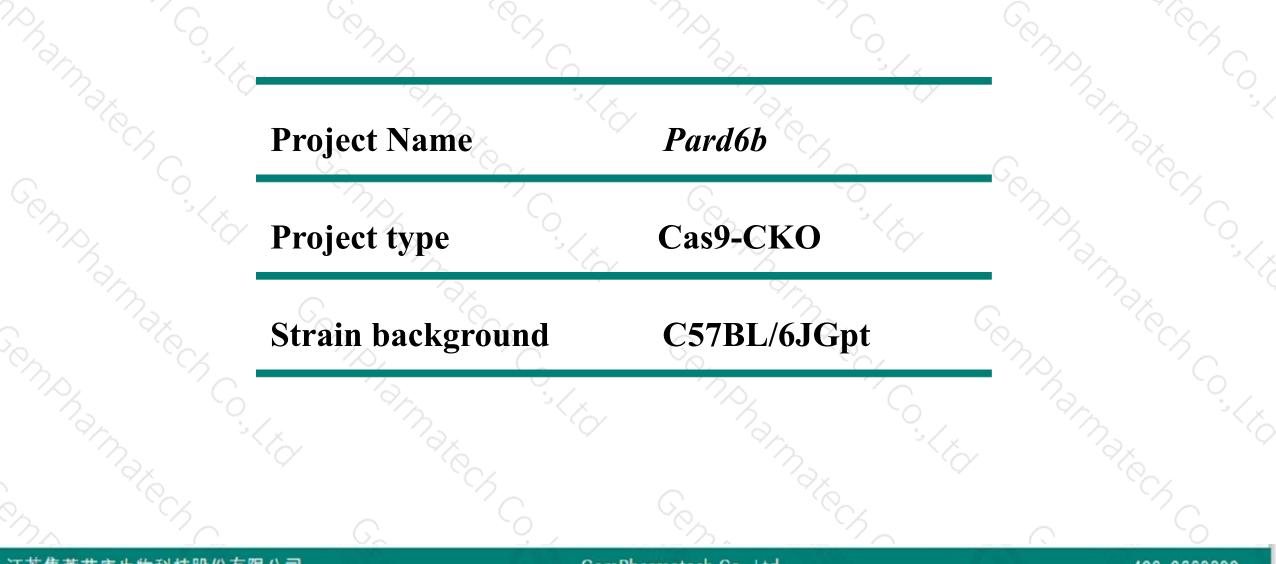


Pard6b Cas9-CKO Strategy

Designer: Reviewer: Design Date: JiaYu Xiaojing Li 2020-3-9

Project Overview





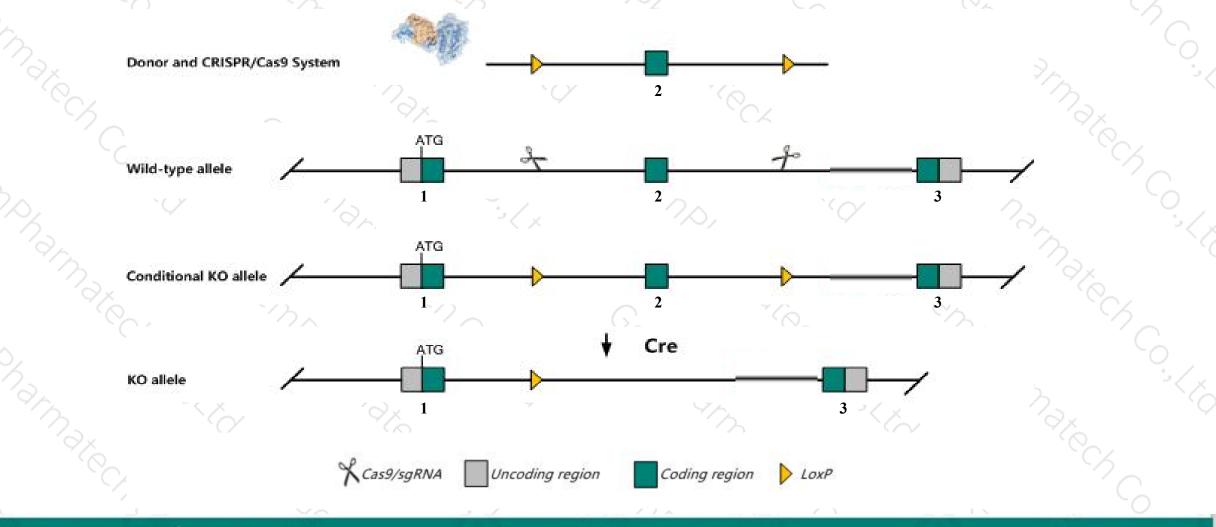
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Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Pard6b* gene. The schematic diagram is as follows:



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The Pard6b gene has 1 transcript. According to the structure of Pard6b gene, exon2 of Pard6b-201 (ENSMUST00000052125.6) transcript is recommended as the knockout region. The region contains 223bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Pard6b* gene. The brief process is as follows:CRISPR/Cas9 system and Donor 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.

The flox mice will be knocked out after mating with mice expressing Cre recombinase, resulting in the loss of function of the target gene in specific tissues and cell types.

Notice



- The Pard6b gene is located on the Chr2. 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 loxp insertion on gene transcription, RNA splicing and protein translation cannot be predicted at existing technological level.

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Gene information (NCBI)

Pard6b par-6 family cell polarity regulator beta [Mus musculus (house mouse)]



- Summary

Official Symbol Pard6b provided by MGI Official Full Name par-6 family cell polarity regulator beta provided by MGI Primary source MGI:MGI:2135605 See related

Ensembl:ENSMUSG0000044641

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 Par6b; AV025615

Expression Broad expression in lung adult (RPKM 10.3), kidney adult (RPKM 8.3) and 23 other tissues See more

Orthologs human all





☆ ?

Transcript information (Ensembl)



The gene has 1 transcript, and the transcript is shown below:

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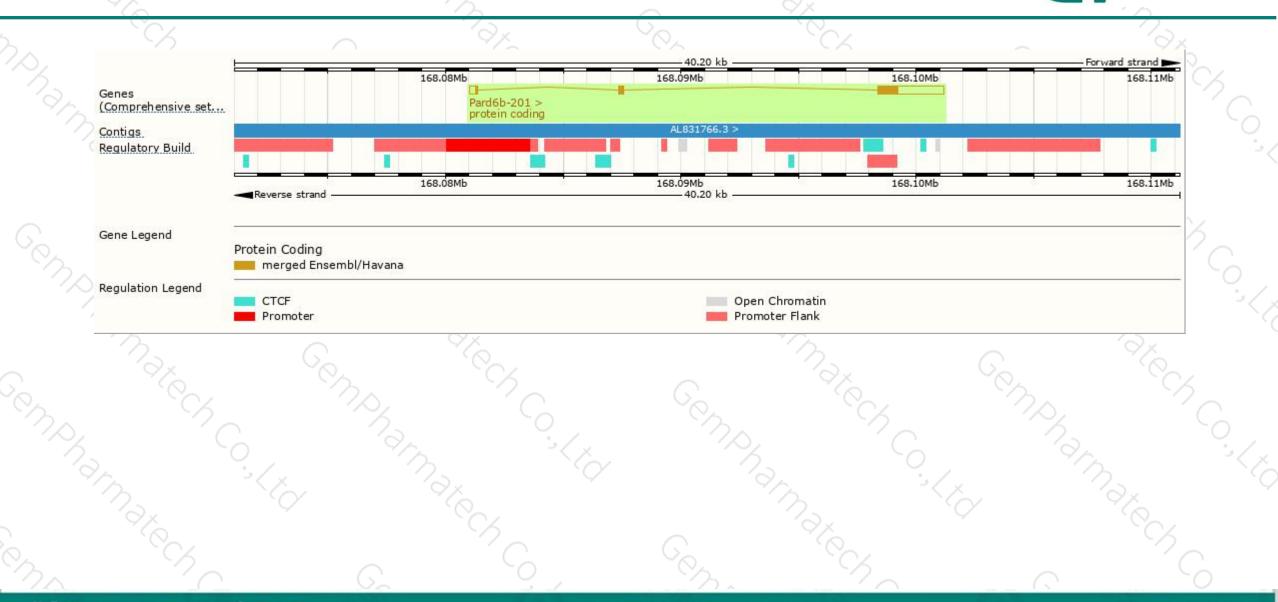
Name 🍦	Transcript ID 👙	bp 🖕	Protein 🖕	Biotype 🍦	CCDS 🖕	UniProt 🖕	Flags 🍦		
Pard6b-201	ENSMUST0000052125.6	3399	<u>371aa</u>	Protein coding	<u>CCDS17108</u> 团	<u>Q9JK83</u> &	TSL:1	GENCODE basic	APPRIS P1

The strategy is based on the design of Pard6b-201 transcript, The transcription is shown below

Pard6b-201 > protein coding

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Genomic location distribution



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400-9660890

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Protein domain



	CA							100
	ENSMUSP00000052 PDB-ENSP mappings MobiDB lite Low complexity (Seg) Superfamily	SSF54277		PDZ superfamily				 %
	SMART	PB1 domain	2	PDZ domain				· · · · · · · · · · · · · · · · · · ·
	Pfam.	PB1 domain		PDZ domain				
	PROSITE profiles	PB1 domain		PDZ domain				
\mathcal{D}	PANTHER	Partitioning defective 6 homologue	e beta					<u> </u>
	Gene3D	PTHR14102 3.10.20.90	2.3	30,42,10				
	CDD	Partitioning defective prot	ein 6, PB1 domain	cd00992				
	All sequence SNPs/i	Sequence variants (dbSNP and	all other sources)	1.1		3	2	<u> </u>
	Variant Legend	synonymous variant						
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If you have any questions, you are welcome to inquire. Tel: 400-9660890



