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***Lca5<sup>hzu7/+</sup> (AB) (CZRC catalog ID: CZ1277)***

**Nature of mutation**

Hzu7 allele was a mutant line, induced by the CRISPR-cas9 in the AB genetic background. 18-21 bp of the wild-type lca5 coding sequence, TAAT, is deleted from exon 2.

**Genotyping assay**

**Primers:**

Hzu7\_forward: 5' AGACAGCTTGAGTGTTGAGTGT 3'

Hzu7\_reverse: 5' TCTGCTCTGTGCATGATCCC 3'

PCR program:

94°C 4min  
94°C 30 sec  
56°C 30 sec      }  
72°C 30 sec      } 30 Cycles  
72°C 5min  
4°C hold

**Product size: 337 bp**

**Reference**

Qu Z, Yimer T A, Xie S, et al. Knocking out Lca5 in zebrafish causes cone-rod dystrophy due to impaired outer segment protein trafficking[J]. Biochimica et Biophysica Acta (BBA)-Molecular Basis of Disease, 2019, 1865(10): 2694-2705.