

Kat6b Cas9-KO Strategy

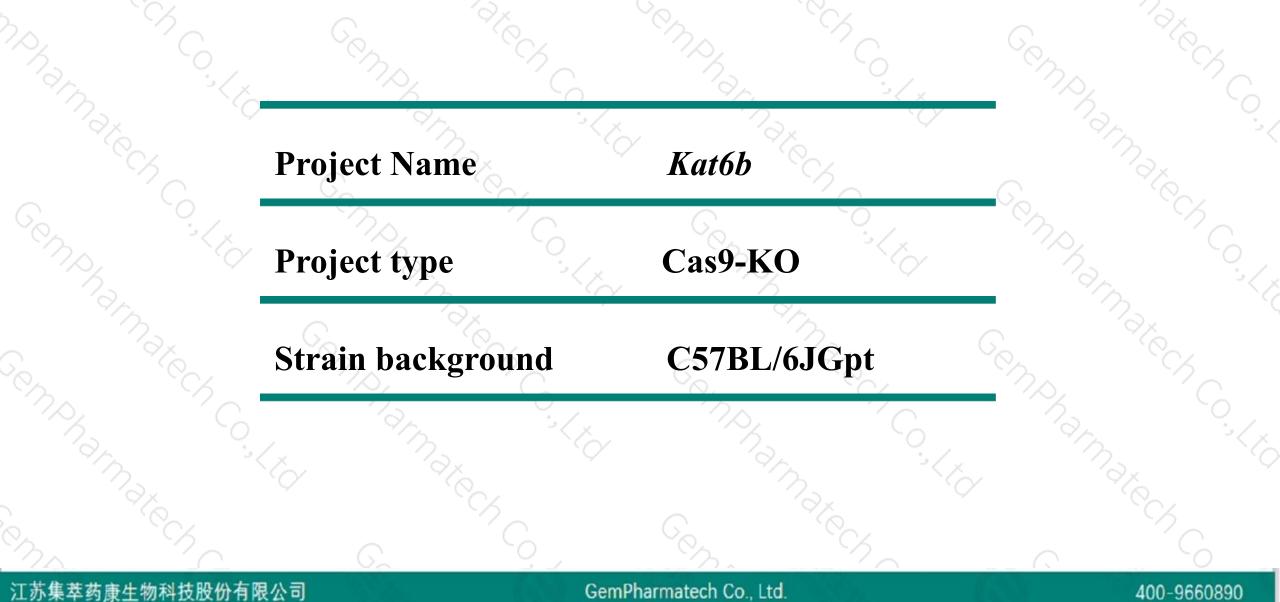
Designer: Reviewer:

Design Date:

Daohua Xu Huimin Su 2019-12-11

Project Overview

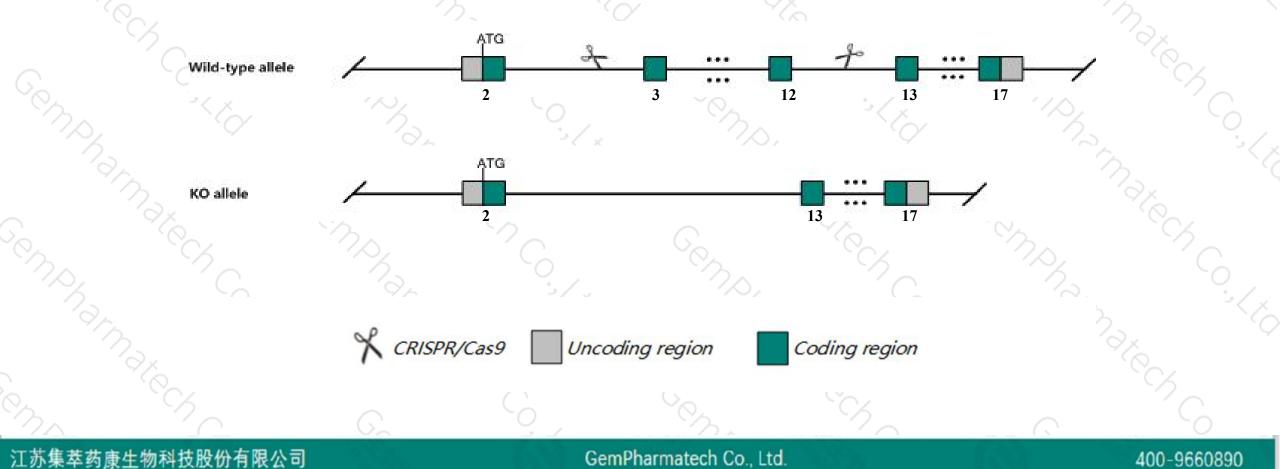




Knockout strategy



This model will use CRISPR/Cas9 technology to edit the Kat6b gene. The schematic diagram is as follows:





- The Kat6b gene has 14 transcripts. According to the structure of Kat6b gene, exon3-exon12 of Kat6b-204 (ENSMUST00000182405.8) transcript is recommended as the knockout region. The region contains 1132bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Kat6b gene. The brief process is as follows: CRISPR/Cas9 system

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- According to the existing MGI data, Reduced expression of this gene results in developmental defects of the skeleton and brain, particularly the cerebral cortex.
- Transcript *Kat6b-203* may not be affected.
- The Kat6b gene is located on the Chr14. 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.

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Gene information (NCBI)



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Kat6b K(lysine) acetyltransferase 6B [Mus musculus (house mouse)]

Gene ID: 54169, updated on 31-Jan-2019

Summary

Official SymbolKat6b provided by MGIOfficial Full NameK(lysine) acetyltransferase 6B provided byMGIPrimary sourceMGI:MGI:1858746Primary sourceInsembl:ENSMUSG0000021767Gene typeprotein codingRefSeq statusVALIDATEDOrganismMus musculusLineageEukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha;
Muroidea; Murinae; Mus; MusAlso knownasAl507552, B130044K16Rik, MYST-4, Morf, Myst4, mKIAA0383, qkf, querkopfExpressionUbiquitous expression in limb E14.5 (RPKM 5.4), frontal lobe adult (RPKM 5.1) and 28 other tissuesSee more
human all

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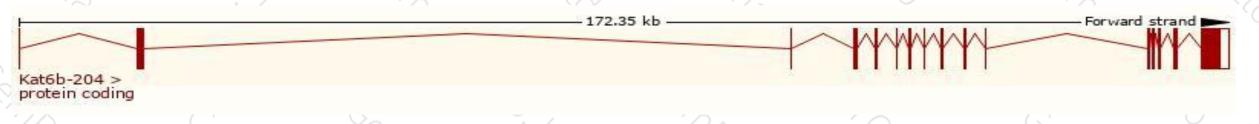
Transcript information (Ensembl)



The gene has 14 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kat6b-204	ENSMUST00000182405.8	6977	<u>1763aa</u>	Protein coding	CCDS59615	Q8BRB7	TSL:1 GENCODE basic APPRIS P2
Kat6b-207	ENSMUST00000182855.7	6791	<u>1763aa</u>	Protein coding	CCDS59615	<u>Q8BRB7</u>	TSL:1 GENCODE basic APPRIS P2
Kat6b-209	ENSMUST00000182964.2	7699	<u>1872aa</u>	Protein coding	18	Q8BRB7	TSL:5 GENCODE basic APPRIS ALT2
Kat6b-201	ENSMUST0000069648.13	7649	<u>1872aa</u>	Protein coding	20 	Q8BRB7	TSL:5 GENCODE basic APPRIS ALT2
Kat6b-203	ENSMUST00000112458.2	997	<u>179aa</u>	Protein coding	₹4	<u>Q14AW2</u>	TSL:5 GENCODE basic
Kat6b-210	ENSMUST00000182996.1	606	<u>79aa</u>	Protein coding	,	<u>S4R2Q8</u>	CDS 3' incomplete TSL:2
Kat6b-206	ENSMUST00000182732.1	4671	No protein	Retained intron	10	-	TSL:1
Kat6b-213	ENSMUST00000183228.1	3873	No protein	Retained intron	20 20	1 12	TSL:1
Kat6b-202	ENSMUST0000096222.11	3508	No protein	Retained intron	74		TSL:1
Kat6b-211	ENSMUST00000183055.1	2680	No protein	Retained intron	,	-	TSL:1
Kat6b-208	ENSMUST00000182859.1	2125	No protein	Retained intron	45	-	TSL:1
Kat6b-205	ENSMUST00000182553.1	839	No protein	Retained intron		1 12	TSL:3
Kat6b-212	ENSMUST00000183201.7	535	No protein	IncRNA	74		TSL:3
Kat6b-214	ENSMUST00000226589.1	18	No protein	IncRNA	-		

The strategy is based on the design of Kat6b-204 transcript, The transcription is shown below

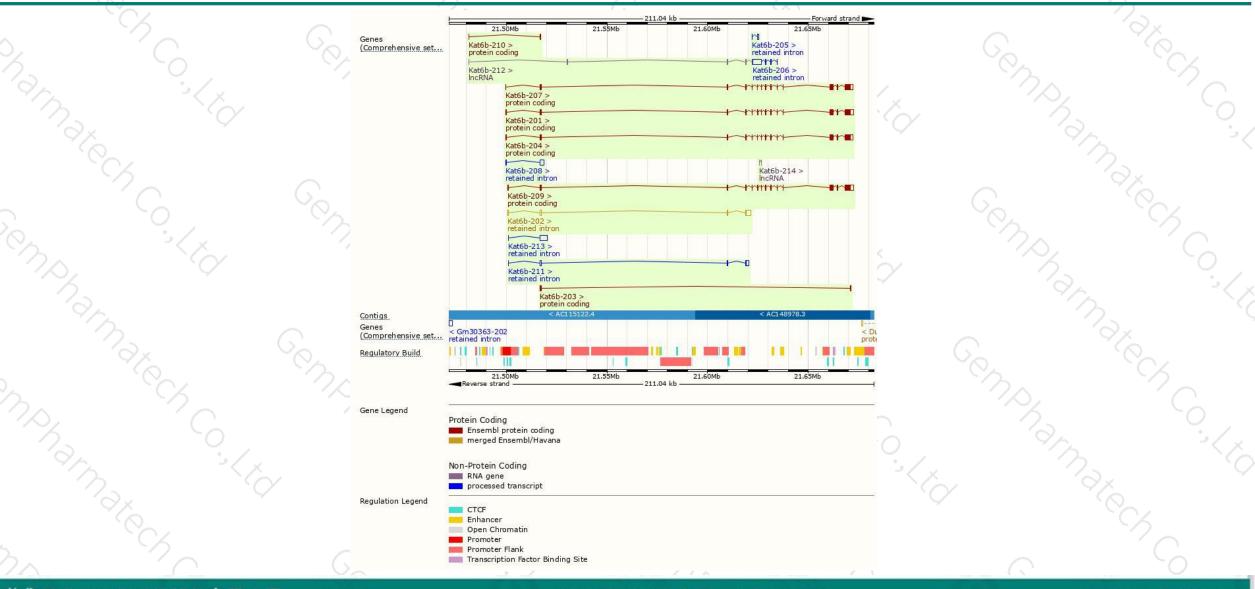


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Genomic location distribution



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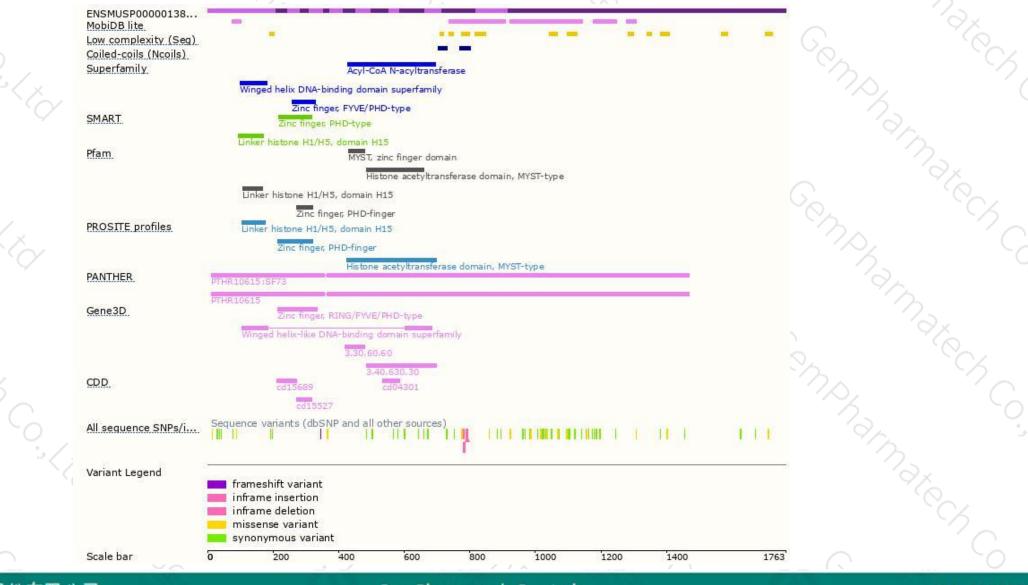
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Protein domain





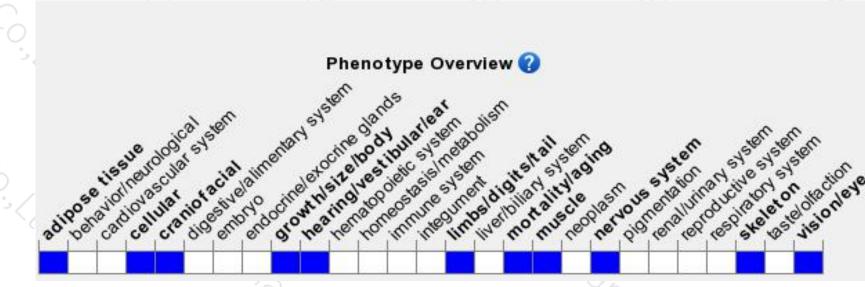
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Mouse phenotype description(MGI)





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, Reduced expression of this gene results in developmental defects of the skeleton and brain, particularly the cerebral cortex.



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



