Fezf1 Cas9-CKO Strategy

Designer: Yanhua Shen
Reviewer: Xueting Zhang
Design Date: 2019-11-12

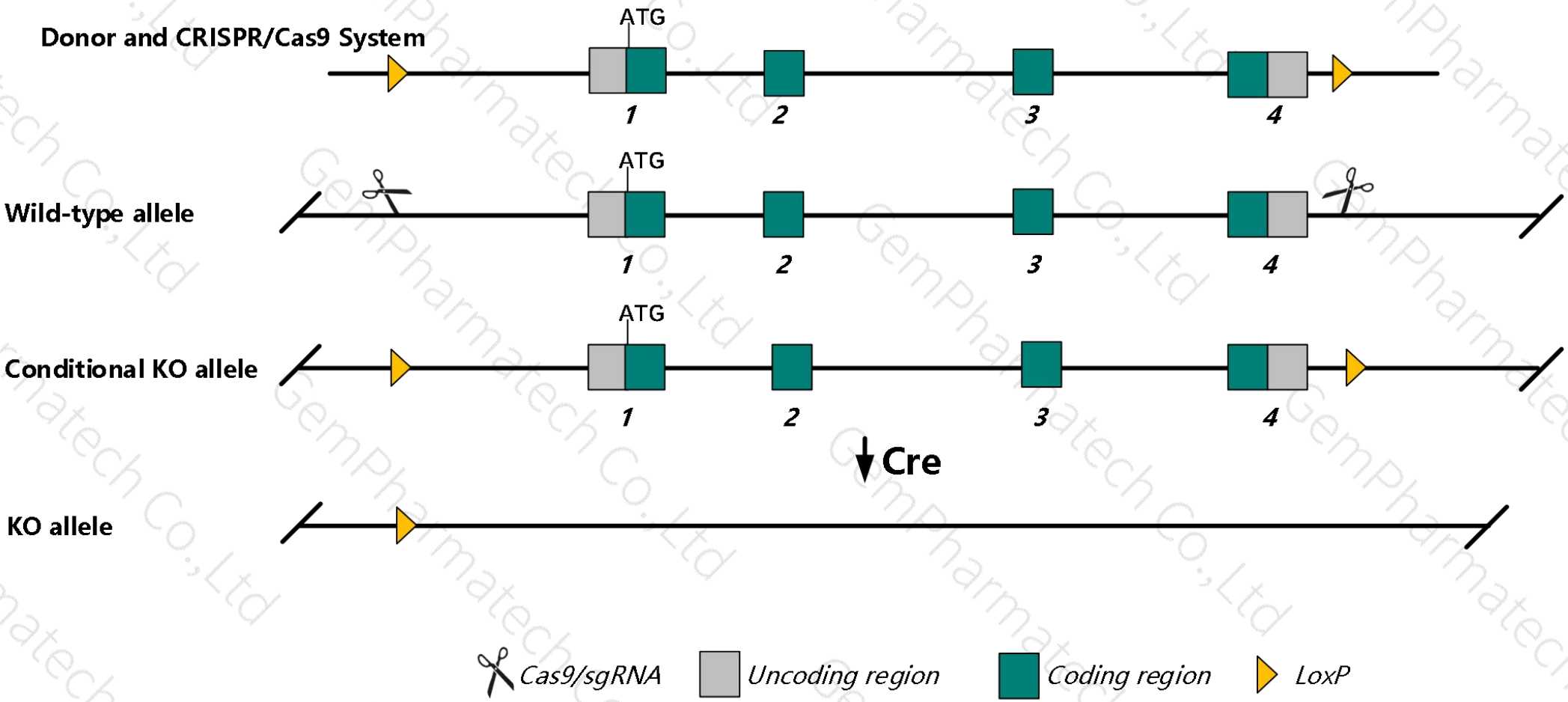
Project Overview

Project Name	***Fezf1***
Project type	**Cas9-CKO**
Strain background	**C57BL/6JGpt**

Conditional Knockout strategy

集萃药康 GemPharmatech

This model will use CRISPR/Cas9 technology to edit the *Fezf1* gene. The schematic diagram is as follows:



Technical routes

- The *Fezf1* gene has 1 transcript. According to the structure of *Fezf1* gene, exon1-exon4 of *Fezf1-201* (ENSMUST00000031709.6) transcript is recommended as the knockout region.The region contains all of coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Fezf1* 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 null mutation of this gene display neonatal lethality, impaired olfactory bulb development and impaired olfactory bulb interneuron migration.

➢ The *Fezf1* gene is located on the Chr6. 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

Fezf1 Fez family zinc finger 1 [*Mus musculus* (house mouse)]

Gene ID: 73191, updated on 10-Oct-2019

Summary

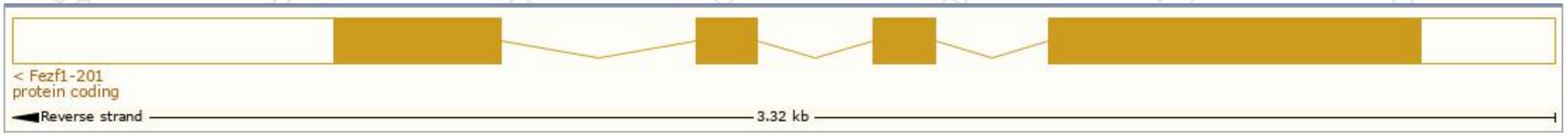
Official Symbol	Fezf1 provided by MGI
Official Full Name	Fez family zinc finger 1 provided by MGI
Primary source	MGI:MGI:1920441
See related	Ensembl:ENSMUSG00000029697
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	Fez; fez-like; Zfp312-like; 3110069A13Rik
Expression	Biased expression in whole brain E14.5 (RPKM 3.0), CNS E11.5 (RPKM 2.7) and 2 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

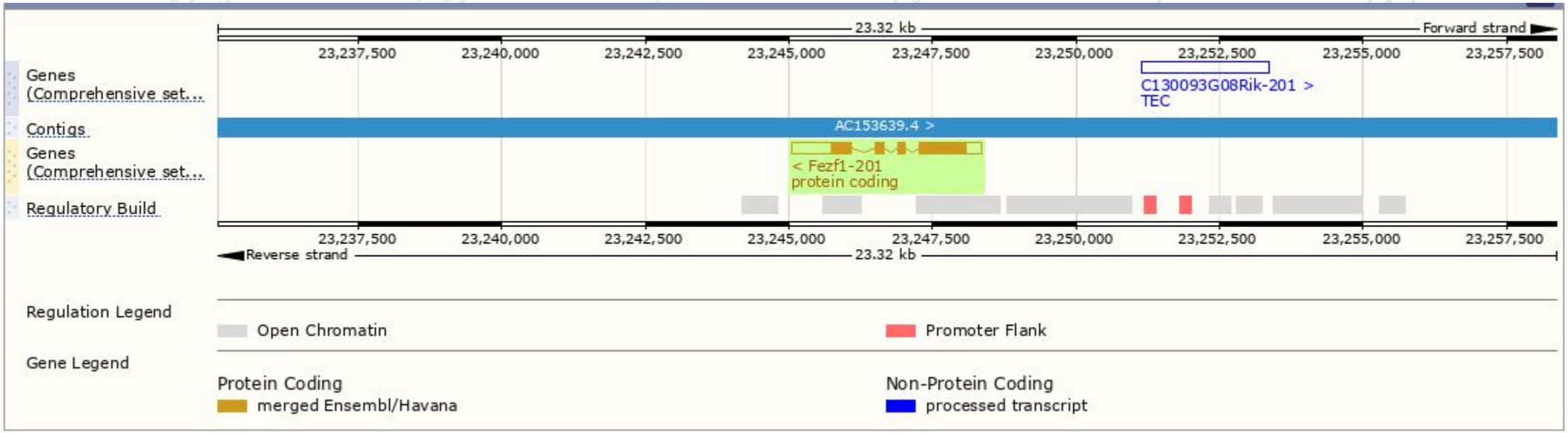
The gene has 1 transcript,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fezf1-201	ENSMUST00000031709.6	2409	475aa	Protein coding	CCDS51725	Q0VDQ9	TSL:1 GENCODE basic APPRIS P1

The strategy is based on the design of *Fezf1-201* transcript,The transcription is shown below

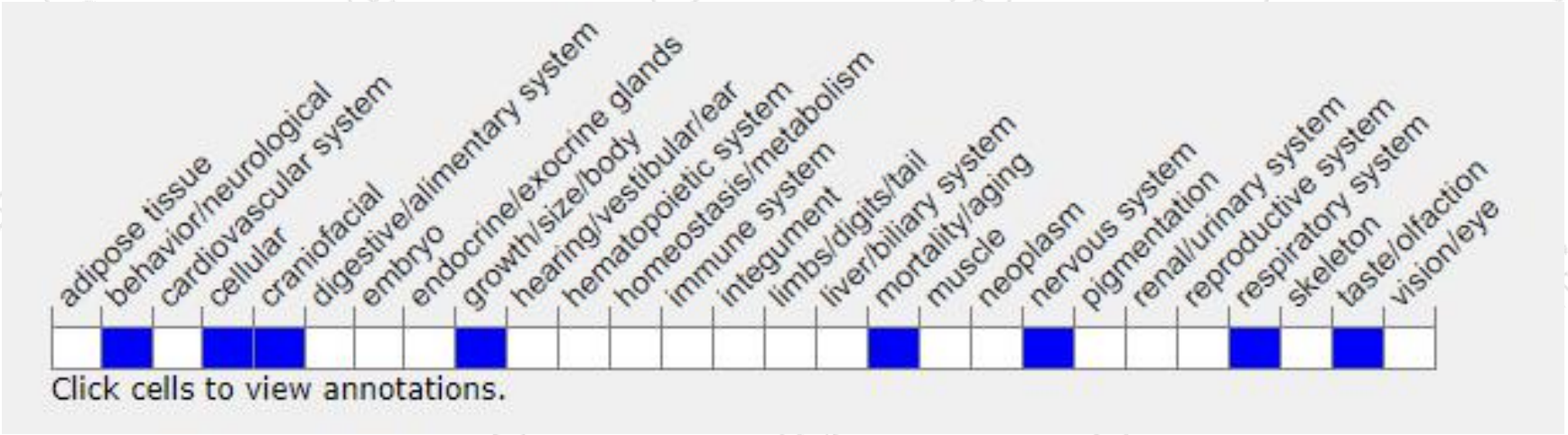


Genomic location distribution



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,Mice homozygous for a null mutation of this gene display neonatal lethality, impaired olfactory bulb development and impaired olfactory bulb interneuron migration.

集萃药康
GemPharmatech

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

Tel: 400-9660890

集萃药康生物科技
GemPharmatech Co.,Ltd