

# ***Map3k15* Cas9-KO Strategy**

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# Project Overview

**Project Name**

*Map3k15*

**Project type**

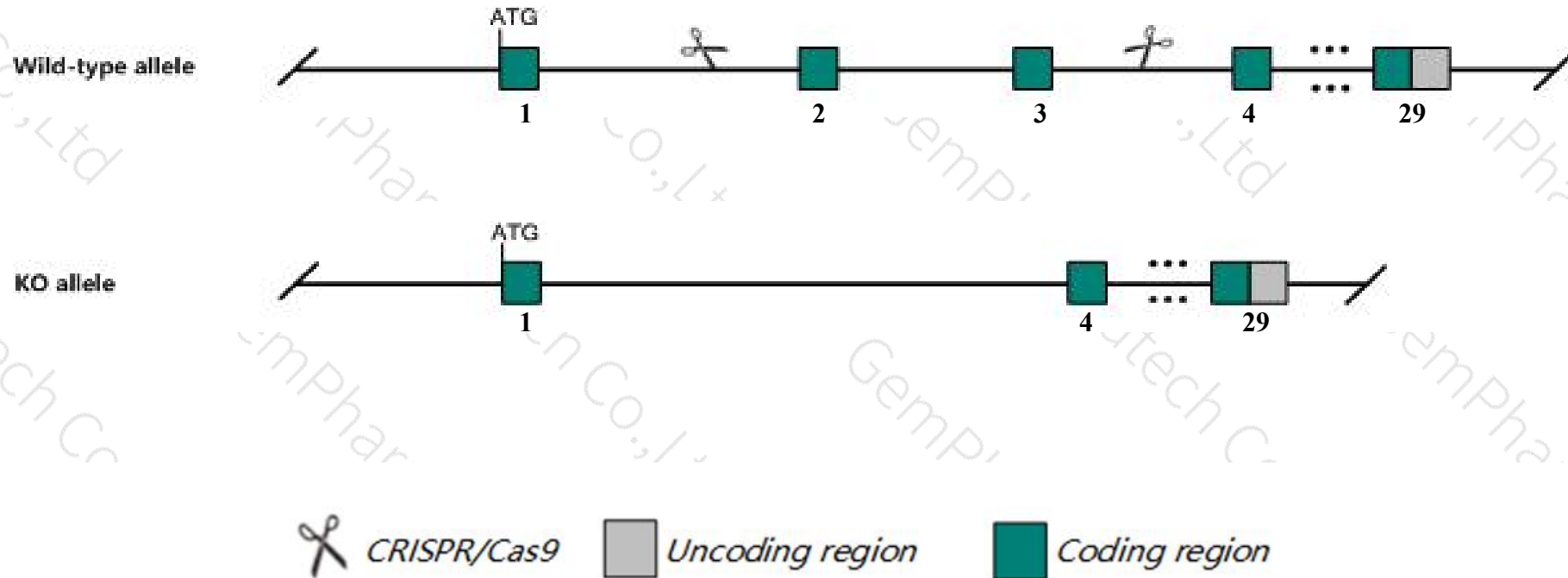
**Cas9-KO**

**Strain background**

**C57BL/6JGpt**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Map3k15* gene. The schematic diagram is as follows:



- The *Map3k15* gene has 1 transcript. According to the structure of *Map3k15* gene, exon2-exon3 of *Map3k15-201* (ENSMUST00000033665.8) transcript is recommended as the knockout region. The region contains 164bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Map3k15* gene. The brief process is as follows: CRISPR/Cas9 syst

- According to the existing MGI data, Male mice hemizygous for a knock-out allele exhibit hyperactivation of the SPAK/OSR1 and Na<sup>+</sup>Cl<sup>-</sup> cotransporter (NCC) pathway in the renal tubules and develop mild hypertension.
- The *Map3k15* gene is located on the ChrX. 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)

## Map3k15 mitogen-activated protein kinase kinase kinase 15 [Mus musculus (house mouse)]

Gene ID: 270672, updated on 31-Jan-2019

### Summary



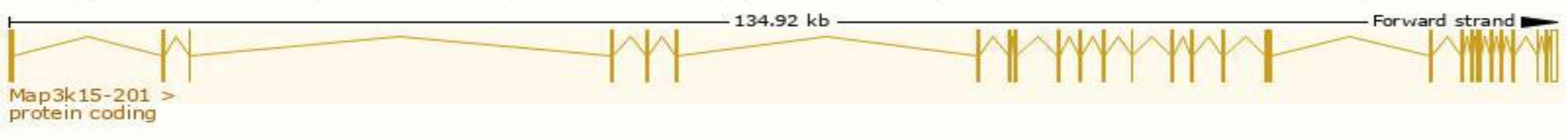
<b>Official Symbol</b>	Map3k15 provided by <a href="#">MGI</a>
<b>Official Full Name</b>	mitogen-activated protein kinase kinase kinase 15 provided by <a href="#">MGI</a>
<b>Primary source</b>	<a href="#">MGI:MGI:2448588</a>
<b>See related</b>	<a href="#">Ensembl:ENSMUSG000000031303</a>
<b>Gene type</b>	protein coding
<b>RefSeq status</b>	VALIDATED
<b>Organism</b>	<a href="#">Mus musculus</a>
<b>Lineage</b>	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
<b>Also known as</b>	BC031147, MEKK15
<b>Expression</b>	Broad expression in heart adult (RPKM 24.0), kidney adult (RPKM 12.9) and 22 other tissues <a href="#">See more</a>
<b>Orthologs</b>	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Map3k15-201	<a href="#">ENSMUST00000033665.8</a>	4348	<a href="#">1331aa</a>	Protein coding	<a href="#">CCDS53236</a>	<a href="#">A2AQW0</a>	TSL:5 GENCODE basic APPRIS P1

The strategy is based on the design of *Map3k15-201* transcript, The transcription is shown below

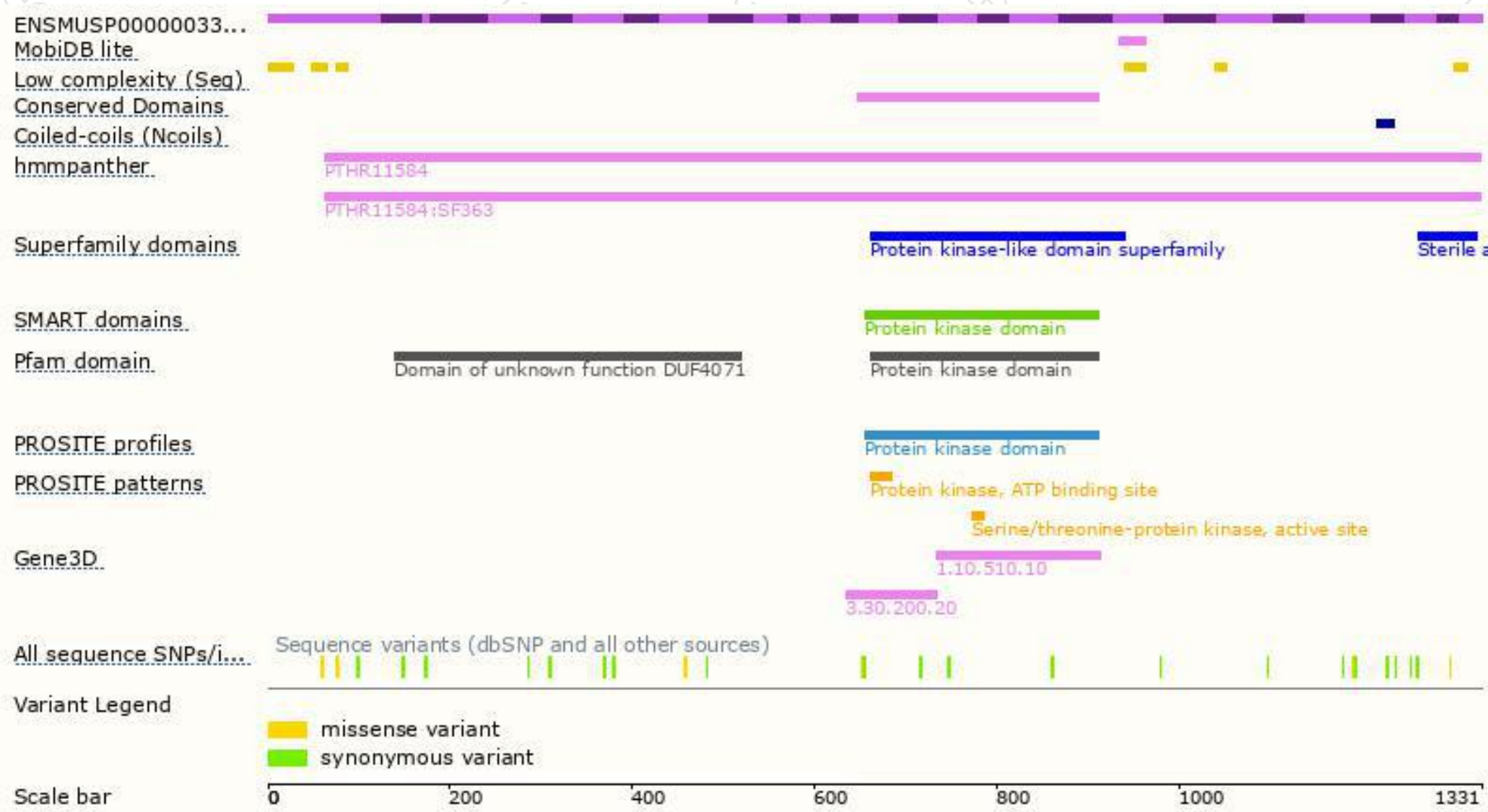


# Genomic location distribution

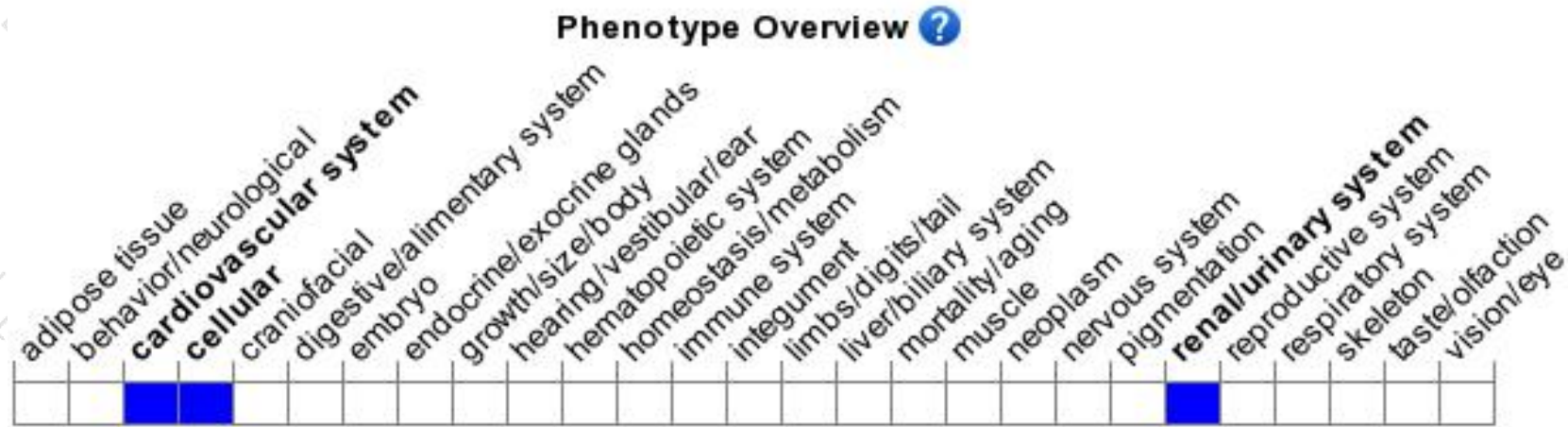




# Protein domain



# 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, Male mice hemizygous for a knock-out allele exhibit hyperactivation of the SPAK/OSR1 and Na<sup>+</sup>Cl<sup>-</sup> cotransporter (NCC) pathway in the renal tubules and develop mild hypertension.

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

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