

# *Ccng1* Cas9-CKO Strategy

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# Project Overview

**Project Name**

*Ccng1*

**Project type**

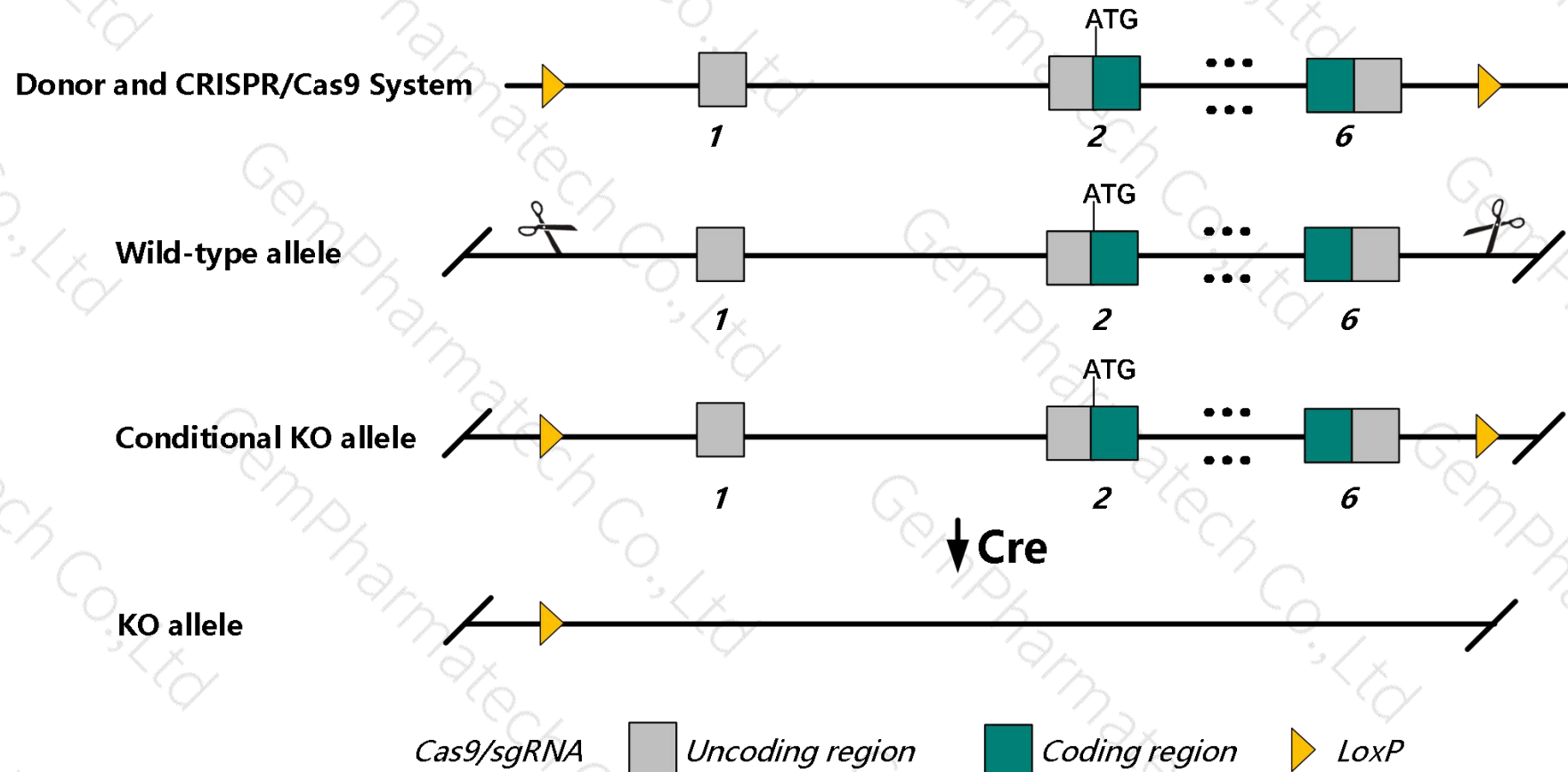
**Cas9-CKO**

**Strain background**

**C57BL/6JGpt**

# Conditional Knockout strategy

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



- The *Ccngl* gene has 2 transcripts. According to the structure of *Ccngl* gene, exon1-exon6 of *Ccngl*-201 (ENSMUST00000020576.7) transcript is recommended as the knockout region. The region contains most 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 *Ccngl* 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.

- According to the existing MGI data, Depending on the allele, homozygous mutants exhibit increased cellular sensitivity to gamma-irradiation or decreased incidence of induced hepatic tumors.
- The *Ccng1* gene is located on the Chr11. 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)

## Ccng1 cyclin G1 [ *Mus musculus* (house mouse) ]

Gene ID: 12450, updated on 12-Nov-2019

### Summary

Official Symbol	Ccng1 provided by MGI
Official Full Name	cyclin G1 provided by MGI
Primary source	MGI:MGI:102890
See related	Ensembl:ENSMUSG00000020326
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<i>Mus musculus</i>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	AI314029
Expression	Broad expression in heart adult (RPKM 67.7), bladder adult (RPKM 21.1) and 22 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

### Genomic context

Location: 11; 11 A5

See Ccng1 in [Genome Data Viewer](#)

Exon count: 6

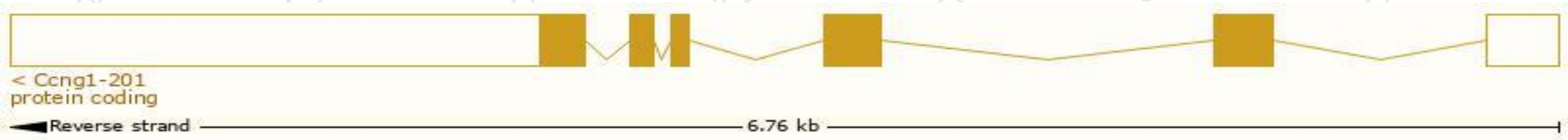
Annotation release	Status	Assembly	Chr	Location
<a href="#">108</a>	current	GRCm38.p6 ( <a href="#">GCF_000001635.26</a> )	11	NC_000077.6 (40748552..40755309, complement)
Build 37.2	previous assembly	MGSCv37 ( <a href="#">GCF_000001635.18</a> )	11	NC_000077.5 (40562054..40568788, complement)

# Transcript information (Ensembl)

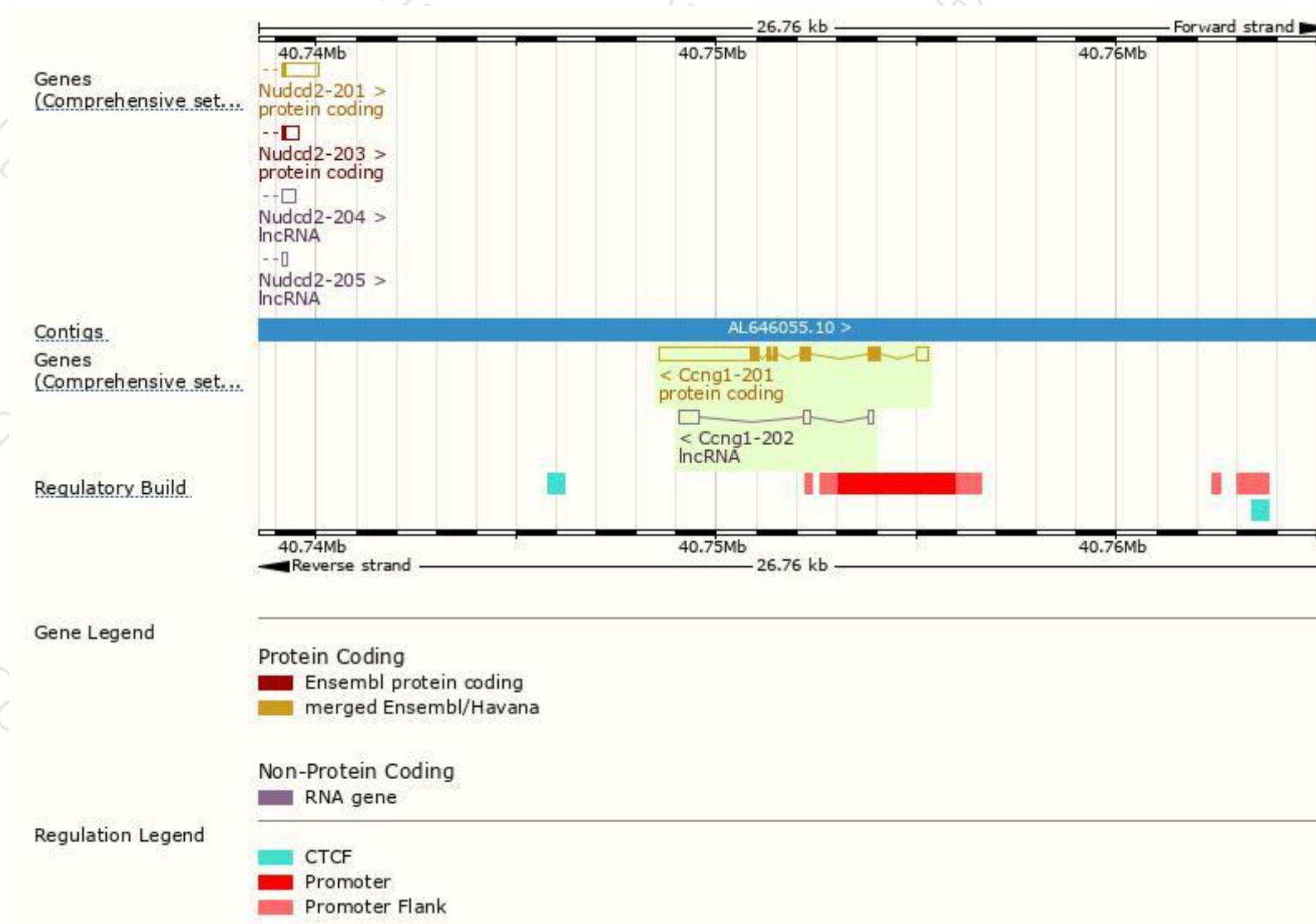
The gene has 2 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Ccng1-201	<a href="#">ENSMUST00000020576.7</a>	3512	<a href="#">294aa</a>	Protein coding	<a href="#">CCDS24550</a>	<a href="#">P51945 Q5NC86</a>	TSL:1 GENCODE basic APPRIS P1
Ccng1-202	<a href="#">ENSMUST00000151359.1</a>	804	No protein	lncRNA	-	-	TSL:5

The strategy is based on the design of *Ccng1-201* transcript,The transcription is shown below

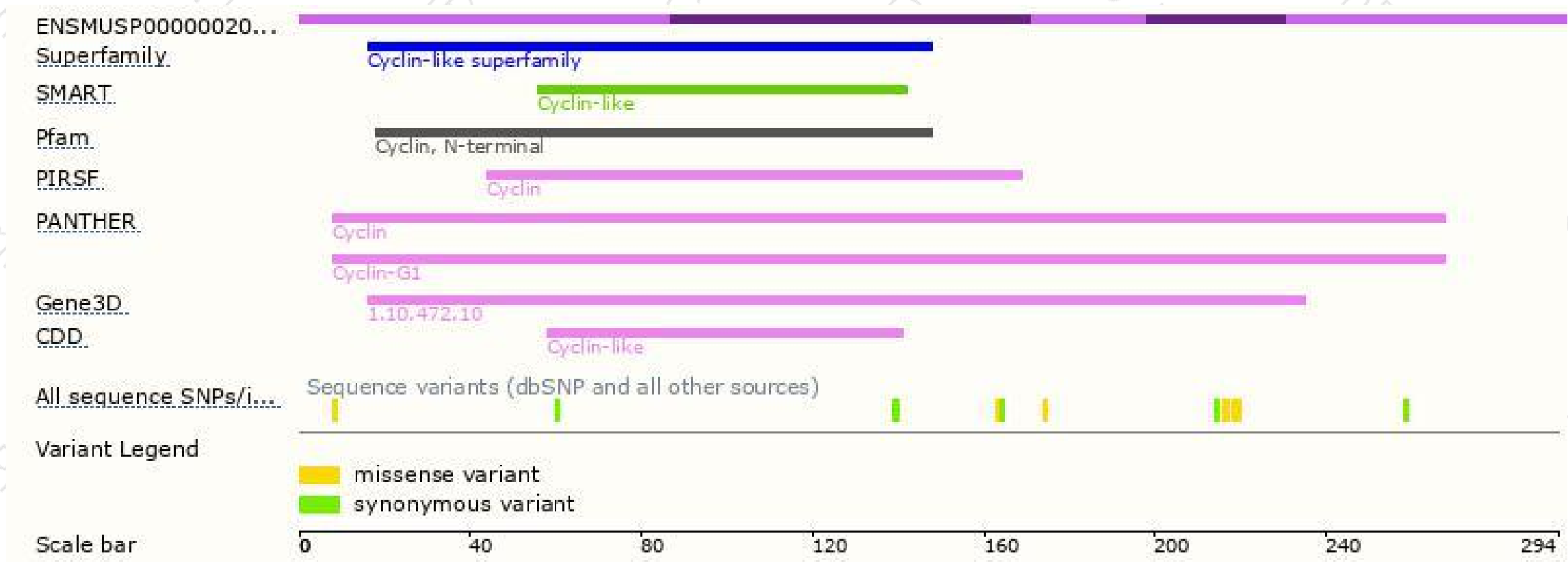


# Genomic location distribution





# 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, Depending on the allele, homozygous mutants exhibit increased cellular sensitivity to gamma-irradiation or decreased incidence of induced hepatic tumors.

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

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