

# Bmp6 Cas9-CKO Strategy

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**Design Date: 2018-11-9** 

## **Project Overview**



Project Name Bmp6

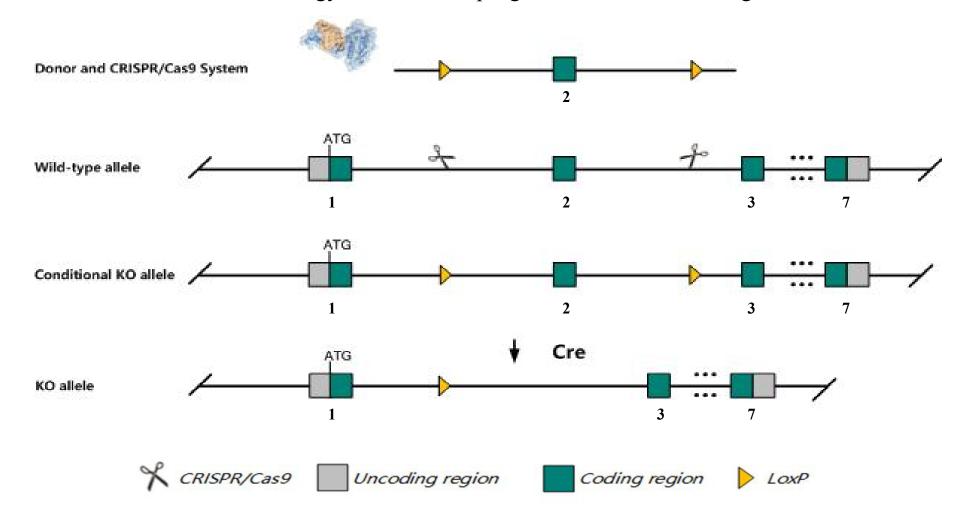
Project type Cas9-CKO

Strain background C57BL/6JGpt

### Conditional Knockout strategy



This model will use CRISPR/Cas9 technology to edit the *Bmp6* gene. The schematic diagram is as follows:



#### **Technical routes**



The *Bmp6* gene has 3 transcripts. According to the structure of *Bmp6* gene, exon2 of *Bmp6-201*(ENSMUST00000171970.2) transcript is recommended as the knockout region. The region contains 193bp coding sequence. Knock out the region will result in disruption of protein function.

In this project we use CRISPR/Cas9 technology to modify *Bmp6* 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, one homozygous null mutant showed delayed ossification in the developing sternum while females of a second null mutant were smaller than normal in size.

The *Bmp6* gene is located on the Chr13. 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



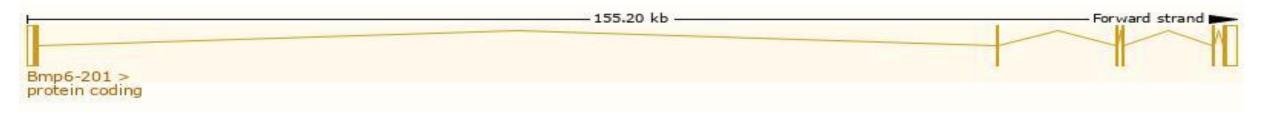
### Transcript information Ensembl



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

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Bmp6-201	ENSMUST00000171970.2	3593	<u>510aa</u>	Protein coding	CCDS26462	P20722	TSL:1 GENCODE basic APPRIS P1
Bmp6-202	ENSMUST00000223628.1	594	<u>71aa</u>	Protein coding	===	A0A286YCE8	CDS 3' incomplete
Bmp6-203	ENSMUST00000224452.1	671	No protein	Processed transcript	20	848	

The strategy is based on the design of Bmp6-201 transcript, the transcription is shown below:



### Genomic location distribution



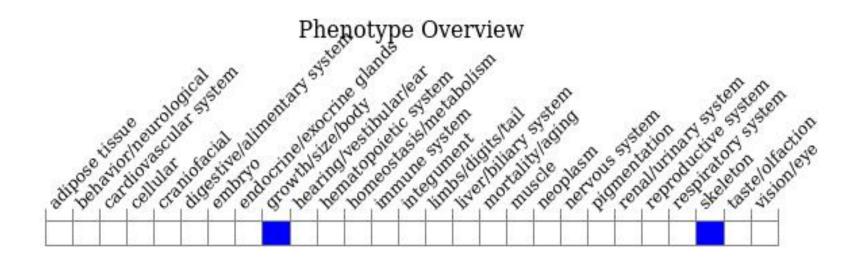
#### Protein domain





# 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, one homozygous null mutant showed delayed ossification in the developing sternum while females of a second null mutant were smaller than normal in size.



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

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