

Grin2d Cas9-KO Strategy

Designer: Reviewer:

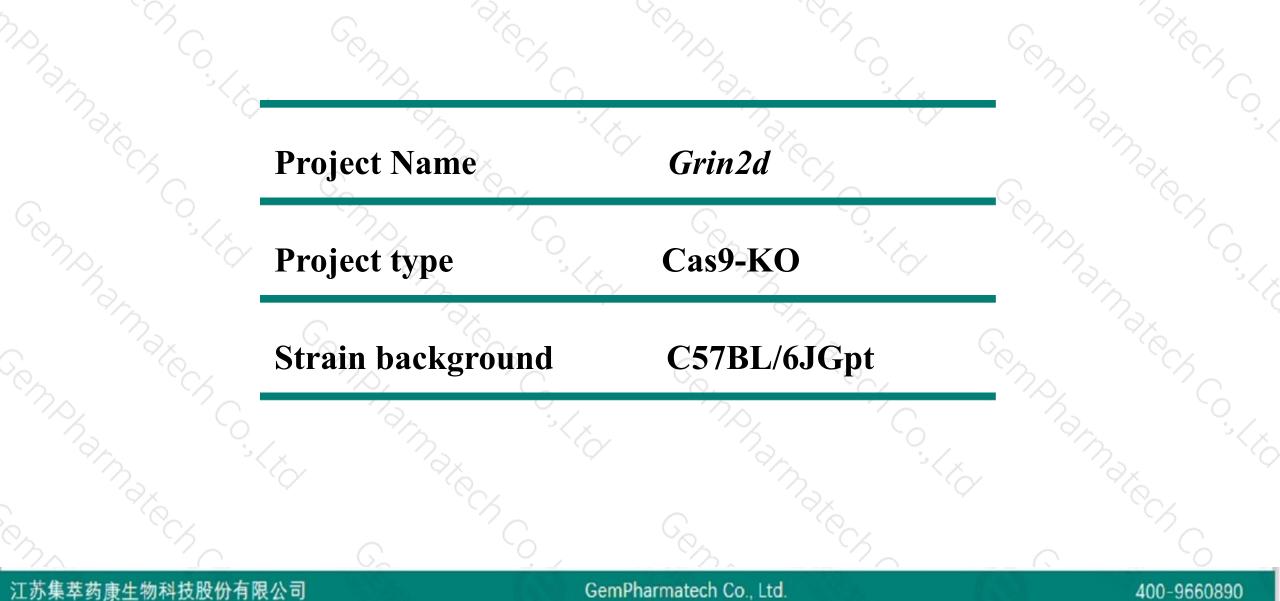
Design Date:

Bingxuan Li Ruirui Zhang

2019-10-31

Project Overview

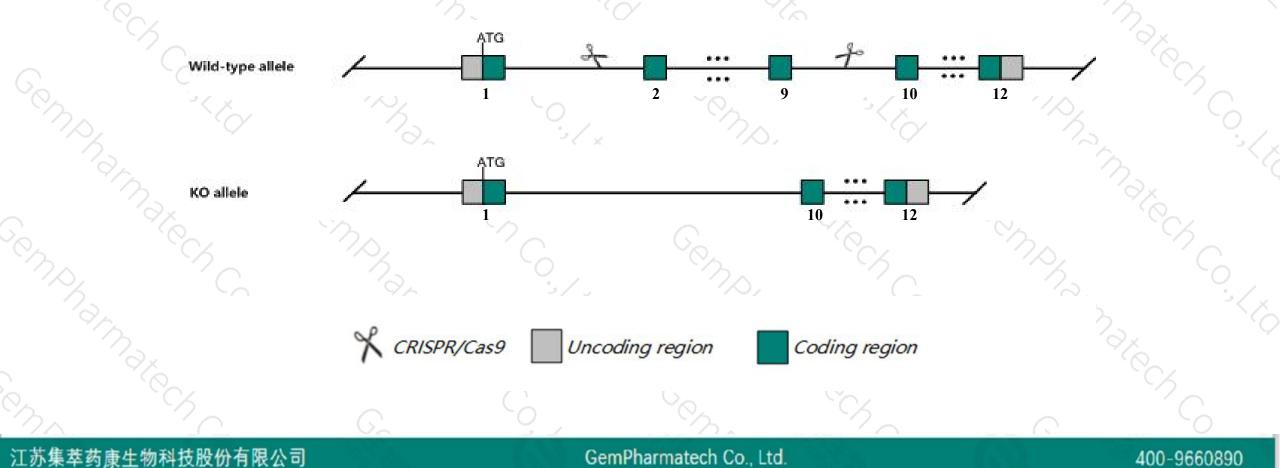




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Grin2d gene. The schematic diagram is as follows:





- The Grin2d gene has 3 transcripts. According to the structure of Grin2d gene, exon2-exon9 of Grin2d-201 (ENSMUST0000002848.9) transcript is recommended as the knockout region. The region contains 1787bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Grin2d gene. The brief process is as follows: CRISPR/Cas9 system

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- According to the existing MGI data, Homozygotes for a targeted null mutation exhibit reduced spontaneous activity and an elevated auditory brainstem response threshold.
- The Grin2d 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.
- > The *Grin2d-202* transcript has not been affected.
- 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.

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Notice

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Gene information (NCBI)



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Grin2d glutamate receptor, ionotropic, NMDA2D (epsilon 4) [Mus musculus (house mouse)]

Gene ID: 14814, updated on 19-Mar-2019

Summary

Official Symbol	Grin2d provided by MGI							
Official Full Name	glutamate receptor, ionotropic, NMDA2D (epsilon 4) provided by MGI							
Primary source	MGI:MGI:95823							
See related	Ensembl:ENSMUSG0000002771							
Gene type	protein coding							
RefSeq status	VALIDATED							
Organism	Mus musculus							
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Ro								
	Muroidea; Muridae; Murinae; Mus; Mus							
Also known as	GluN2D, NMDAR2D, NR2D							
Expression	Broad expression in testis adult (RPKM 12.8), CNS E18 (RPKM 5.9) and 19 other tissuesSee more							
Orthologs	human all							

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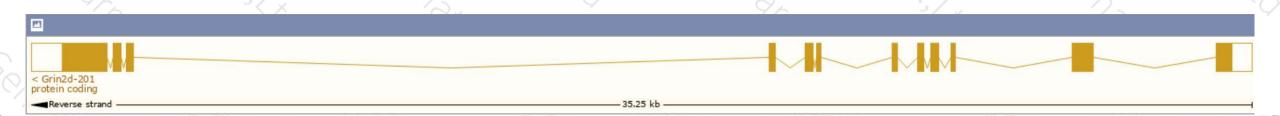
Transcript information (Ensembl)



The gene has 3 transcripts, all transcripts are shown below:

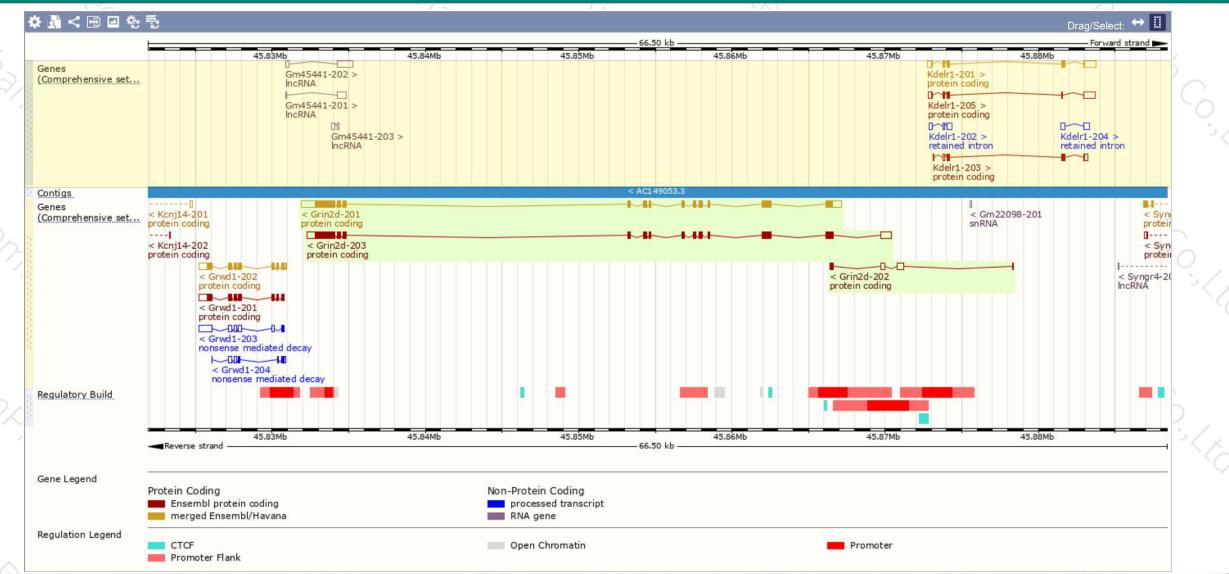
Show/hide columns (1 hidden)									
Name 🍦	Transcript ID 🔹	bp 🖕	Protein 🖕	Biotype 🍦	CCDS 🍦	UniProt 💧	Flags		
Grin2d-203	ENSMUST00000211713.1	5244	<u>1323aa</u>	Protein coding	<u>CCDS21267</u> &	<u>Q03391</u> &	TSL:5 GENCODE basic APPRIS P1		
Grin2d-202	ENSMUST00000211250.1	1009	<u>63aa</u>	Protein coding	(1)	<u>A0A1B0GRF9</u> &	CDS 3' incomplete TSL:5		
Grin2d-201	ENSMUST0000002848.9	5431	<u>1323aa</u>	Protein coding	<u>CCDS21267</u> &	<u>Q03391</u> &	TSL:1 GENCODE basic APPRIS P1		

The strategy is based on the design of Grin2d-201 transcript, The transcription is shown below



Genomic location distribution





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Protein domain



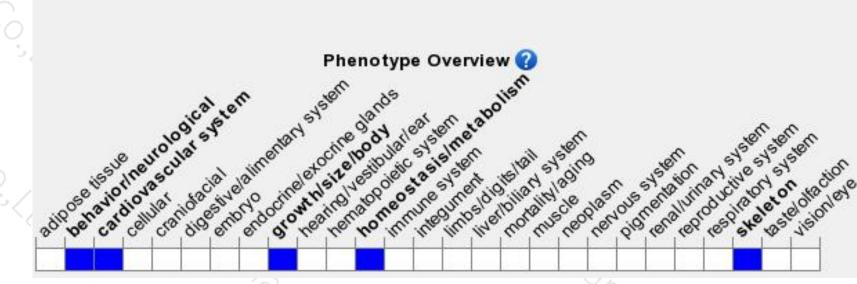
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	Cleavage site (Sign Superfamily	Periplasmic binding protein-like	1	SSF53850					
	CMART								
	SMART			Ionotropic glutamate receptor	ceptor, L-glutamate and glycine-bi	ndina domain			
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Mouse phenotype 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, Homozygotes for a targeted null mutation exhibit reduced spontaneous activity and an elevated auditory brainstem response threshold.

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



