



Odf1 Cas9-CKO Strategy

Designer: Jia Yu

Reviewer: Xiaojing Li

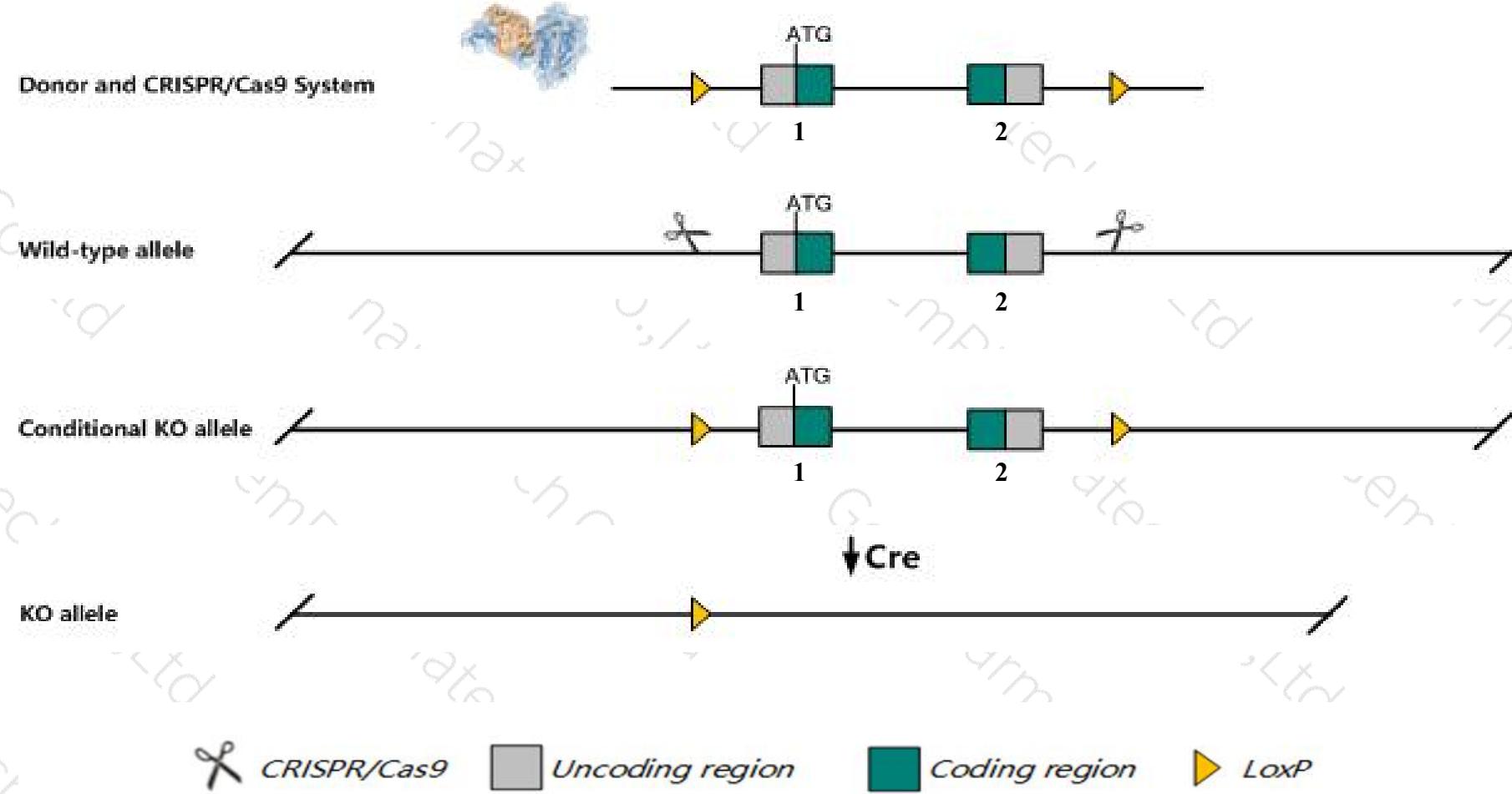
Design Date: 2020-8-1

Project Overview

Project Name	<i>Odf1</i>
Project type	Cas9-CKO
Strain background	C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Odf1* gene. The schematic diagram is as follows:



Technical routes

- The *Odf1* gene has 1 transcript. According to the structure of *Odf1* gene, exon1-exon2 of *Odf1*-201(ENSMUST00000081966.4) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Odf1* 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 exhibit male infertility associated with asthenozoospermia, oligozoospermia, detached sperm head and absent acrosome reaction. Mice heterozygous for this allele exhibit asthenozoospermia without altered fertility.
- The *Odf1* gene is located on the Chr15. 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)

Odf1 outer dense fiber of sperm tails 1 [Mus musculus (house mouse)]

Gene ID: 18285, updated on 13-Mar-2020

Summary



Official Symbol Odf1 provided by [MGI](#)

Official Full Name outer dense fiber of sperm tails 1 provided by [MGI](#)

Primary source [MGI:MG1:97424](#)

See related [Ensembl:ENSMUSG00000061923](#)

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

Expression Restricted expression toward testis adult (RPKM 1293.5)[See more](#)

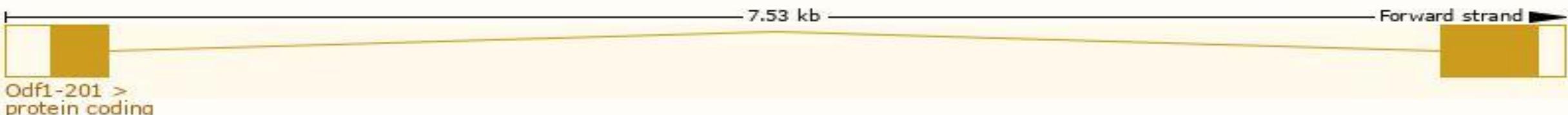
Orthologs [human](#) [all](#)

Transcript information (Ensembl)

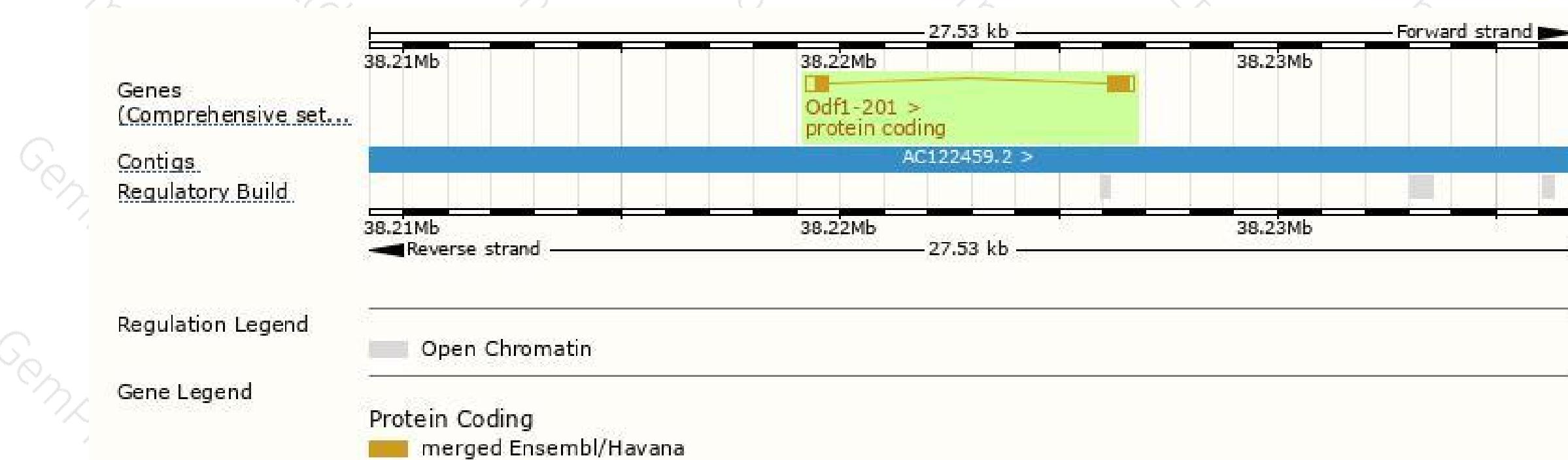
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Odf1-201	ENSMUST00000081966.4	1105	248aa	Protein coding	CCDS49596	A6H656	TSL:1 GENCODE basic APPRIS P1

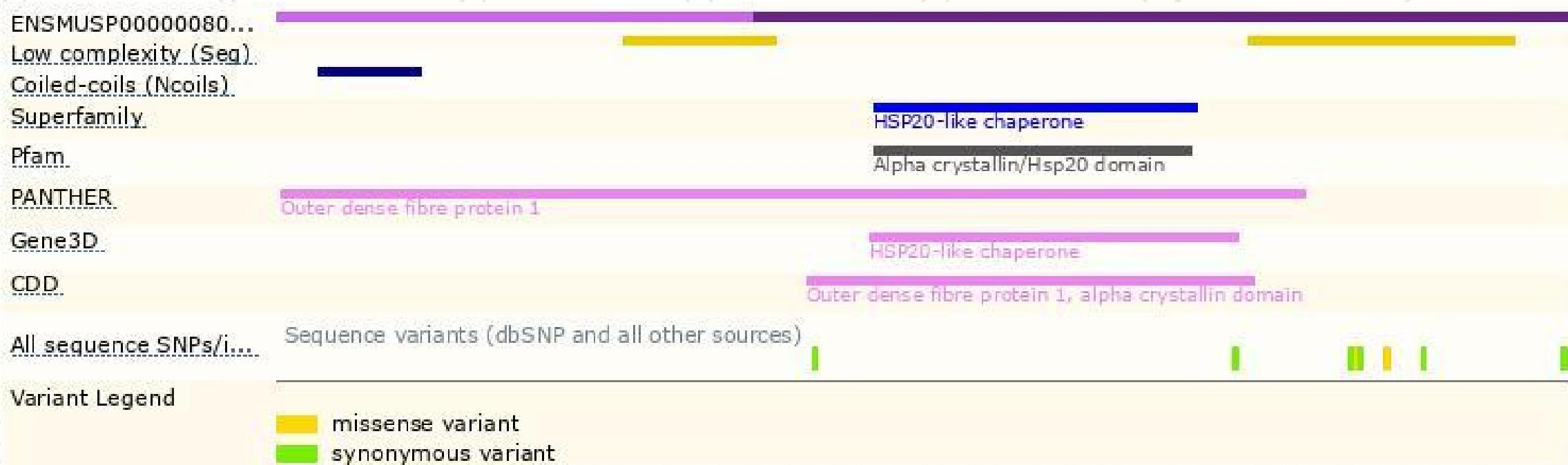
The strategy is based on the design of *Odf1-201* transcript, the transcription is shown below:



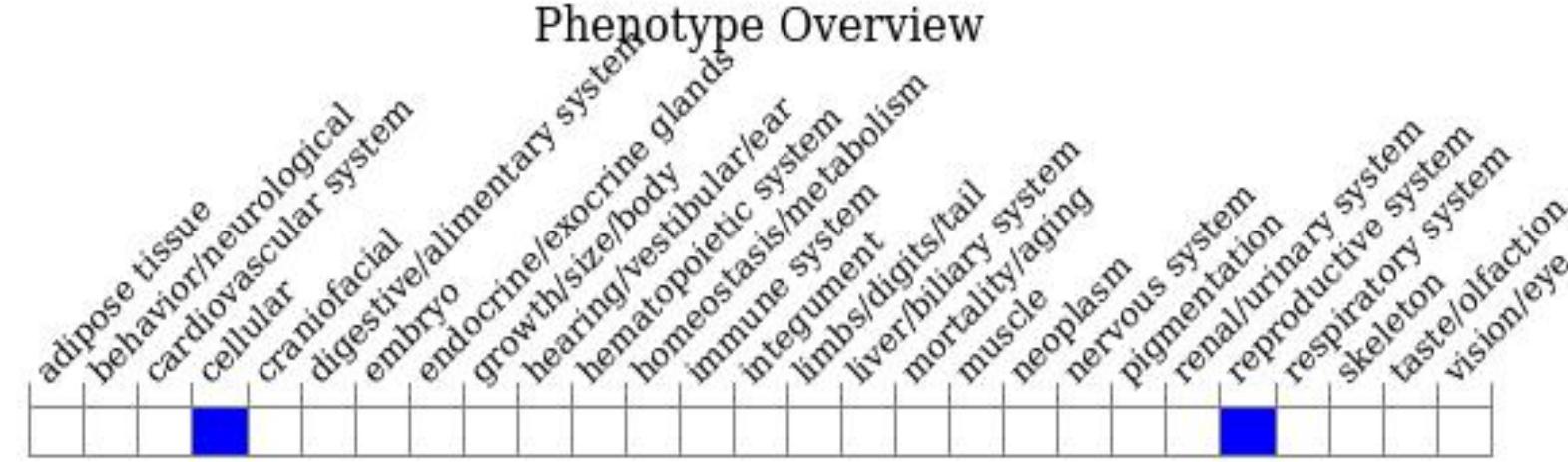
Genomic location distribution



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 knock-out allele exhibit male infertility associated with asthenozoospermia, oligozoospermia, detached sperm head and absent acrosome reaction. Mice heterozygous for this allele exhibit asthenozoospermia without altered fertility.



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

Tel: 400-9660890



集萃药康生物科技
GemPharmatech Co.,Ltd

