

Exoc6 Cas9-KO Strategy

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Project Overview

Project Name

Exoc6

Project type

Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Exoc6* gene. The schematic diagram is as follows:



- The *Exoc6* gene has 1 transcript. According to the structure of *Exoc6* gene, exon2-exon7 of *Exoc6-201* (ENSMUST00000066439.7) transcript is recommended as the knockout region. The region contains 718bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Exoc6* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Homozygotes for a spontaneous mutation exhibit severe microcytic anemia, erythrocyte hyperchromia, and markedly increased levels of red cell protoporphyrin.
- The *Exoc6* gene is located on the Chr19. 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Exoc6 exocyst complex component 6 [Mus musculus (house mouse)]

Gene ID: 107371, updated on 31-Jan-2019

Summary



Official Symbol	Exoc6 provided by MGI
Official Full Name	exocyst complex component 6 provided by MGI
Primary source	MGI:MGI:1351611
See related	Ensembl:ENSMUSG00000053799
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	4833405E05Rik, AW413330, C430002C19, Sec15, Sec15l1, hbd, msec15
Expression	Ubiquitous expression in placenta adult (RPKM 12.6), liver E14 (RPKM 11.8) and 28 other tissues See more
Orthologs	human all

Transcript information (Ensembl)

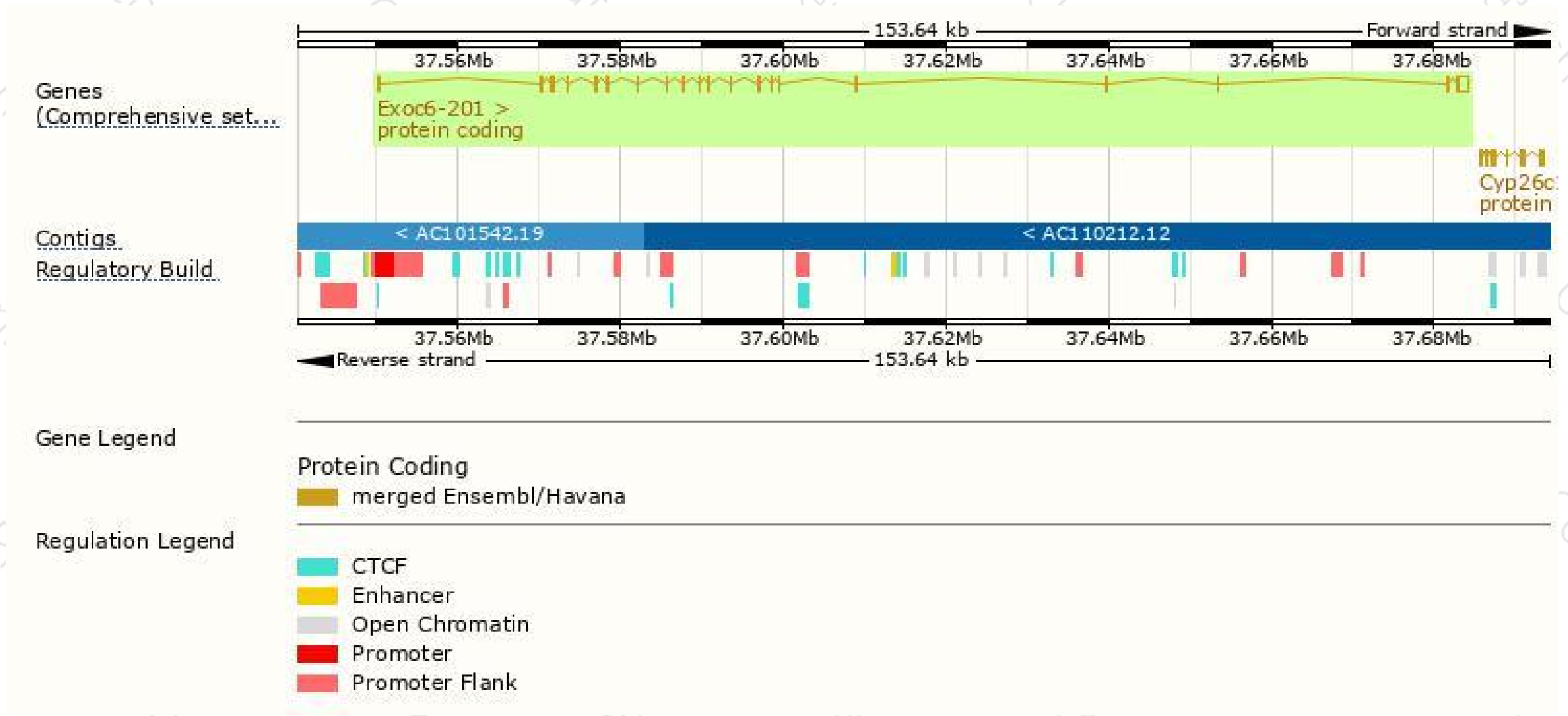
The gene has 1 transcript, and the transcript is shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Exoc6-201	ENSMUST00000066439.7	3543	804aa	Protein coding	CCDS29779	Q3U9D6	TSL:1 GENCODE basic APPRIS P1

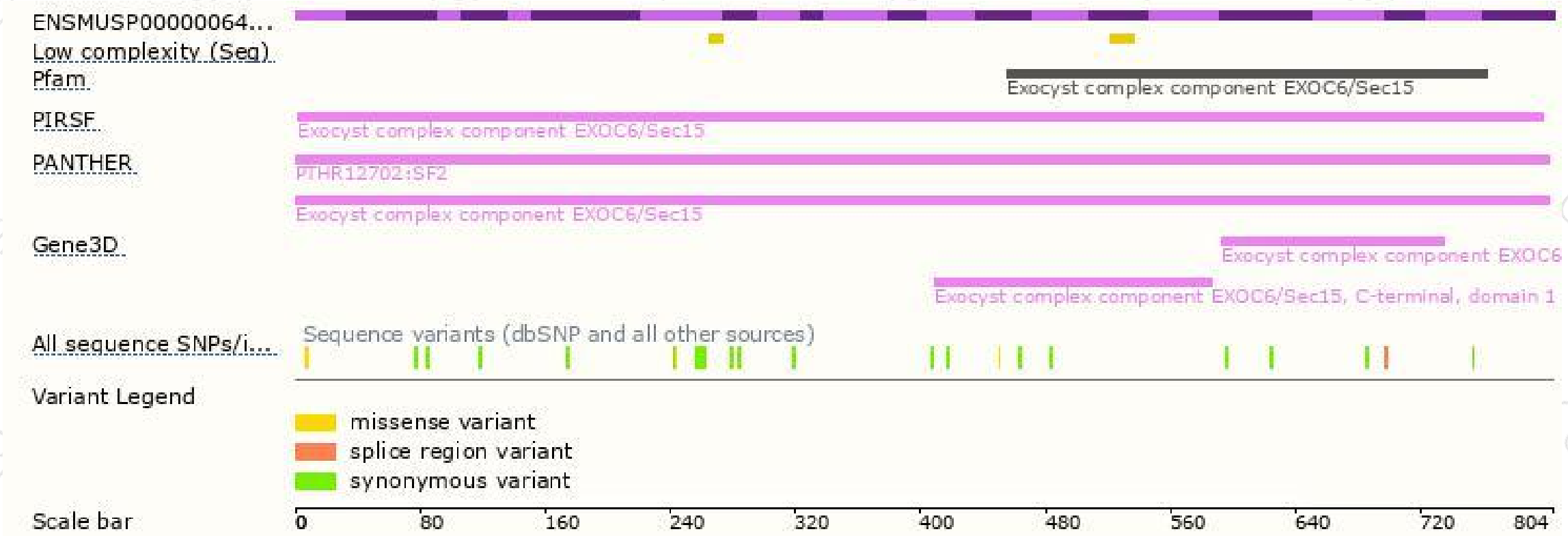
The strategy is based on the design of *Exoc6-201* transcript, The transcription is shown below



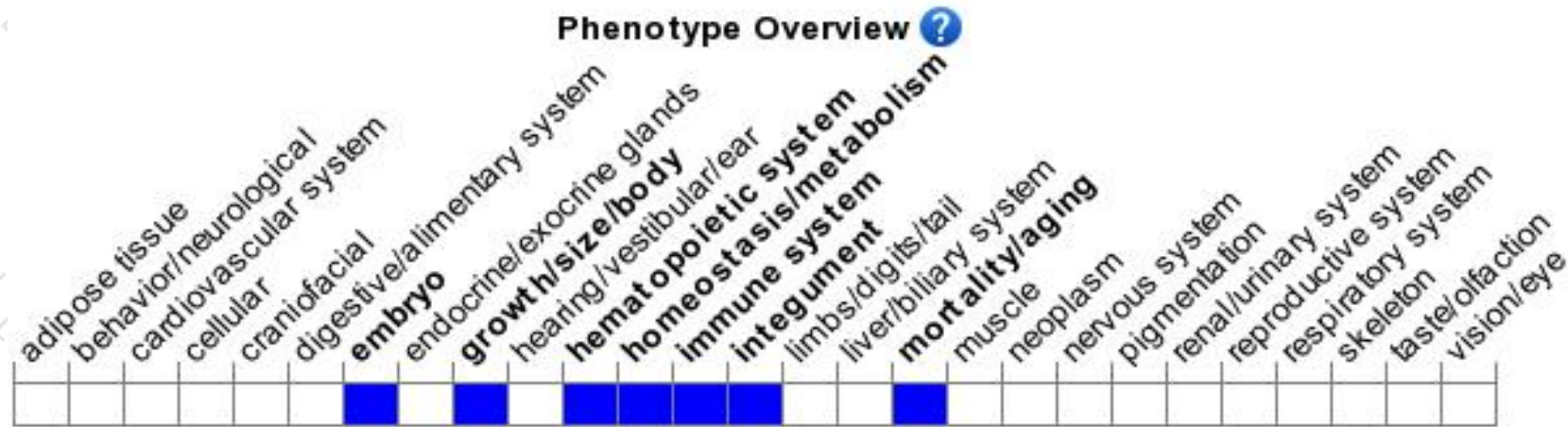
Genomic location distribution



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, Homozygotes for a spontaneous mutation exhibit severe microcytic anemia, erythrocyte hyperchromia, and markedly increased levels of red cell protoporphyrin.

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

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