

Odc1 Cas9-KO Strategy RAMPHAMAKON CO.

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Project Overview



Project Name Odc1

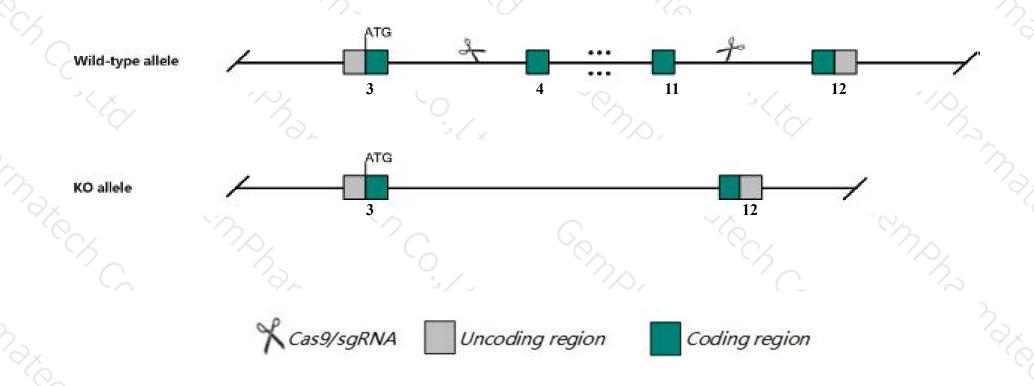
Project type Cas9-KO

Strain background C57BL/6J

Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Odc1* gene. The schematic diagram is as follows:



Technical routes



- ➤ The *Odc1* gene has 8 transcripts. According to the structure of *Odc1* gene, exon4-exon11 of *Odc1-201*(ENSMUST00000171737.2) transcript is recommended as the knockout region. The region contains 1139bp coding sequence Knock out the region will result in disruption of protein function.
- ➤ In this project we use CRISPR/Cas9 technology to modify *Odc1* gene. The brief process is as follows: sgRNA was transcribed in vitro.Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J 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/6J mice.

Notice



- > According to the existing MGI data, Homozygous null embryos die prior to gastrulation.
- ➤ The *Odc1* 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)



Odc1 ornithine decarboxylase, structural 1 [Mus musculus (house mouse)]

Gene ID: 18263, updated on 2-Apr-2019

Summary

☆ ?

Official Symbol Odc1 provided by MGI

Official Full Name ornithine decarboxylase, structural 1 provided by MGI

Primary source MGI:MGI:97402

See related Ensembl: ENSMUSG00000011179

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 ODC

Expression Broad expression in liver E14 (RPKM 226.2), liver E14.5 (RPKM 221.0) and 28 other tissuesSee more

Orthologs <u>human all</u>

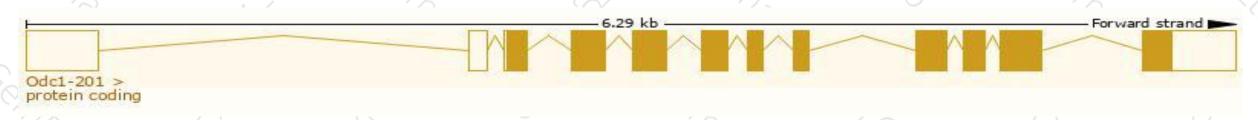
Transcript information (Ensembl)



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

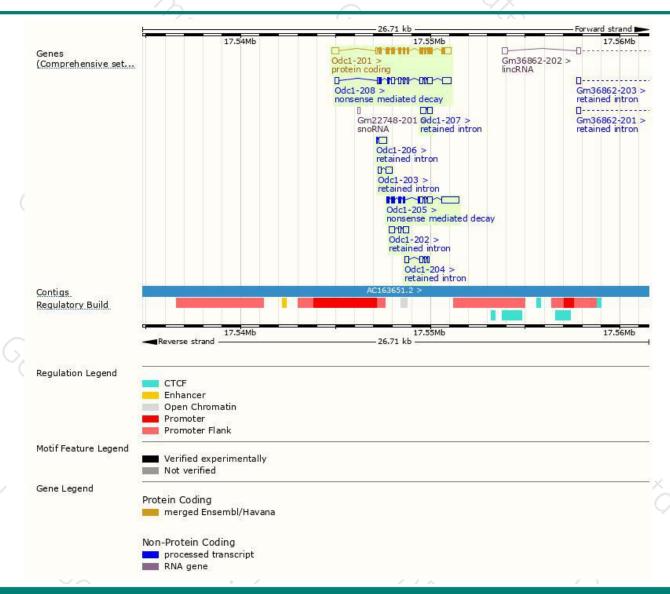
)		
Transcript ID	bp	Protein	Biotype	ccds	UniProt	Flags
ENSMUST00000171737.2	2208	<u>461aa</u>	Protein coding	CCDS25829	P00860	TSL:1 GENCODE basic APPRIS P1
ENSMUST00000221613.1	1982	<u>197aa</u>	Nonsense mediated decay	-	A0A1Y7VJC5	CDS 5' incomplete TSL:5
ENSMUST00000222617.1	1884	<u>42aa</u>	Nonsense mediated decay	ų.	A0A1Y7VMF9	TSL:5
ENSMUST00000220849.1	764	No protein	Retained intron	2	728	TSL:3
ENSMUST00000221354.1	561	No protein	Retained intron		1871	TSL:2
ENSMUST00000222250.1	538	No protein	Retained intron	-		TSL:2
ENSMUST00000220947.1	504	No protein	Retained intron	ū.	12)	TSL:2
ENSMUST00000221701.1	472	No protein	Retained intron	2	758	TSL:2
	ENSMUST00000171737.2 ENSMUST00000221613.1 ENSMUST00000222617.1 ENSMUST00000220849.1 ENSMUST00000221354.1 ENSMUST0000022250.1 ENSMUST00000220947.1	ENSMUST00000171737.2 2208 ENSMUST00000221613.1 1982 ENSMUST00000222617.1 1884 ENSMUST00000220849.1 764 ENSMUST00000221354.1 561 ENSMUST0000022250.1 538 ENSMUST00000220947.1 504	ENSMUST00000171737.2 2208 461aa ENSMUST000000221613.1 1982 197aa ENSMUST000000222617.1 1884 42aa ENSMUST000000220849.1 764 No protein ENSMUST00000221354.1 561 No protein ENSMUST00000222250.1 538 No protein ENSMUST00000220947.1 504 No protein	ENSMUST00000171737.2 2208 461aa Protein coding ENSMUST00000221613.1 1982 197aa Nonsense mediated decay ENSMUST00000222617.1 1884 42aa Nonsense mediated decay ENSMUST00000220849.1 764 No protein Retained intron ENSMUST00000221354.1 561 No protein Retained intron ENSMUST00000222250.1 538 No protein Retained intron ENSMUST00000220947.1 504 No protein Retained intron	ENSMUST00000171737.2 2208 461aa Protein coding CCDS25829 ENSMUST00000221613.1 1982 197aa Nonsense mediated decay - ENSMUST00000222617.1 1884 42aa Nonsense mediated decay - ENSMUST00000220849.1 764 No protein Retained intron - ENSMUST00000221354.1 561 No protein Retained intron - ENSMUST00000222250.1 538 No protein Retained intron - ENSMUST00000220947.1 504 No protein Retained intron -	ENSMUST00000171737.2 2208 461aa Protein coding CCDS25829 P00860 ENSMUST00000221613.1 1982 197aa Nonsense mediated decay - A0A1Y7VJC5 ENSMUST00000222617.1 1884 42aa Nonsense mediated decay - A0A1Y7VMF9 ENSMUST00000220849.1 764 No protein Retained intron - - ENSMUST00000221354.1 561 No protein Retained intron - - ENSMUST00000222250.1 538 No protein Retained intron - - ENSMUST000000220947.1 504 No protein Retained intron - -

The strategy is based on the design of *Odc1-201* transcript, The transcription is shown below



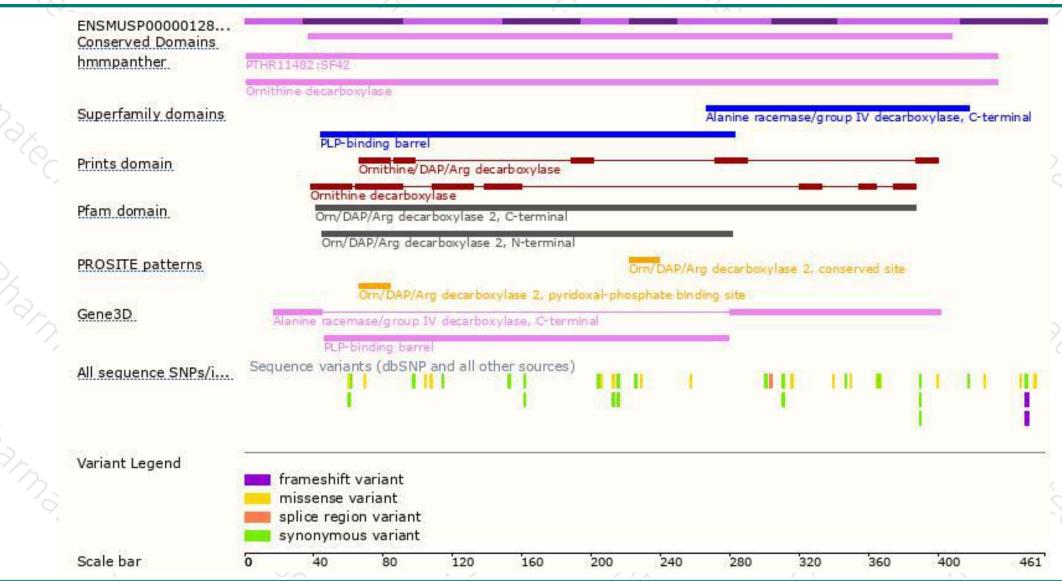
Genomic location distribution





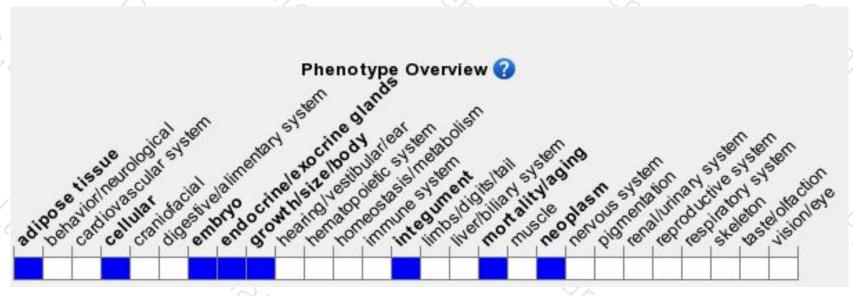
Protein domain





Mouse phenotype description(MGI)





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

According to the existing MGI data, Homozygous null embryos die prior to gastrulation.



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

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