Bpi Cas9-CKO Strategy

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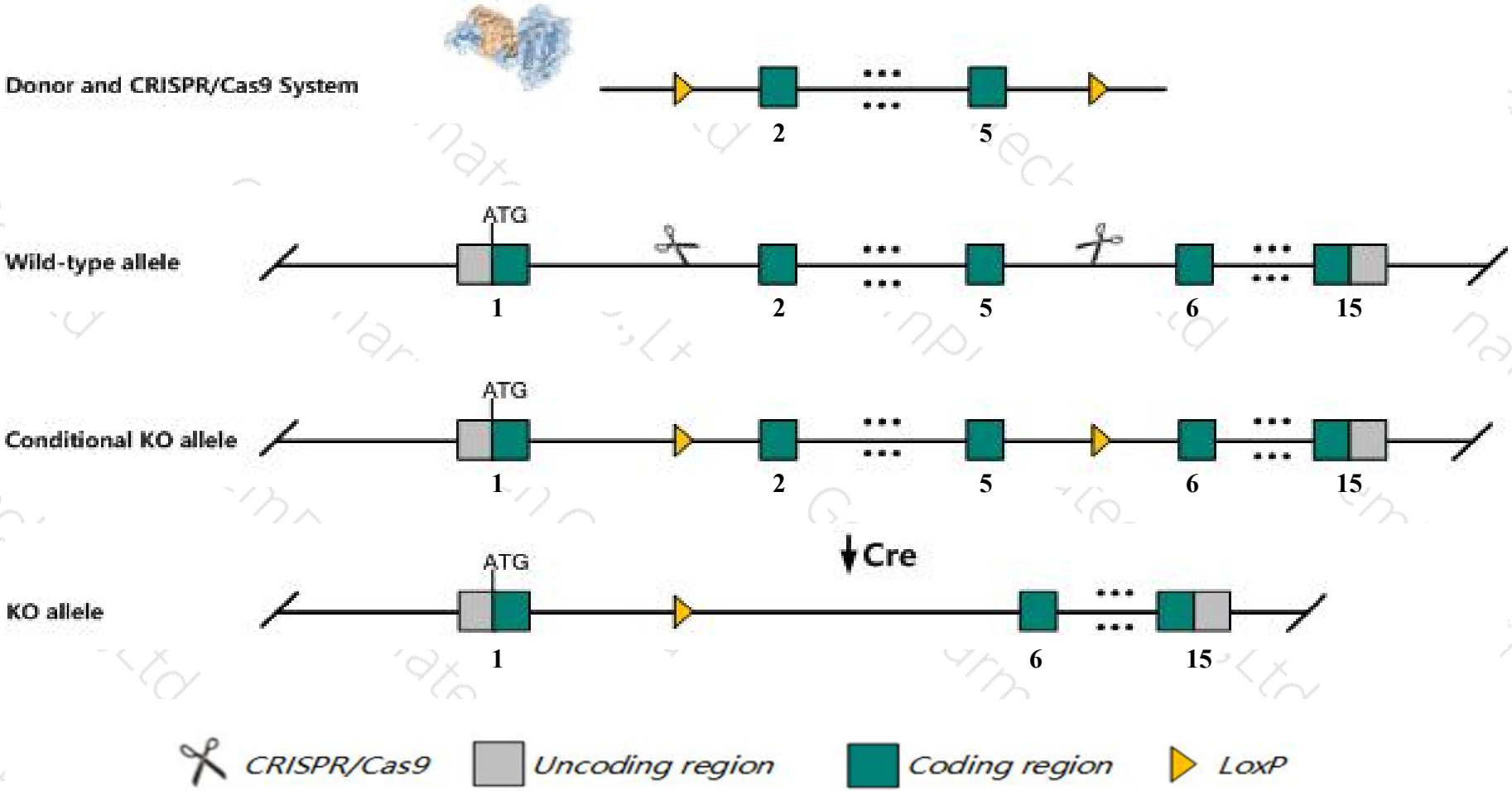
Project Overview

Project Name	***Bpi***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

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



Technical routes

- The *Bpi* gene has 3 transcripts. According to the structure of *Bpi* gene, exon2-exon5 of *Bpi-201* (ENSMUST00000065039.2) transcript is recommended as the knockout region.The region contains 467bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Bpi* 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 *Bpi* 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）

集萃药康 GemPharmatech

Bpi bactericidal permeablility increasing protein [Mus musculus (house mouse)]

Gene ID: 329547, updated on 31-Jan-2019

Summary

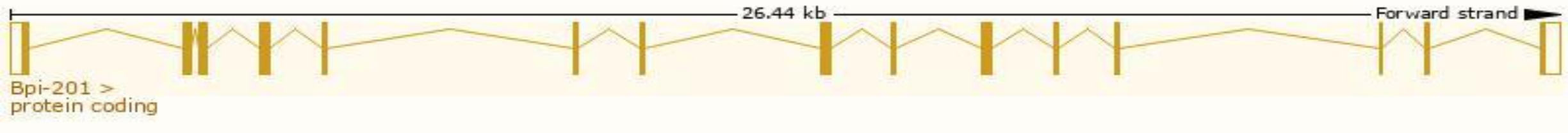
Official Symbol	Bpi provided by MGI
Official Full Name	bactericidal permeablility increasing protein provided byMGI
Primary source	MGI:MGI:3045315
See related	Ensembl:ENSMUSG00000052922
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	9230105K17Rik, Bpifd1
Expression	Biased expression in genital fat pad adult (RPKM 29.4) and testis adult (RPKM 24.9)See more
Orthologs	human all

Transcript information（Ensembl）

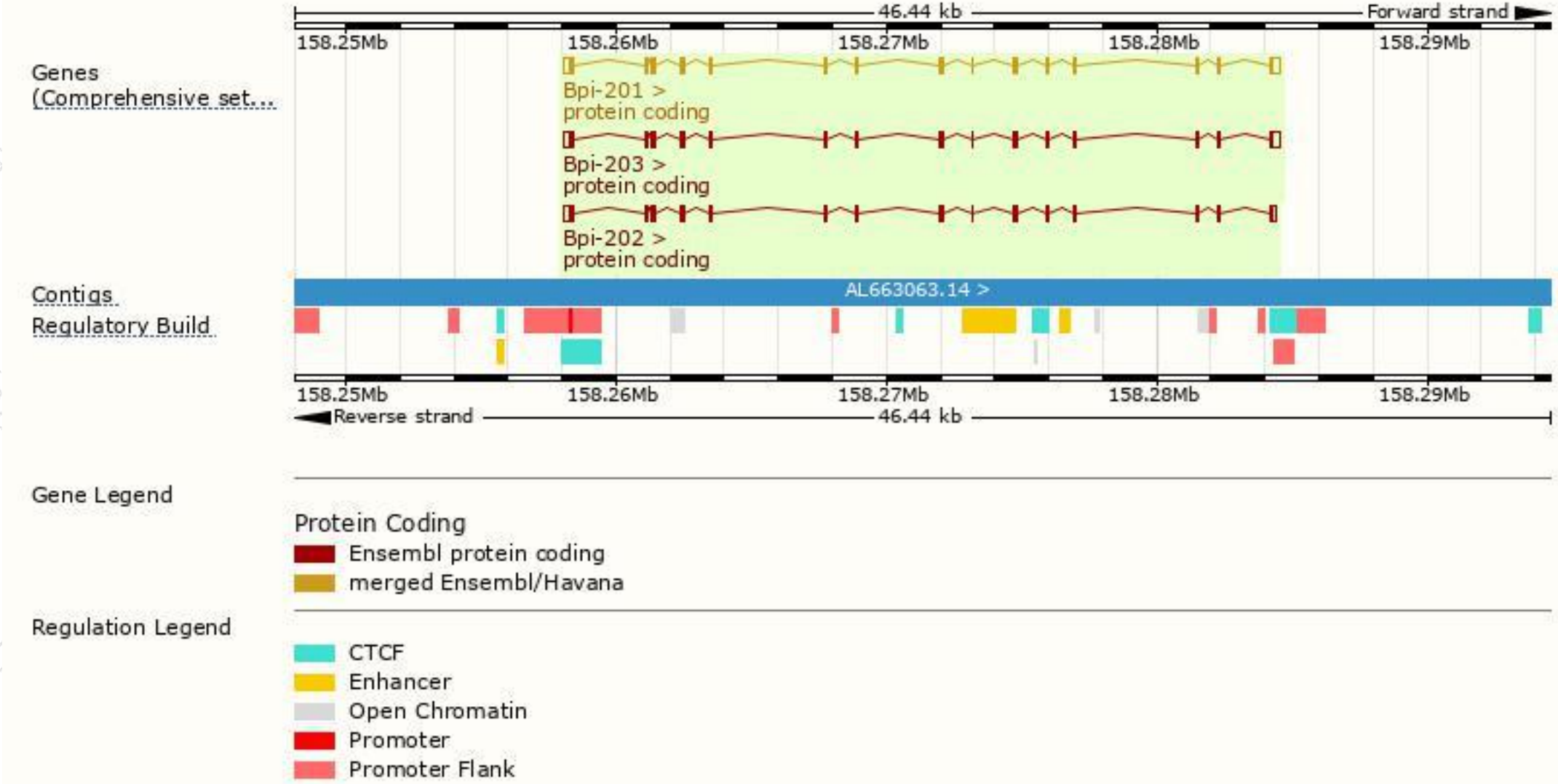
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Bpi-201	ENSMUST00000065039.2	1926	486aa	Protein coding	CCDS16987	Q67E05	TSL:1 GENCODE basic APPRIS P2
Bpi-203	ENSMUST00000109500.7	1911	482aa	Protein coding	-	Q67E05	TSL:5 GENCODE basic APPRIS ALT2
Bpi-202	ENSMUST00000109499.7	1778	483aa	Protein coding	-	Q67E05	TSL:1 GENCODE basic APPRIS ALT2

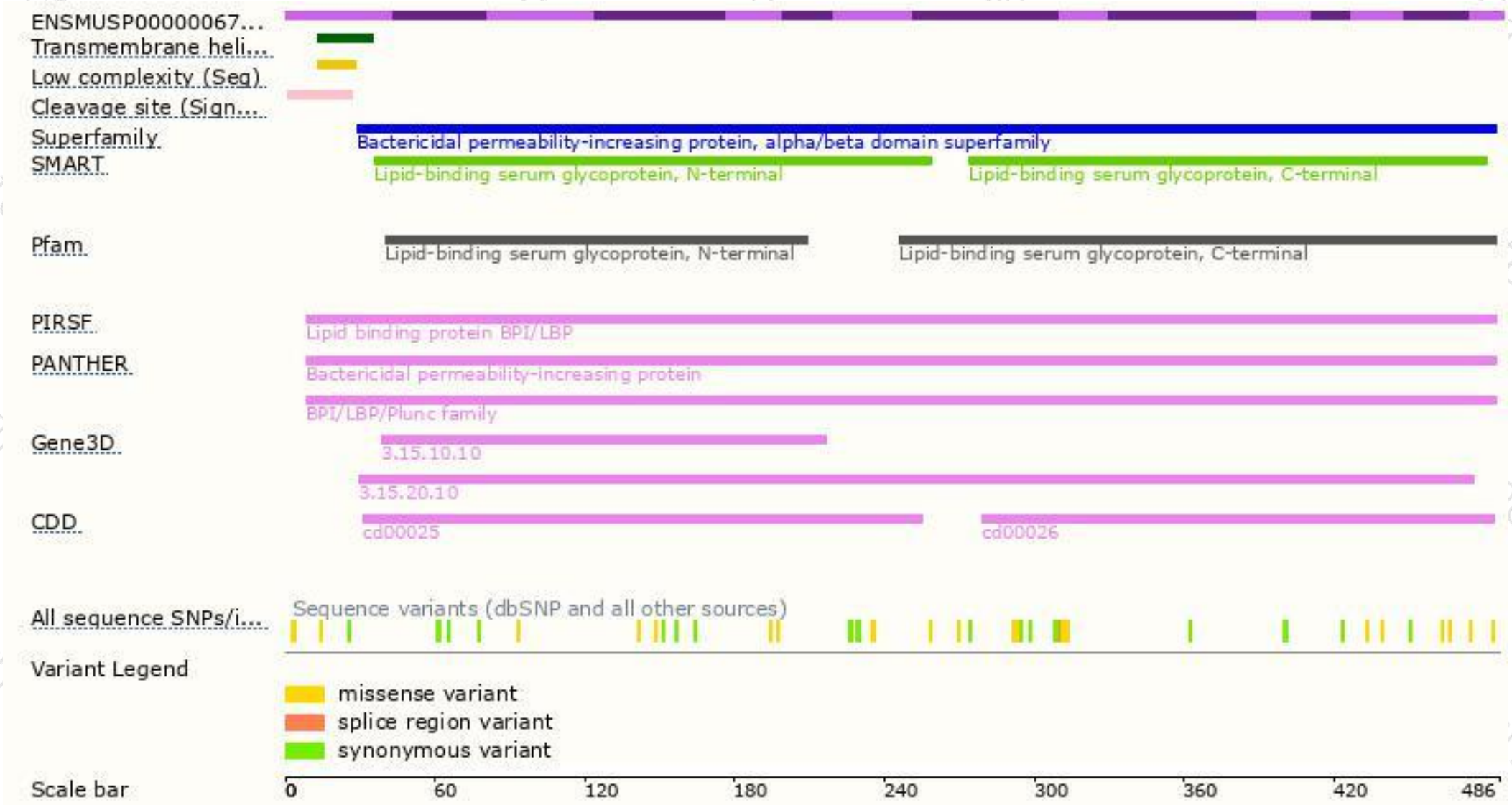
The strategy is based on the design of *Bpi-201* transcript,The transcription is shown below



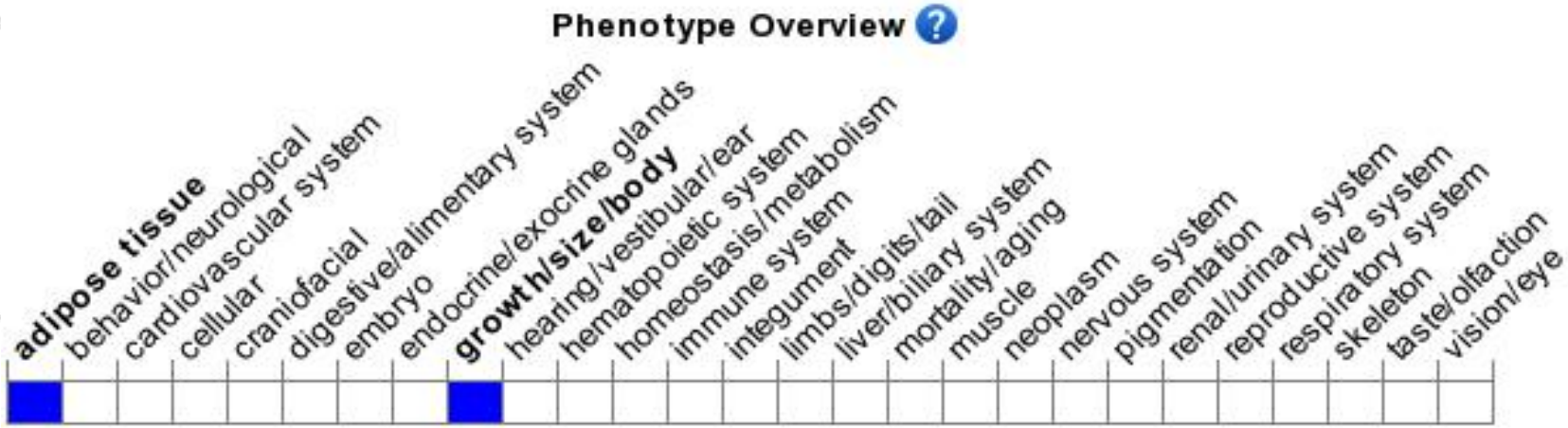
Genomic location distribution



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/).

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

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