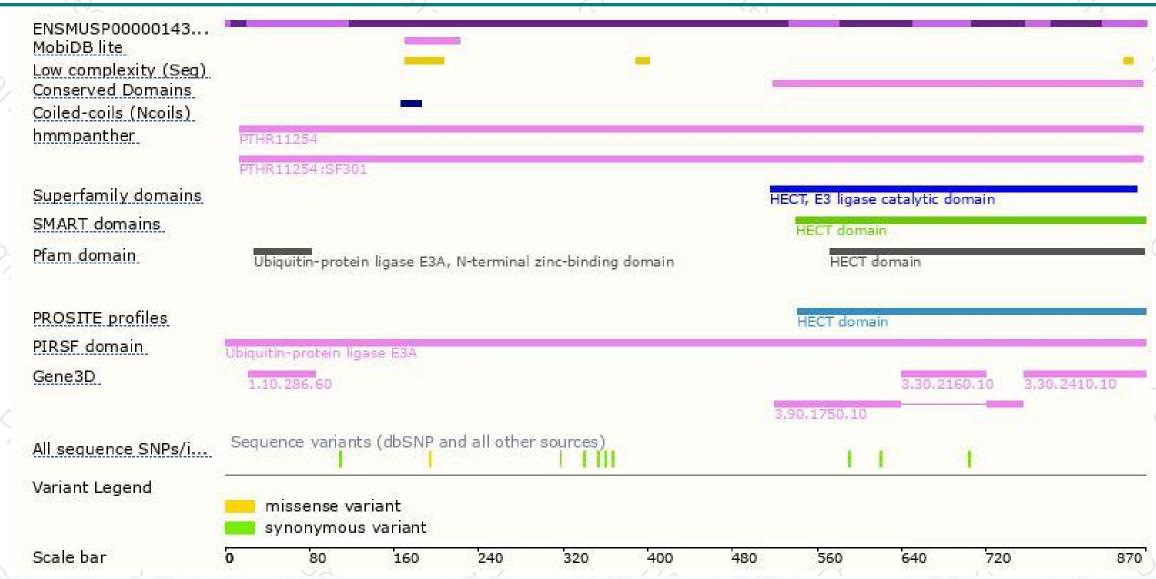
Genomic location distribution





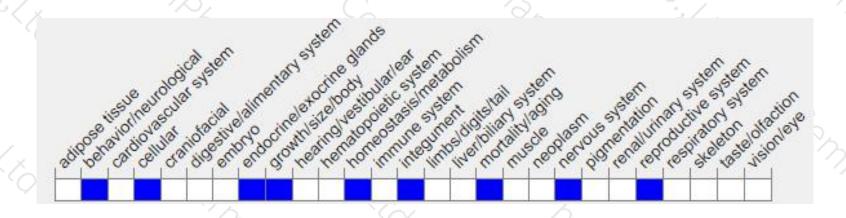
Protein domain





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 with maternally inherited targeted null mutations exhibit reduced brain weight, impaired motor function, inducible seizures, learning deficits, abnormal hippocampal electroencephalographic recordings, and severely impaired long-term potentiation.



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





