Dcps Cas9-CKO Strategy

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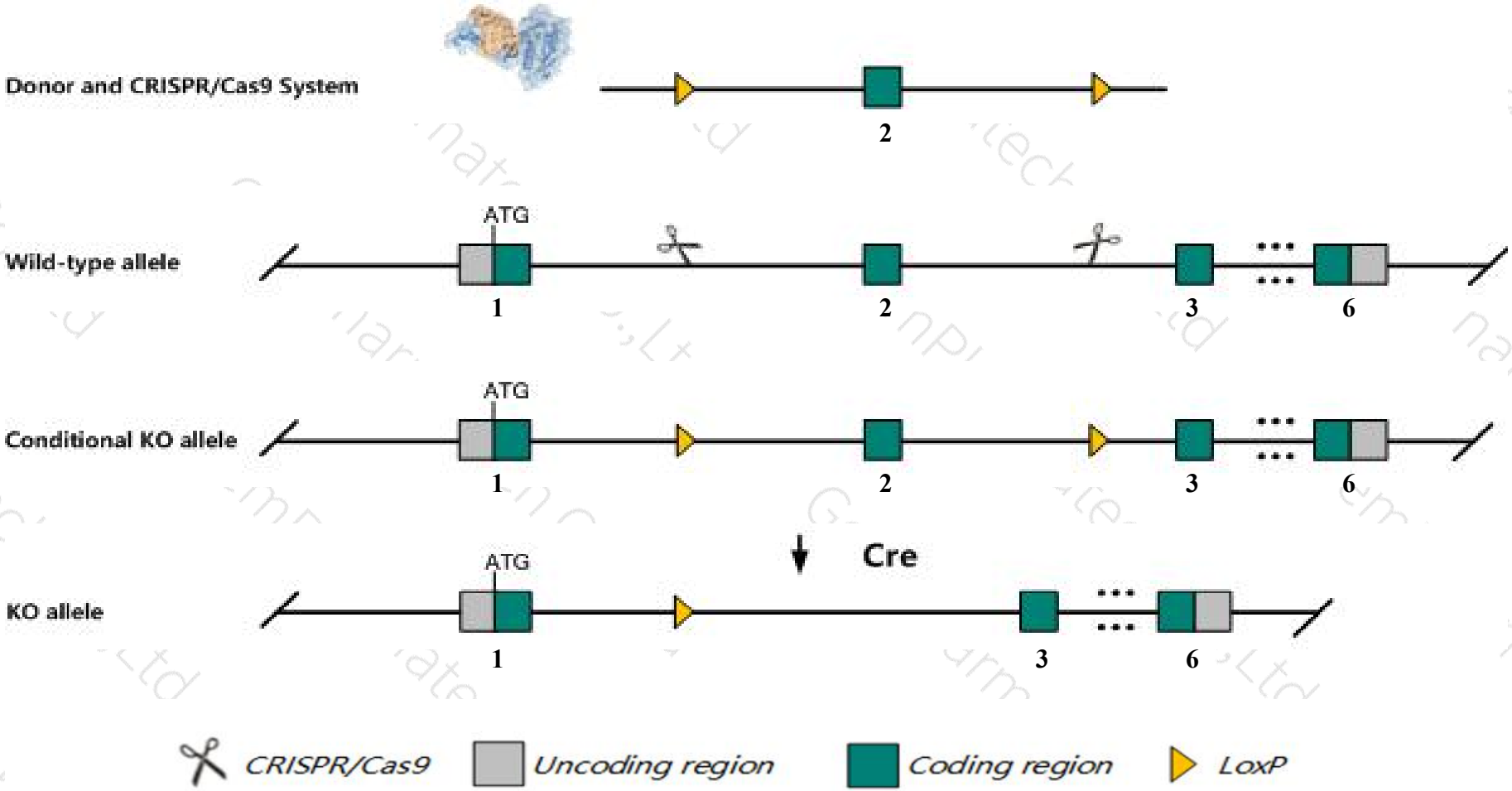
Project Overview

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

Conditional Knockout strategy

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



Technical routes

- The *Dcps* gene has 3 transcripts. According to the structure of *Dcps* gene, exon2 of *Dcps-201* (ENSMUST00000034539.11) transcript is recommended as the knockout region.The region contains 175bp coding sequence. Knock out the region will result in disruption of protein function.

- In this project we use CRISPR/Cas9 technology to modify *Dcps* 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,A mutant mouse line was generated from gene-trapped ES cells, but no phenotypic information is available.

- The *Dcps* gene is located on the Chr9. 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

Dcps decapping enzyme, scavenger [Mus musculus (house mouse)]

Gene ID: 69305, updated on 31-Jan-2019

Summary

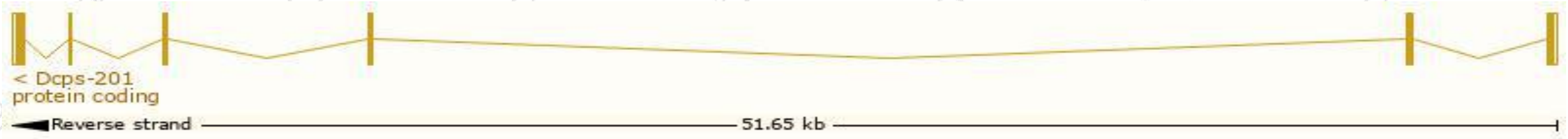
Official Symbol	Dcps provided by MGI
Official Full Name	decapping enzyme, scavenger provided by MGI
Primary source	MGI:MGI:1916555
See related	Ensembl:ENSMUSG00000032040
Gene type	protein coding
RefSeq status	PROVISIONAL
Organism	Mus musculus
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	1700001E16Rik, AA408441
Expression	Ubiquitous expression in liver E14.5 (RPKM 26.3), liver E14 (RPKM 24.3) and 28 other tissues See more
Orthologs	human all

Transcript information（Ensembl）

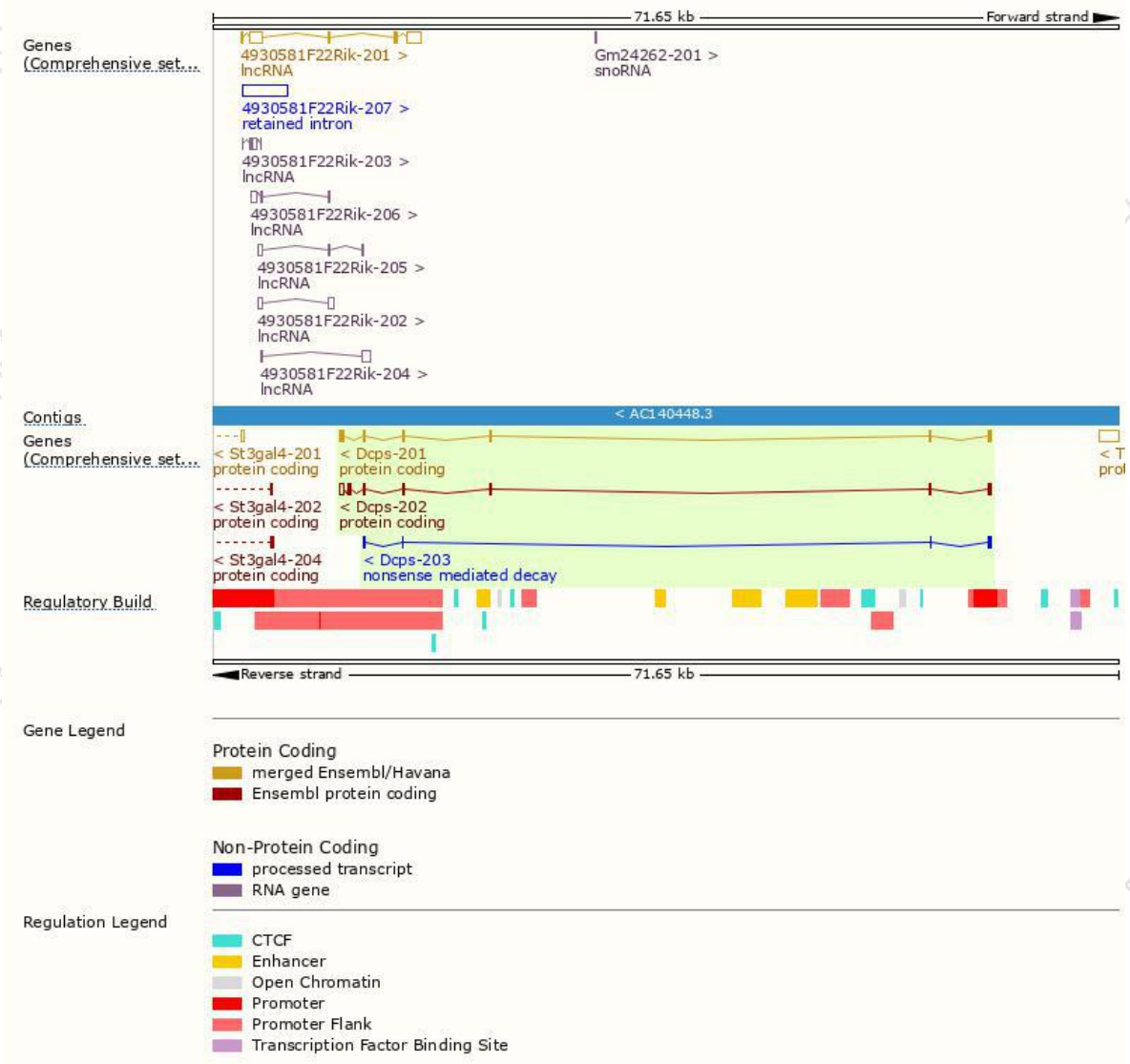
The gene has 3 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Dcps-201	ENSMUST00000034539.11	1274	338aa	Protein coding	CCDS22957	Q9DAR7	TSL:1 GENCODE basic APPRIS P1
Dcps-202	ENSMUST00000119847.1	1445	291aa	Protein coding	-	Q3TBW9	TSL:1 GENCODE basic
Dcps-203	ENSMUST00000155139.1	426	102aa	Nonsense mediated decay	-	D6RFQ0	TSL:2

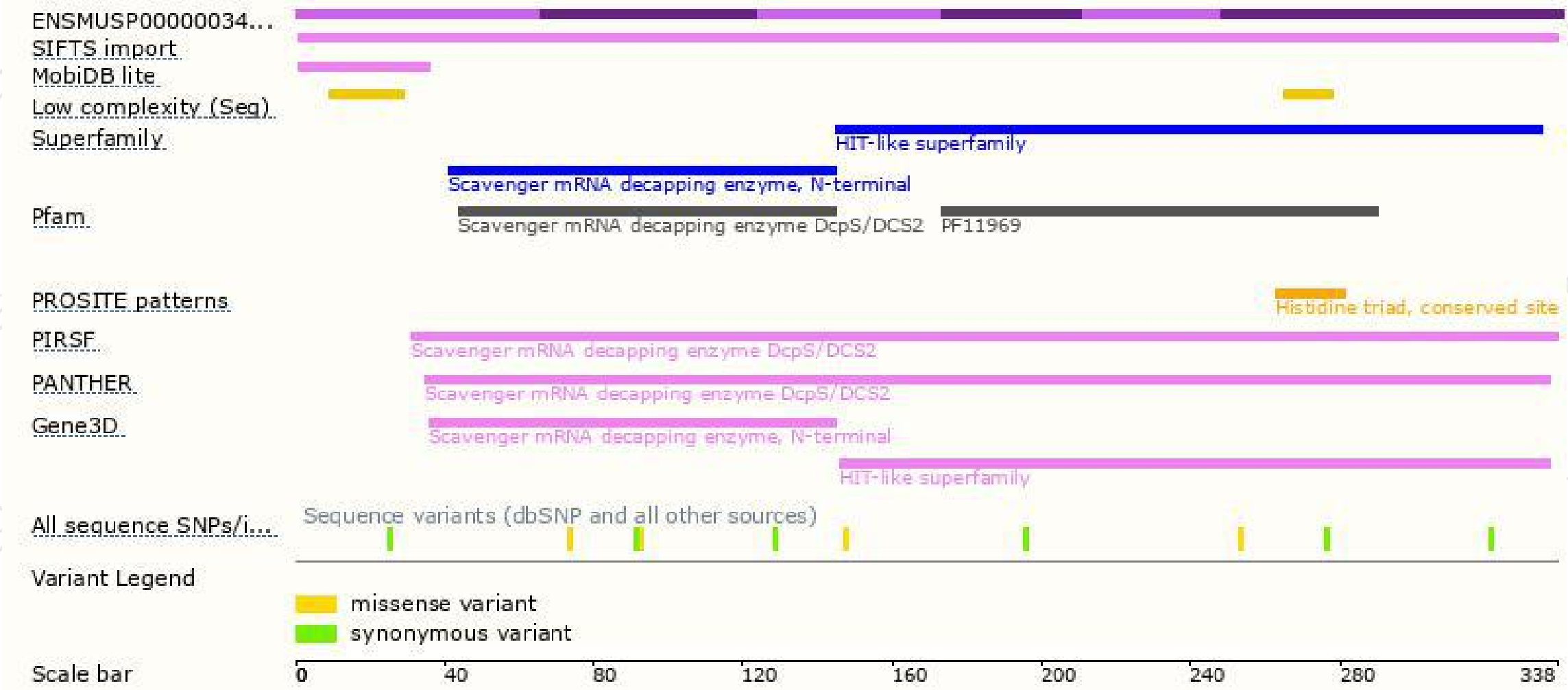
The strategy is based on the design of *Dcps-201* transcript,The transcription is shown below



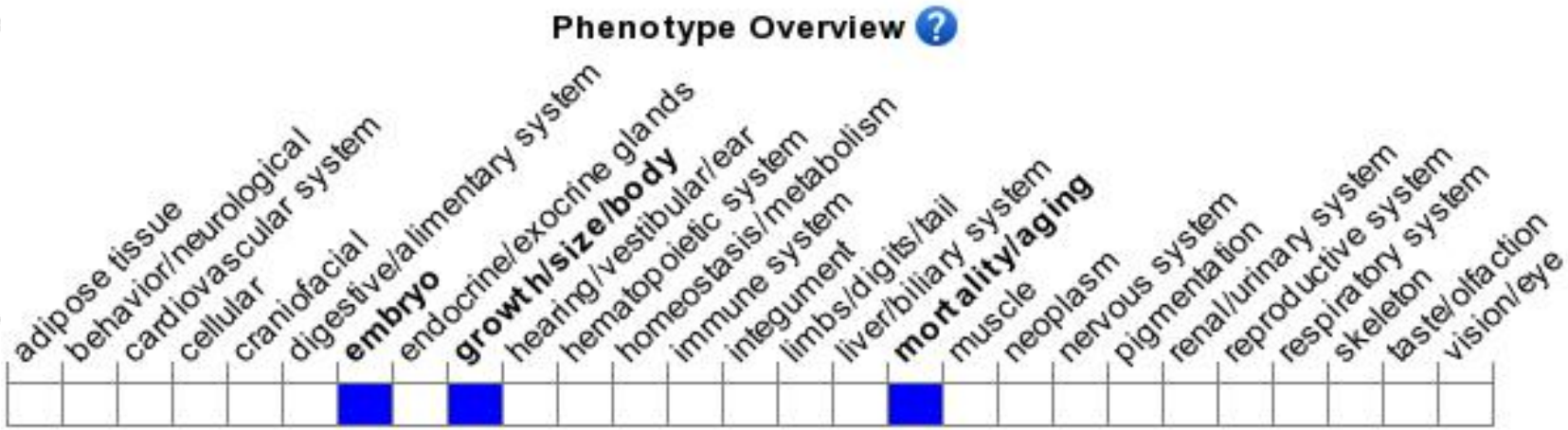
Genomic location distribution



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/).

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

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