

Ell Cas9-CKO Strategy

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Reviewer:

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Design Date:

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Project Overview

Project Name

Ell

Project type

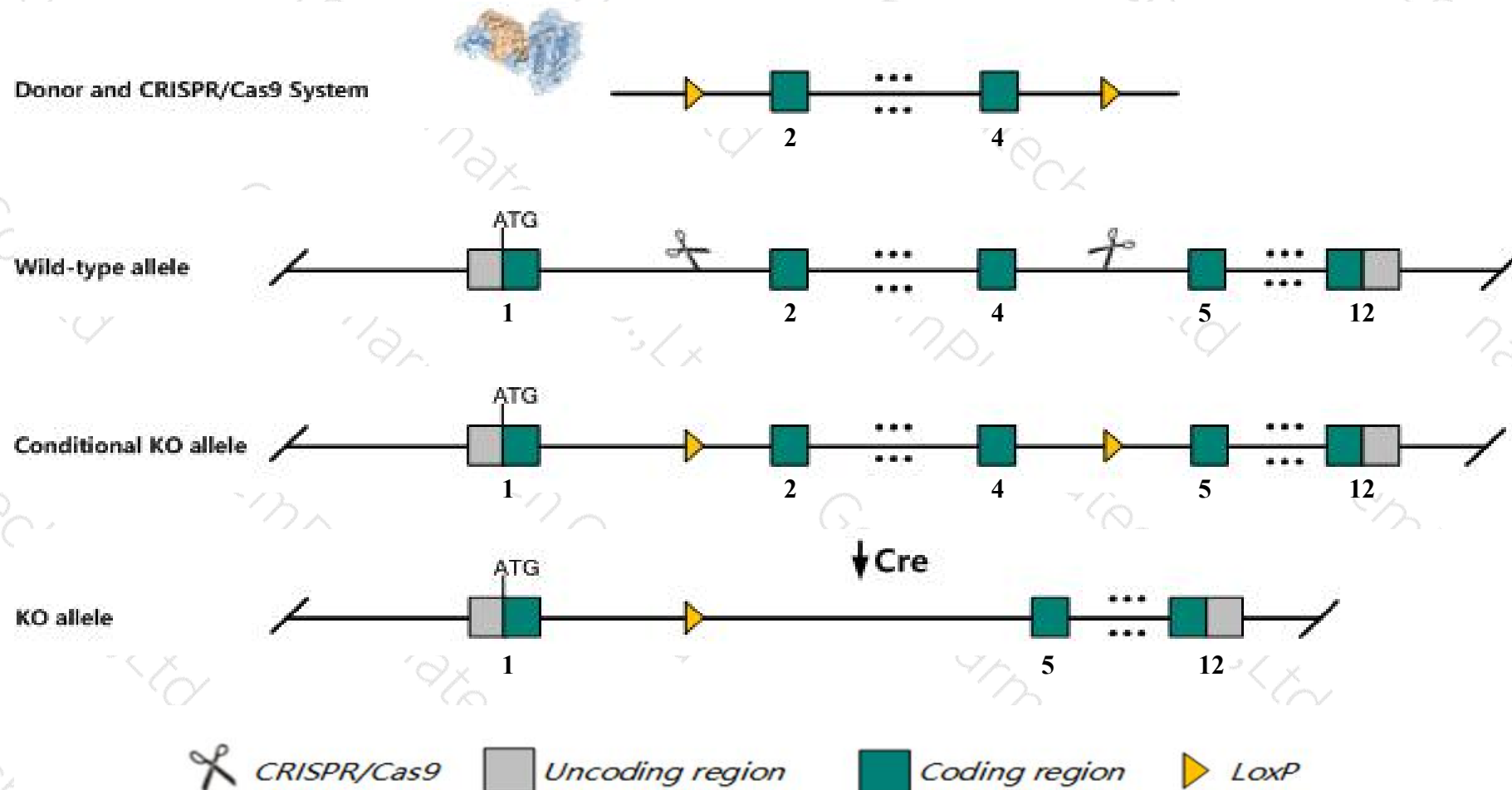
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

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



Technical routes

- The *Ell* gene has 2 transcripts. According to the structure of *Ell* gene, exon2-exon4 of *Ell*-201 (ENSMUST00000093454.7) transcript is recommended as the knockout region. The region contains 334bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Ell* 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

- According to the existing MGI data, Mice homozygous for a knock-out allele die prior to E6.5 but after implantation.
- The *Ell* gene is located on the Chr8. 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)

EII elongation factor RNA polymerase II [Mus musculus (house mouse)]

Gene ID: 13716, updated on 31-Jan-2019

Summary



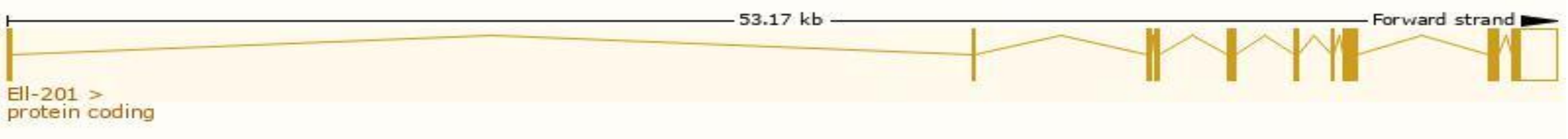
Official Symbol	EII provided by MGI
Official Full Name	elongation factor RNA polymerase II provided by MGI
Primary source	MGI:MGI:109377
See related	Ensembl:ENSMUSG00000070002
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	EII1, Men
Expression	Ubiquitous expression in testis adult (RPKM 19.6), colon adult (RPKM 17.4) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

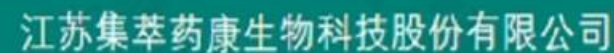
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
EII-201	ENSMUST00000093454.7	3072	602aa	Protein coding	CCDS22372	O08856	TSL:1 GENCODE basic APPRIS P1
EII-202	ENSMUST00000210155.1	457	52aa	Protein coding	-	A0A1B0GR04	CDS 3' incomplete TSL:5

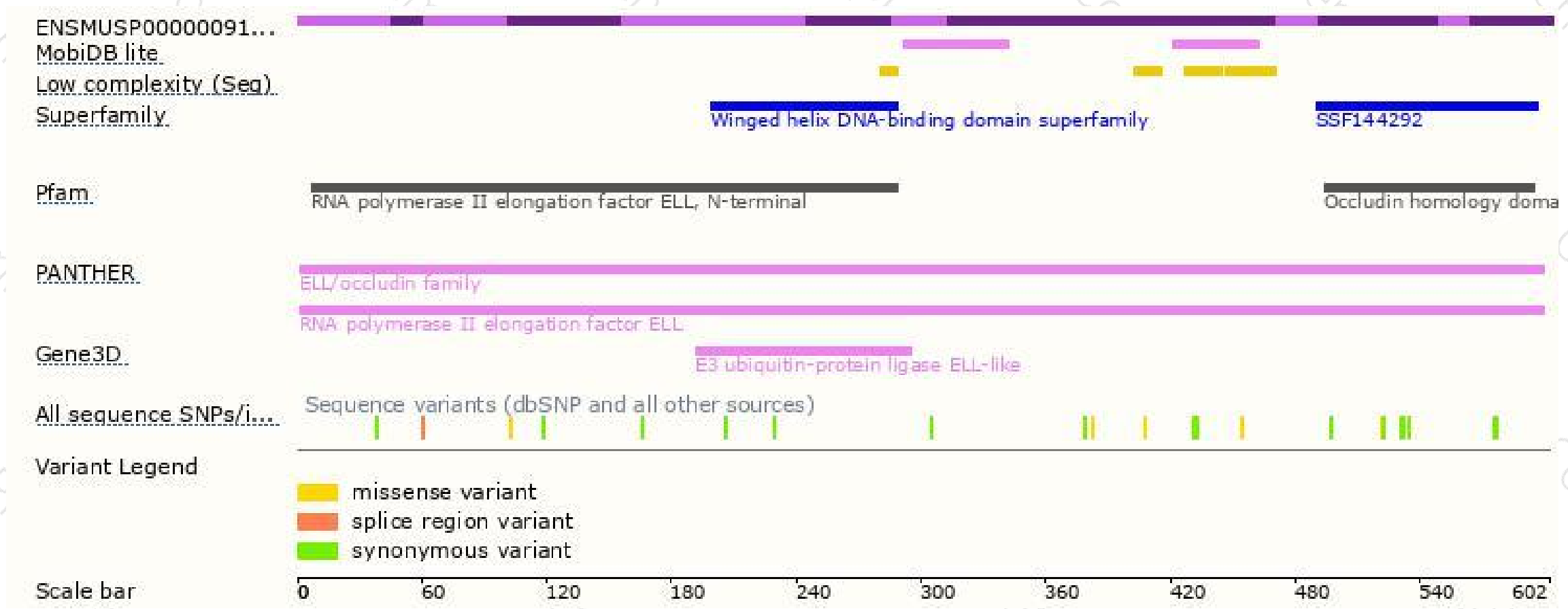
The strategy is based on the design of *EII-201* transcript,The transcription is shown below



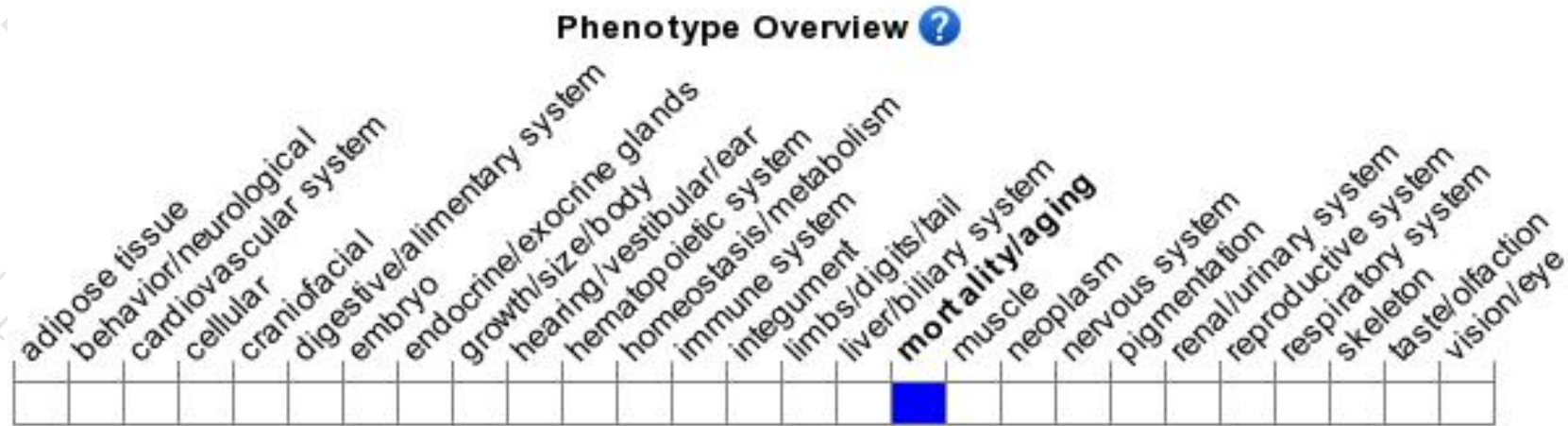
集萃药康
GemPharmatech



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 knock-out allele die prior to E6.5 but after implantation.

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

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