

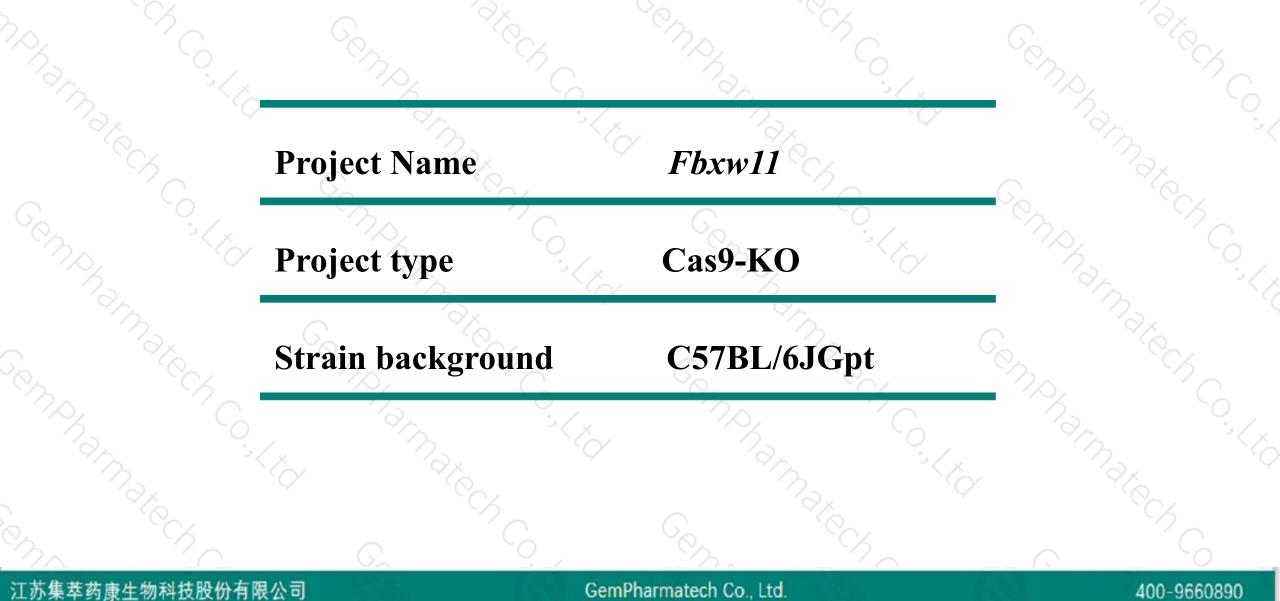
# Fbxw11 Cas9-KO Strategy

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### **Project Overview**

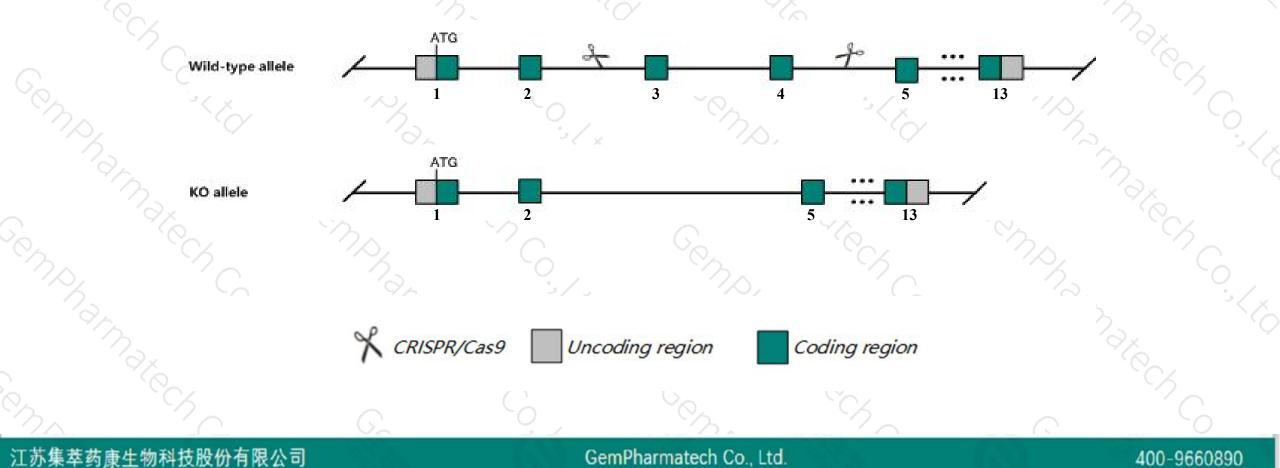




# **Knockout** strategy



This model will use CRISPR/Cas9 technology to edit the *Fbxw11* gene. The schematic diagram is as follows:





- The *Fbxw11* gene has 5 transcripts. According to the structure of *Fbxw11* gene, exon3-exon4 of *Fbxw11-201* (ENSMUST00000076383.7) transcript is recommended as the knockout region. The region contains 289bp coding sequence. Knock out the region will result in disruption of protein function.
- > In this project we use CRISPR/Cas9 technology to modify Fbxw11 gene. The brief process is as follows: CRISPR/Cas9 system

- The *Fbxw11* 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 the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Notice

# **Gene information (NCBI)**



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### Fbxw11 F-box and WD-40 domain protein 11 [Mus musculus (house mouse)]

Gene ID: 103583, updated on 3-Feb-2019

#### Summary

Official Symbol	Fbxw11 provided by MGI
Official Full Name	F-box and WD-40 domain protein 11 provided by MGI
<b>Primary source</b>	MGI:MGI:2144023
See related	Ensembl:ENSMUSG00000020271
Gene type	protein coding
<b>RefSeq status</b>	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	2310065A07Rik, AA536858, BTRC2, BTRCP2, Fbxw1b, HOS
Expression	Ubiquitous expression in cerebellum adult (RPKM 24.5), whole brain E14.5 (RPKM 24.2) and 28 other tissues See more
Orthologs	human all

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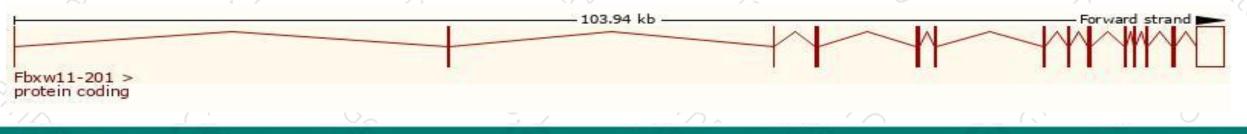
## **Transcript information (Ensembl)**



### The gene has 5 transcripts, all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Fbxw11-201	ENSMUST0000076383.7	4019	<u>563aa</u>	Protein coding	CCDS36125	Q5SRY7	TSL:1 GENCODE basic APPRIS P3
Fbxw11-202	ENSMUST0000093205.12	3938	<u>542aa</u>	Protein coding	CCDS70157	Q5SRY7	TSL:1 GENCODE basic APPRIS ALT1
Fbxw11-203	ENSMUST00000109366.7	3821	<u>529aa</u>	Protein coding	CCDS70158	Q5SRY7	TSL:1 GENCODE basic APPRIS ALT1
Fbxw11-205	ENSMUST00000143290.1	620	<u>43aa</u>	Nonsense mediated decay	<u>64</u>	F7BGU3	CDS 5' incomplete TSL:3
Fbxw11-204	ENSMUST00000137898.1	309	No protein	IncRNA	15		TSL:1

The strategy is based on the design of *Fbxw11-201* transcript, The transcription is shown below

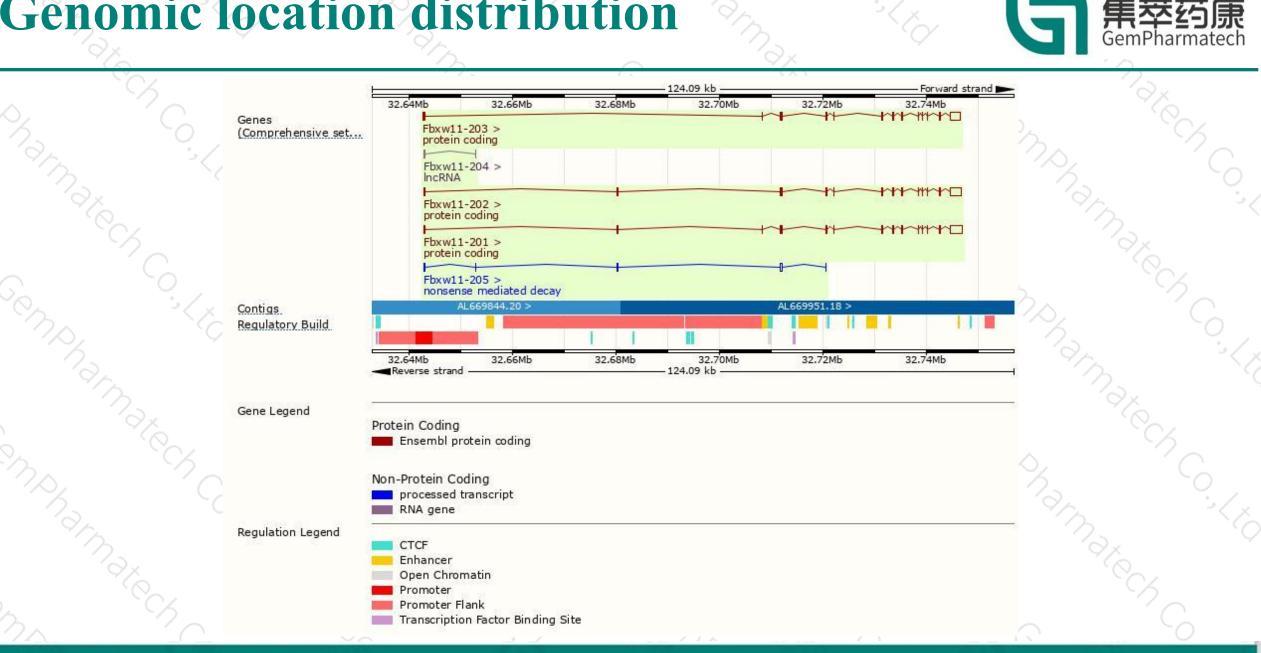


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### **Genomic location distribution**

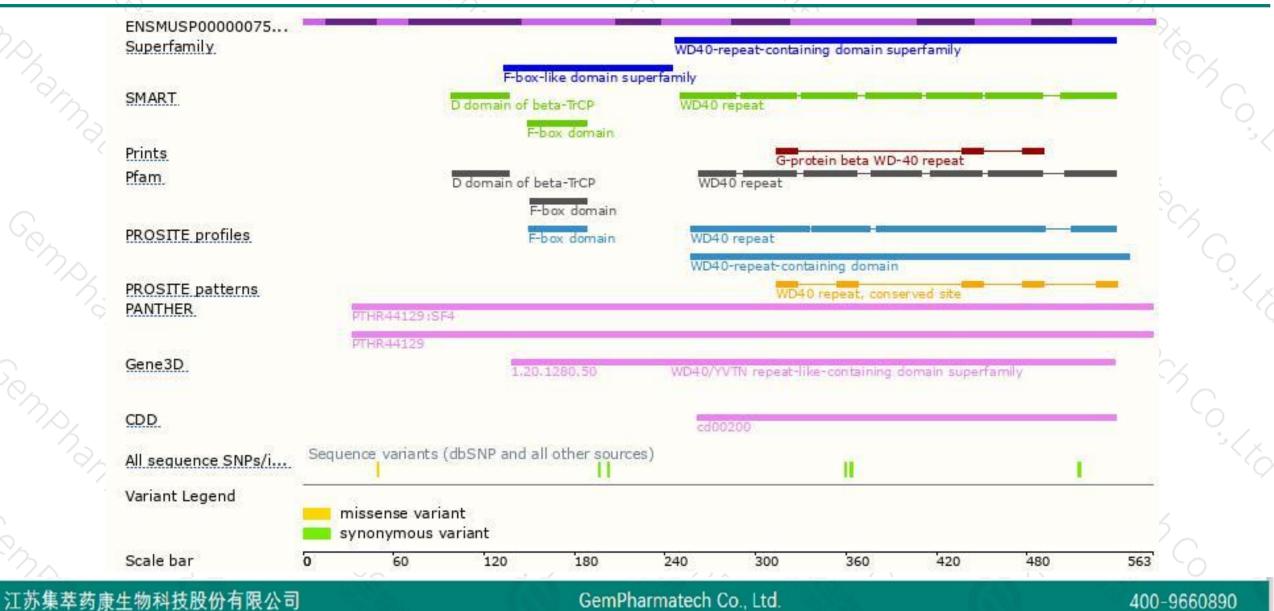


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### **Protein domain**







If you have any questions, you are welcome to inquire. Tel: 400-9660890



