

Genotyping protocol

General information:

Strain name	C57BL/6-Fxr1<em2Geno>/Cnbc
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Primers:

Name	Sequence	Primer type
Fxr1-Fw	5'-TCCAGAGGGCAGTGCTTTAT-3'	Gene-specific
Fxr1-Rv	5'-TGCATTTTCTTGAGCAGCAT-3'	Gene-specific
Fxr1-seq Rv	5'-TGCAATGCAAATTCTGAATGT-3'	Gene-specific

In case more than two primers are introduced, please indicate how they should be combined:

	Forward primer	Reverse primer
PCR amplification	Fxr1-Fw	Fxr1-Rv
Sequencing		Fxr1-seq Rv

Reaction mix:

ddH ₂ O	18,5	μl
PCR Buffer 15mM MgCl ₂	2,5	μl
dNTPs 2,5mM	2,5	μl
Primer 100μM	0,125	μl
Primer 100μM	0,125	μl
Taq Polymerase 5U/μl	0,250	μl
DNA	1	μl
Final volume	25	μl

PCR program:

94.0 °C	5	min	
94.0 °C	30	sec	X10
62.5 °C	30	sec	
72.0 °C	45	sec	
94.0 °C	30	sec	X25
61.5 °C	30	sec	
72.0 °C	45	sec	
72.0 °C	5	min	

Expected fragment size:

PCR product length: 420 bp for wt and mutant.

Comments/Additonal information:

It's necessary to sequence with Fxr1-seq Rv primer to differentiate the mutation

Sequencing and PCR amplification primers

Part of *Mus musculus* wild type Fxr1 sequence (exon 15 and delACAG mutation)

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GCGGTCTCGTAGACGGAGGACTGATGAAGATGCTGTTCTGATGGATGGACTGACTGAATCTGATACAGCC
TCAGTTAATGAGAATGGGCTAGGTATGTAAGCACTTAGGGAAGAGTTGTGAGTGTGTGGTTGCATTGT
ATTGATAATGGAGGAAGCCTTTCTTTTTTTGTATAGAATTCACAAATGGTTTTTTTTTTCTTTCTTT
TCTTTTTAGTGTTTACTTTAAATGAGGGATATGGCTGCTATCTTGGGCACTCATTTTCATATCTGTCAGA
CTATCTGAACTGGGAGGATTCCTTAAGGTGCCCTTTCATACCATTAAATCAACCTAAATACTCAGTTGAT
CTGTCTACAAAACAAAACCTTAAGTGTTTTAGTCCCTCCCCCTTTAAAAATATAGTACATATTTATGGGGG
CTGCAAAAACCTAGCATTATTAATAAGGGGCTTTGAGATCCAGGGTAATGTTGGAGAGATATCTTATAT
TTCTGAATGATGGTGGTTGTTTCAATTACATGGCAGTTTGCTTTATATGAGTTATTAAGCCCTCTTAA
TTTCCAGAGGGCAGTGCCTTATGCTGCAAAAATGTGTATAATCTTTGTCAGTATTGTATATGCTTTAT
CTTTGATGTTTAATTAAGGTTTCTCTTTCCCTTAAATAAGGAAAGAAATTTTTGTTTGTTTTTTT
GTTTATTTTTTGTGAGAATAGTTGATGGTAAACTTTGCAATTACAGATGATAGTGAAAAAAAAACCCAGC
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AGAAGTGTCTTTTTCTTTAAAGAGGATGCATGTTAATGTTTTTGTGTTAATGTAGTGTGGTGCTTTTTC
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CAGCTTTGAAAGTAGCACAGCAGTGAAGATTTTTGACCCTACTTTTTTGAAGATAGAAAATTCAGGCT
ATTAGTATGGGCTCACTAAGTGAATGACTTTCAATCTAAGCCAGTGCATGTTAAATGTAC
AGACACAAAATATCAAAACAGGATTTCTTCCAAAAGAGATGATTTTGAATGGAATTAACATAAAATGAAT
AATTTGGAGTTTTGGTAATTTTAAACAGTGTGTGCTGTTGCATACTATATCTGATTTGTTTTACTGTTC
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