

Adam30 Cas9-CKO Strategy

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Design Date: 2020-9-14

Project Overview

Project Name

Adam30

Project type

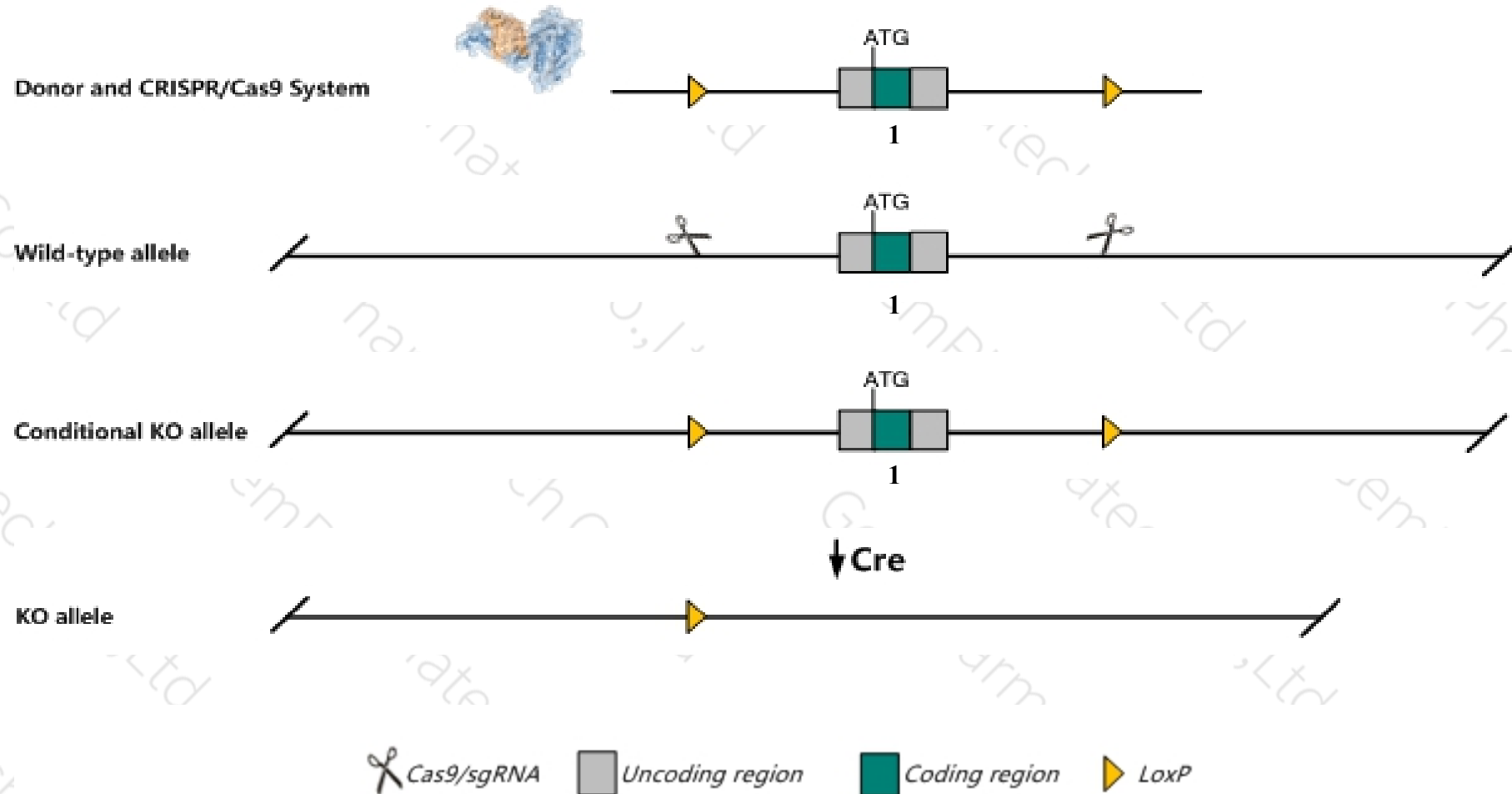
Cas9-CKO

Strain background

C57BL/6JGpt

Conditional Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Adam30* gene. The schematic diagram is as follows:



- The *Adam30* gene has 2 transcripts. According to the structure of *Adam30* gene, exon1 of *Adam30-201*(ENSMUST00000050342.4) 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 *Adam30* gene. The brief process is as follows: sgRNA was transcribed in vitro, donor vector was constructed. Cas9, sgRNA 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 was 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.

- The *Adam30* gene is located on the Chr3. 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)

Adam30 a disintegrin and metallopeptidase domain 30 [Mus musculus (house mouse)]

Gene ID: 71078, updated on 13-Mar-2020

Summary



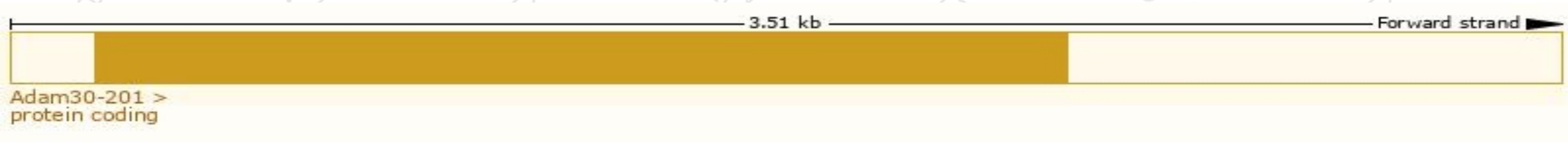
Official Symbol	Adam30 provided by MGI
Official Full Name	a disintegrin and metallopeptidase domain 30 provided by MGI
Primary source	MGI:MGI:1918328
See related	Ensembl:ENSMUSG00000043468
Gene type	protein coding
RefSeq status	VALIDATED
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	4933424D07Rik, svph4
Orthologs	human all

Transcript information (Ensembl)

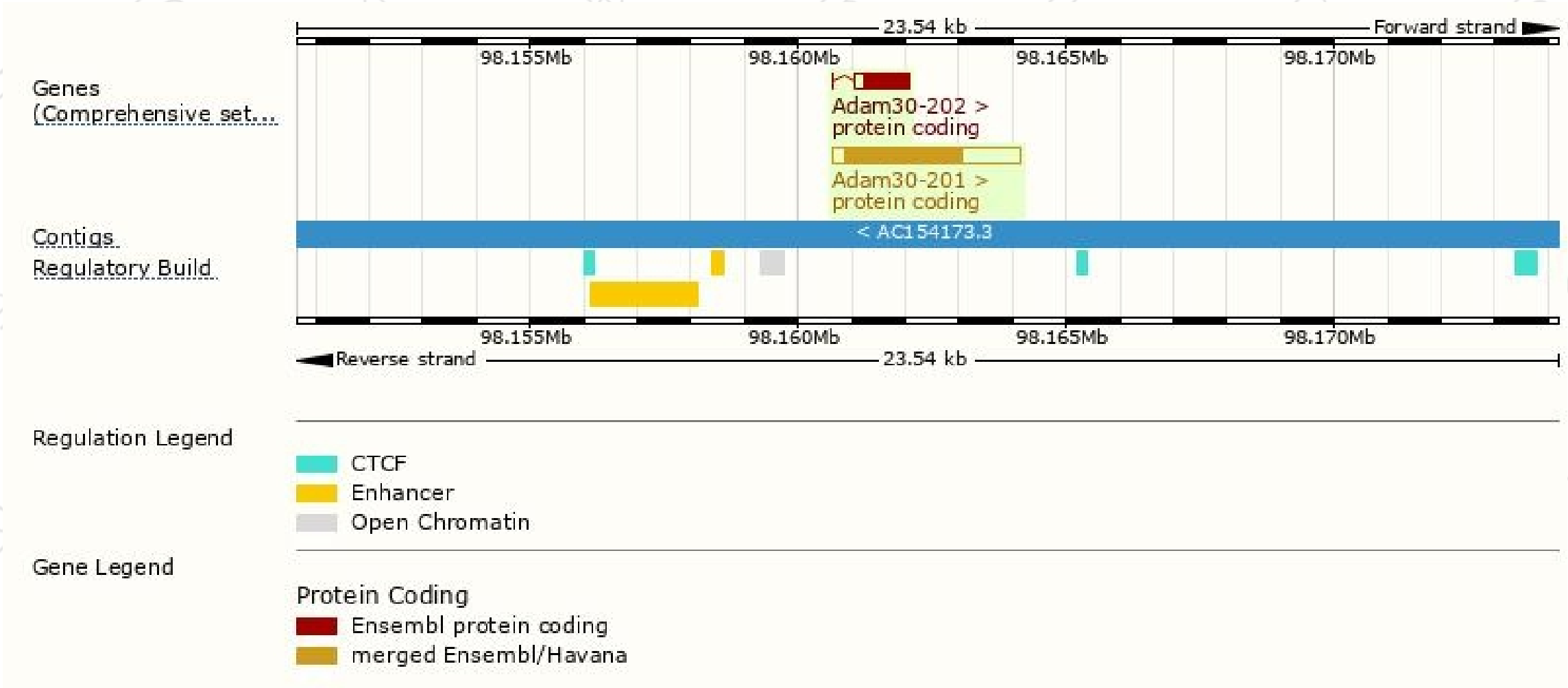
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Adam30-201	ENSMUST00000050342.4	3510	732aa	Protein coding	CCDS17660	Q811Q3	TSL:NA GENCODE basic APPRIS P1
Adam30-202	ENSMUST00000198363.1	1042	278aa	Protein coding	-	Q9CUC4	CDS 3' incomplete TSL:1

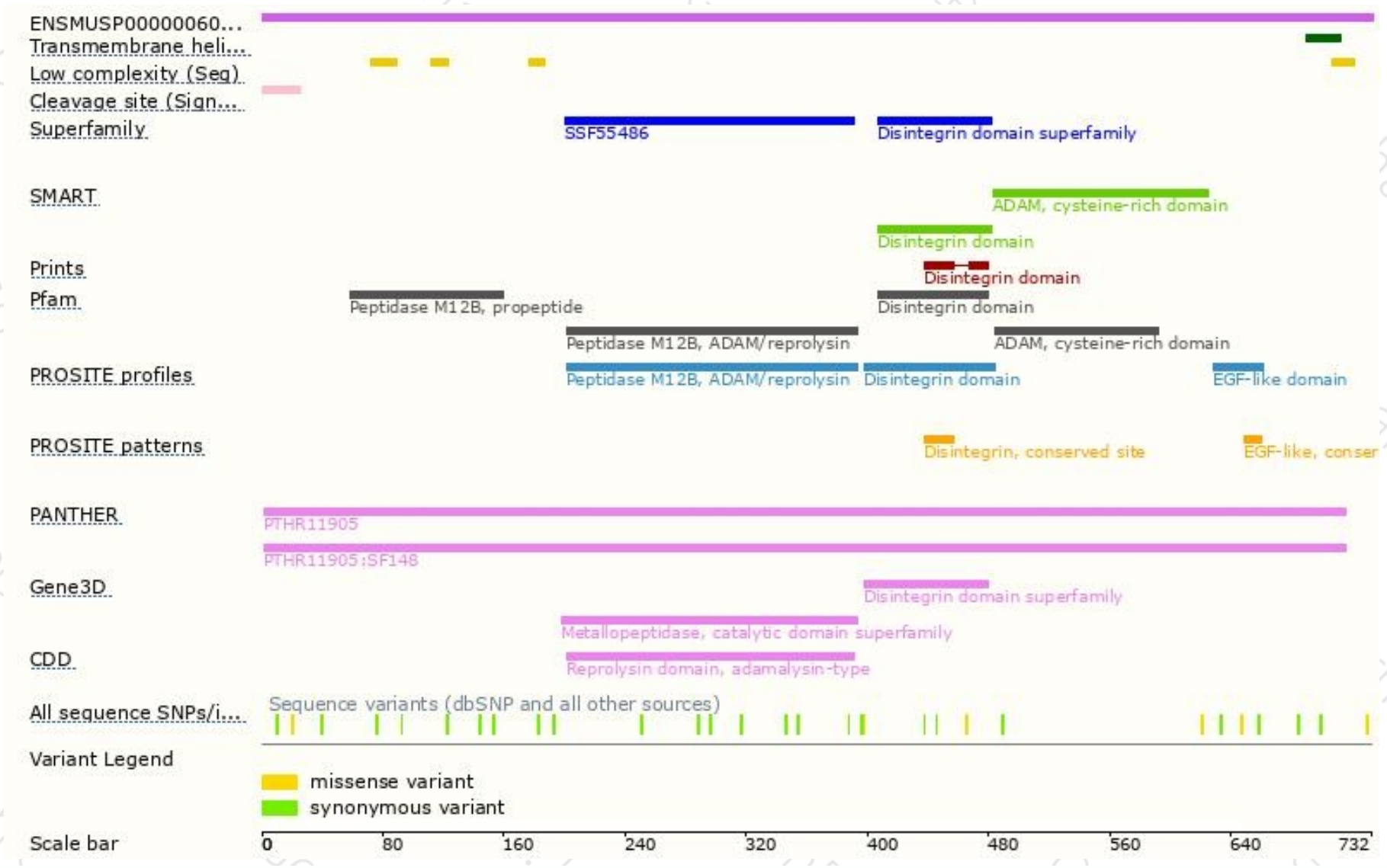
The strategy is based on the design of *Adam30-201* transcript,the transcription is shown below:



Genomic location distribution



Protein domain



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

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