

Kcnq2 Cas9-CKO Strategy

Designer:

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Project Overview

Project Name

Kcnq2

Project type

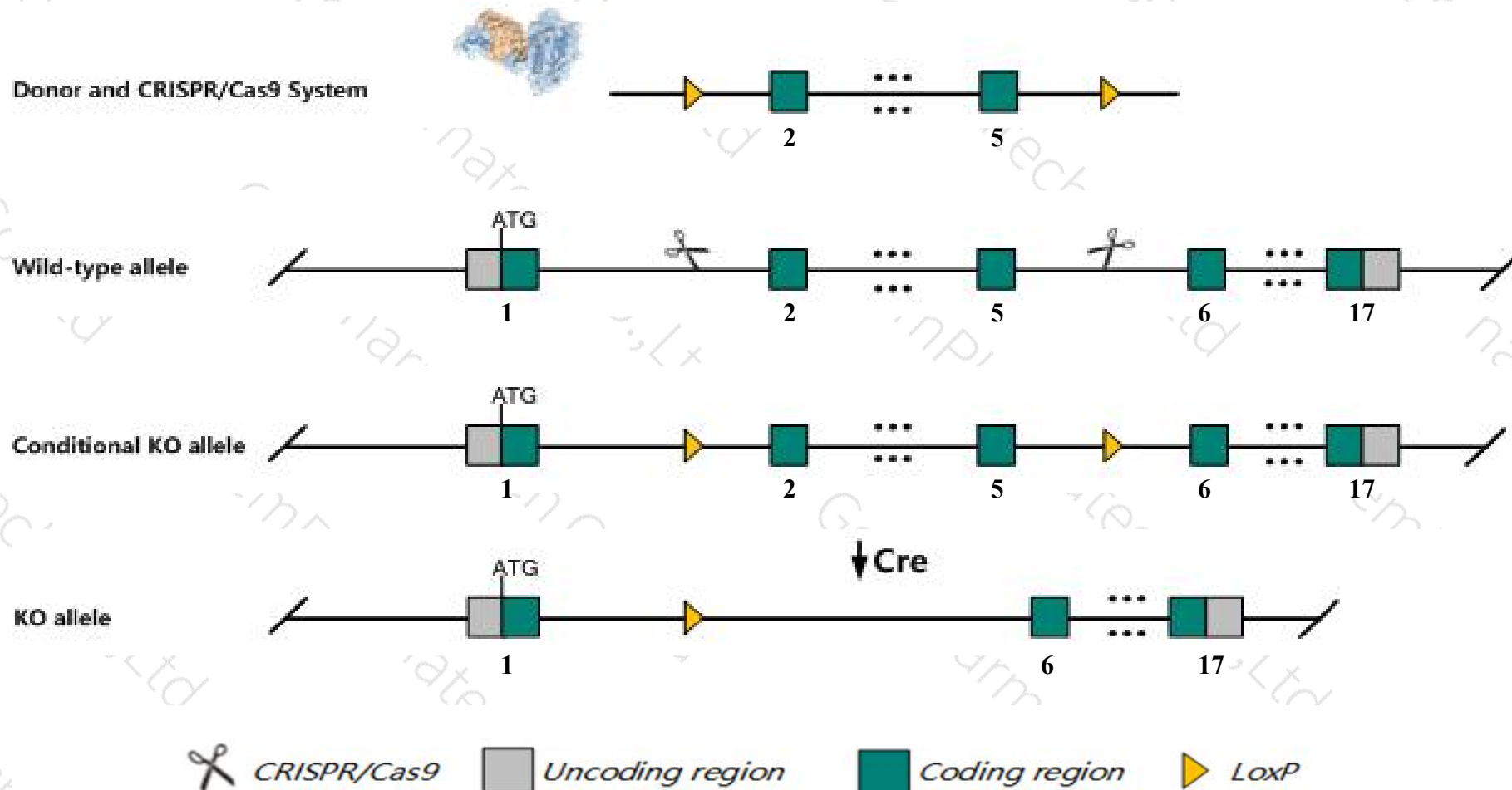
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Kcnq2* gene. The schematic diagram is as follows:



- The *Kcnq2* gene has 21 transcripts. According to the structure of *Kcnq2* gene, exon2-exon5 of *Kcnq2*-217 (ENSMUST00000149964.8) transcript is recommended as the knockout region. The region contains 520bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Kcnq2* 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.

- According to the existing MGI data, Mice homozygous for a null mutation die perinatally with pulmonary atelectasis. Heterozygous mice exhibit a hypersensitivity to the epileptic inducer pentylenetetrazole. Mice homozygous for a knock-in allele exhibit spontaneous seizures and premature death.
- The *Kcnq2* 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.

Gene information (NCBI)

Kcnq2 potassium voltage-gated channel, subfamily Q, member 2 [Mus musculus (house mouse)]

Gene ID: 16536, updated on 16-Feb-2019

Summary



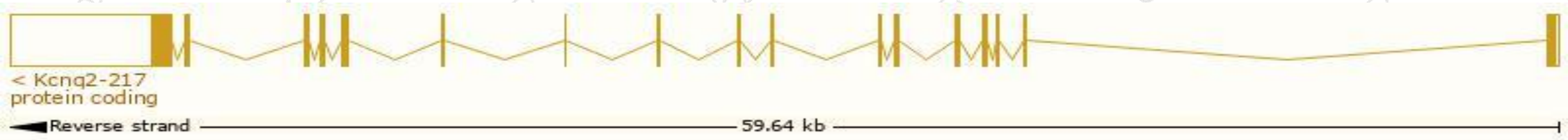
Official Symbol	Kcnq2 provided by MGI
Official Full Name	potassium voltage-gated channel, subfamily Q, member 2 provided by MGI
Primary source	MGI:MGI:1309503
See related	Ensembl:ENSMUSG00000016346
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	HNSPC, KQT2, Nmf134
Expression	Biased expression in cortex adult (RPKM 26.0), frontal lobe adult (RPKM 26.0) and 5 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

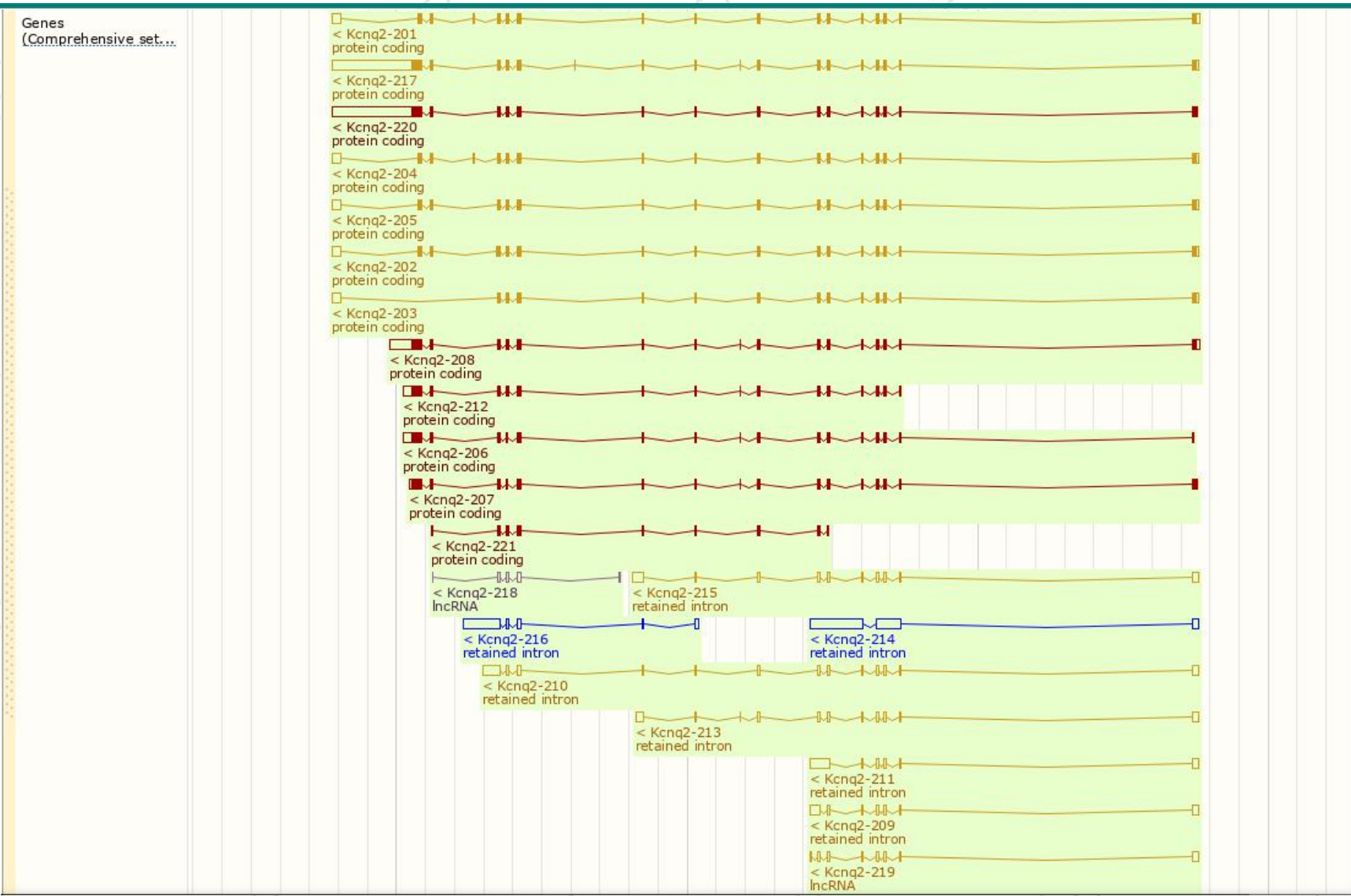
The gene has 21 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Kcnq2-217	ENSMUST00000149964.8	8209	870aa	Protein coding	CCDS17198	B7ZBV9	TSL:1 Gencode basic APPRIS P2
Kcnq2-201	ENSMUST0000016491.13	3067	759aa	Protein coding	CCDS17193	B7ZBV4	TSL:1 Gencode basic
Kcnq2-202	ENSMUST0000049792.14	2920	754aa	Protein coding	CCDS17194	B7ZBV6	TSL:1 Gencode basic
Kcnq2-204	ENSMUST00000103047.9	2899	747aa	Protein coding	CCDS17195	B7ZBV8	TSL:1 Gencode basic
Kcnq2-205	ENSMUST00000103048.9	2827	723aa	Protein coding	CCDS17196	B7ZBV7	TSL:1 Gencode basic
Kcnq2-203	ENSMUST00000081528.12	2382	570aa	Protein coding	CCDS17197	B7ZBV5	TSL:1 Gencode basic
Kcnq2-220	ENSMUST00000197015.4	8031	842aa	Protein coding	-	A0A0G2JFQ2	TSL:5 Gencode basic APPRIS ALT2
Kcnq2-208	ENSMUST00000103051.8	4200	852aa	Protein coding	-	B7ZBW1	TSL:5 Gencode basic APPRIS ALT2
Kcnq2-206	ENSMUST00000103049.9	2936	795aa	Protein coding	-	F6UC55	CDS 5' incomplete TSL:1
Kcnq2-212	ENSMUST00000129695.7	2735	726aa	Protein coding	-	B7ZBV3	CDS 5' incomplete TSL:1
Kcnq2-207	ENSMUST00000103050.9	2713	839aa	Protein coding	-	B7ZBW2	TSL:5 Gencode basic APPRIS ALT2
Kcnq2-221	ENSMUST00000197599.1	855	285aa	Protein coding	-	A0A0G2JGA6	CDS 5' and 3' incomplete TSL:5
Kcnq2-214	ENSMUST00000140789.1	5796	No protein	Retained intron	-	-	TSL:2
Kcnq2-216	ENSMUST00000145861.7	3131	No protein	Retained intron	-	-	TSL:1
Kcnq2-210	ENSMUST00000129073.7	2758	No protein	Retained intron	-	-	TSL:2
Kcnq2-211	ENSMUST00000129361.2	2243	No protein	Retained intron	-	-	TSL:2
Kcnq2-215	ENSMUST00000144592.7	2011	No protein	Retained intron	-	-	TSL:2
Kcnq2-213	ENSMUST00000139458.7	1816	No protein	Retained intron	-	-	TSL:2
Kcnq2-209	ENSMUST00000123336.7	1686	No protein	Retained intron	-	-	TSL:2
Kcnq2-219	ENSMUST00000154164.7	1226	No protein	lncRNA	-	-	TSL:2
Kcnq2-218	ENSMUST00000152099.6	513	No protein	lncRNA	-	-	TSL:5

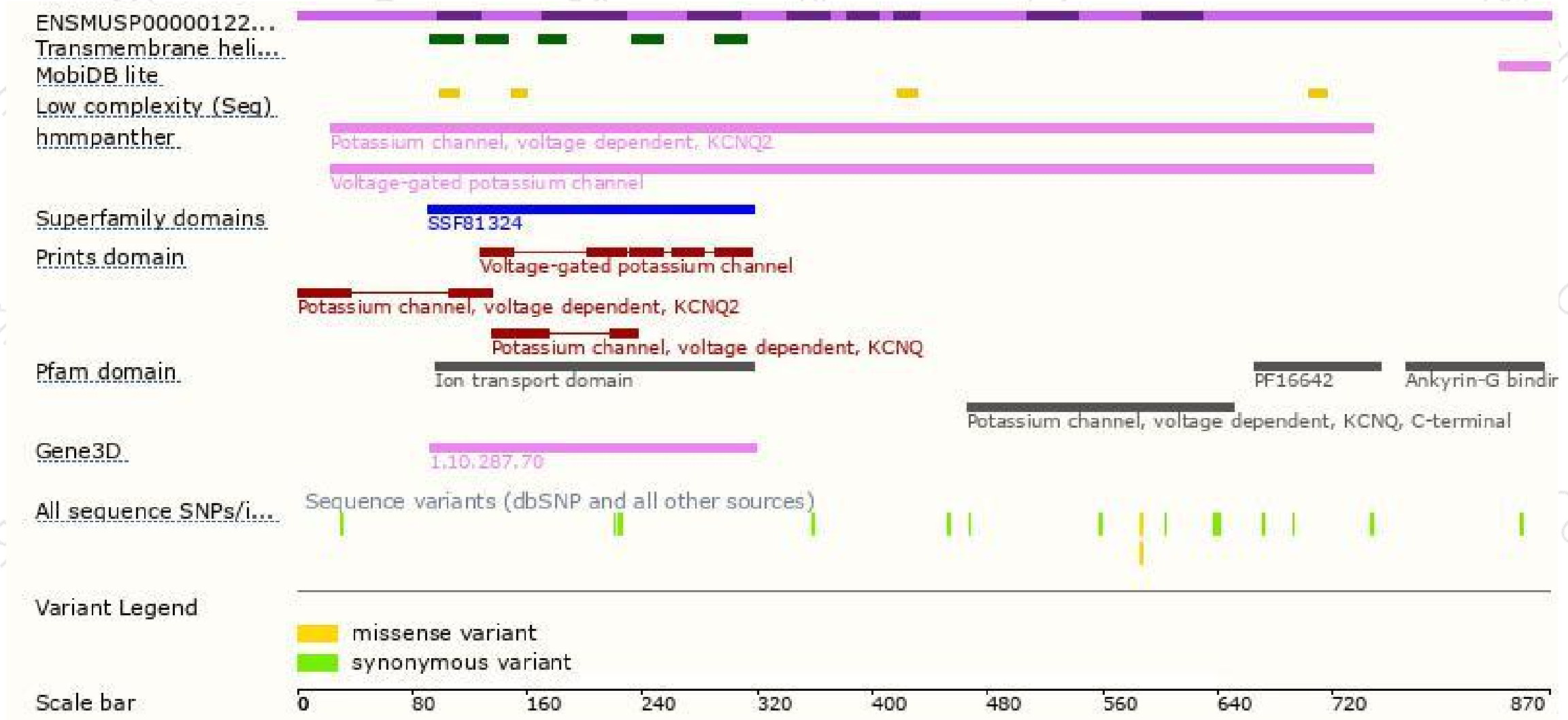
The strategy is based on the design of *Kcnq2-217* transcript,The transcription is shown below



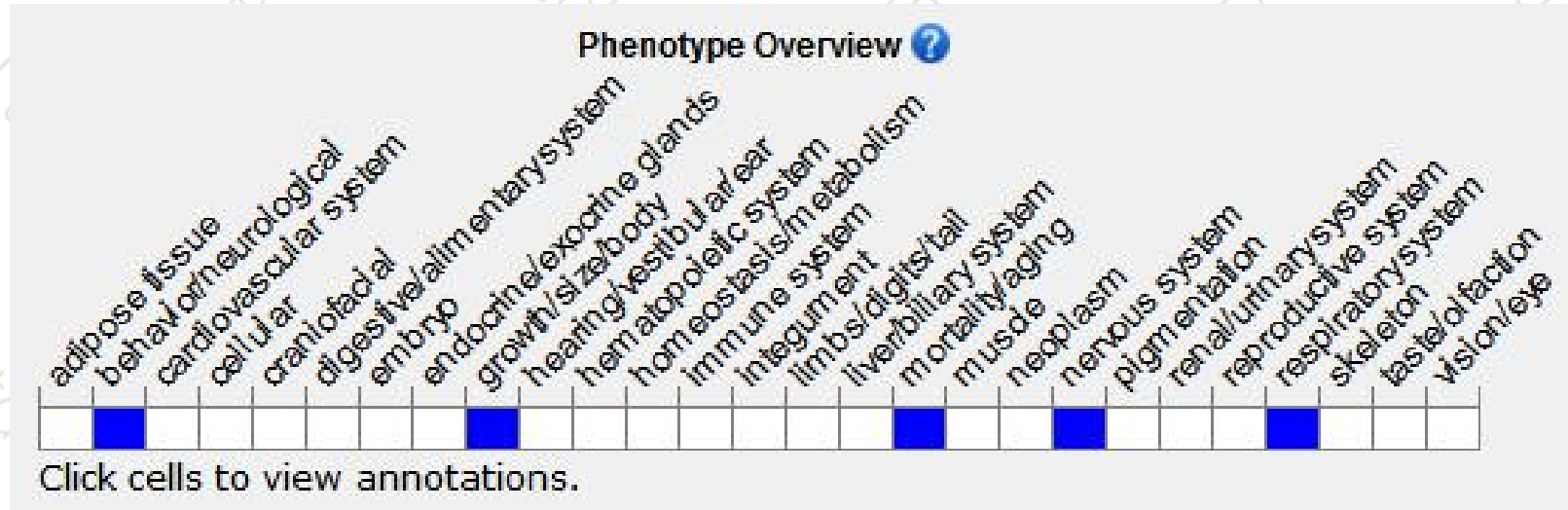
Genomic location distribution



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, Mice homozygous for a null mutation die perinatally with pulmonary atelectasis.

Heterozygous mice exhibit a hypersensitivity to the epileptic inducer pentylenetetrazole. Mice homozygous for a knock-in allele exhibit spontaneous seizures and premature death.

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

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