



Erratum

Rapid and efficient identification of the mouse leptin receptor mutation (C57BLKS/J-*Lepr^{db}*) by tetra-primer amplification refractory mutation system-polymerase chain reaction (ARMS-PCR) analysis

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We designed 4 primers for amplification of allele-specific PCR products from leptin receptor mutant and normal alleles using primer design web service for tetra-primer ARMS-PCR (Figure 1) i.e., *Lepr*-forward outer primer (*Lepr*-FO); 5'-AGGATACAATACAAGAACA AAAGCCTG-3', *Lepr*-forward inner primer (*Lepr*-FI);

5'-TATTAGAAGATGTTTACATTTTGATGGAGGG-3', *Lepr*-reverse inner primer (*Lepr*-RI); 5'-GTCATTCAAA CCAATAGITTAGGTTTIGTTTA-3', *Lepr*-reverse outer primer (*Lepr*-RO); 5'-ATGCAGAGTCCATGAATATCAACTTT AA-3'.

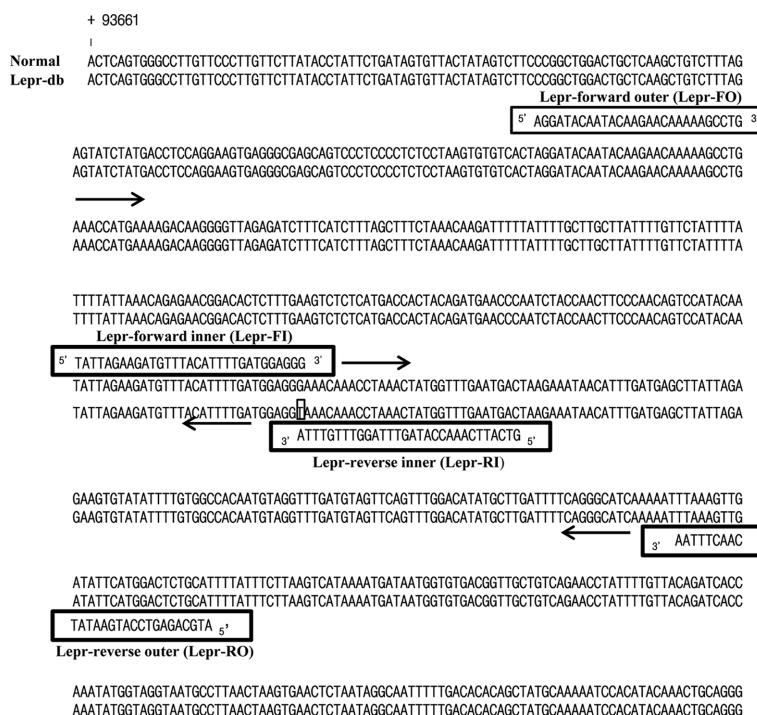


Figure 1. Sequences of the leptin receptor mutation and normal allele (NCBI, Gene ID: 16847). The primers for ARMS-PCR were designed using the primer design web service for tetra-primer ARMS-PCR. Underline indicates transverse point mutation (G→T) in intron 18 of the leptin receptor.

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