

## ***klf2b*<sup>ihb249/+</sup> (CZRC catalog ID: CZ456)**

### **Nature of the mutation**

Between 655 bp to 661 bp of the wild-type *klf2b* coding sequence, GCGTATC is mutated to TGA in exon 2.

### **Sense Strand Sequence**

GGCAGAGCTCCTGCGATCTGATGTAGACTCCACGTATGACACCACAGTGCAGGGGAGATTCCTGCTCAACTCCAGCGGCTTCCCCAGACAGGAGTTTCCTGAAATCAAAGTGGAGCCGCCGATGGATGGCTACGGTCCGGTGATAGGCATGGTGCCACAAACCTGCCAAAAAATCAAGCAGGAAGGCAACGTTTCATGCATGATGTCTTTCGAGCAGCCCAGACTAGCAGTTCTCCGCAGGCCACCGGGAATGACTCCTCCGCTGAGTCTTGACGACTCCCACCTCCGTCAAACGACATATACACAGAGCTACCATCACTCTCCTCCGGCGTATCCGCAGGTGCCAATGCAGTTCACCGCGCCACACCAGTTTGCCATGTATGAGGAGGCAATGGGGATGCAACCCAGCATGCAGCGGGCTTTCTCACTCCTCCGTCTCTCCGTTAGAGCTAATGGAGTCTAAACCCAAGCGAGGGCGACGCACCTGGCCGAGGAAGCGCATGGCGACACATACATGCACATACGCCGGCTGCGGGAAGACGTACACAAAAGCTCC

Uppercase: Exon/coding sequence

Lowercase: intron/noncoding sequence

atcg : Forward/Reverse primer

atcg : Crispr/cas9 target site

### **Genotyping assay**

#### **Primers:**

**klf2b\_forward:** 5' GGCAGAGCTCCTGCGATCTG 3'

**klf2b\_reverse:** 5' GGAGCTTTTGGTGTACGTCT 3'

#### **PCR program:**

95°C 5min

95°C 30 sec

60°C 30 sec

72°C 30 sec

72°C 8min

4°C hold

} 30 Cycles

**Product size: 562 bp**

### The sequencing results of the CZ456:

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WT          GGCCACCGGGAATATGACTCCTCCGCTGAGTCTGACGACTCCCACCTCCGTCAAACGAC
CZ456       GGCCACCGGGAATATGACTCCTCCGCTGAGTCTGACGACTCCCACCTCCGTCAAACGAC
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WT          ATATACACAGAGCTACCATCACTCTCCTCCGGCGTATCCGCAGGTGCCAATGCAGTTCAC
CZ456       ATATACACAGAGCTACCATCACTCTCCTCCGTGA----CGCAGGTGCCAATGCAGTTCAC
*****

WT          CGCGCCACACCAGTTTGCCATGTATGAGGAGGCAATGGGGATGCAACCCAGCATGCAGCG
CZ456       CGCGCCACACCAGTTTGCCATGTATGAGGAGGCAATGGGGATGCAACCCAGCATGCAGCG
*****
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### Reference:

None