EMMA ID: *02345*
Gene: *P2rx5*
Common name: *LEXKO-1615, CHA391N1*
Allele: *P2rx5* ^{tm1Lex}

Genotyping Information

Genotyping by end-point PCR based on gel is composed of a genespecific short range PCR using primers on wild type allele and a mutant allele-specific short range PCR. The combined results show the genotype of the mice. For example: mutant positive, wild type positive = Heterozygous.

PCR primer pairs and expected size bands

Assay	Forward Primer	Reverse Primer	Expected Size Band (bp)
Wildtype	LEXKO-1615-1	LEXKO-1615-2	638
Mutant	Neo3a	LEXKO-1615-4	288

Primer sequences

Primer Name	Sequence 5' --> 3'
Neo3a	GCAGCGCATCGCCTTCTATC
LEXKO-1615-4	GCCCACGTTTAACAAGTCAG
LEXKO-1615-2	CCTATACATCTGCCTGACAG
LEXKO-1615-1	TGCCCAAGAACTCACATGAG

PCR setup (Qiagen, Hot Start Plus)

Component	Volume (µl) 1x	Final conc.
DNA (~ 50-100 ng)	2	
Q-Solution (5x)	2,5	0,5
PCR-Buffer (10x)	2,5	1
DNTP mix (10 mM)	0,5	0,2
$MgCl_2$ (25 mM)	1,5	1,5
Primer 1 (10 pmol/µl)	1	0,4
Primer 2 (10 pmol/µl)	1	0,4
Taq Polymerase (5 U/µl)	0,3	0,06
H_2O*	13,7	
Final volume	25	

*** The amount of H_2O is adjusted with the number of primer.**

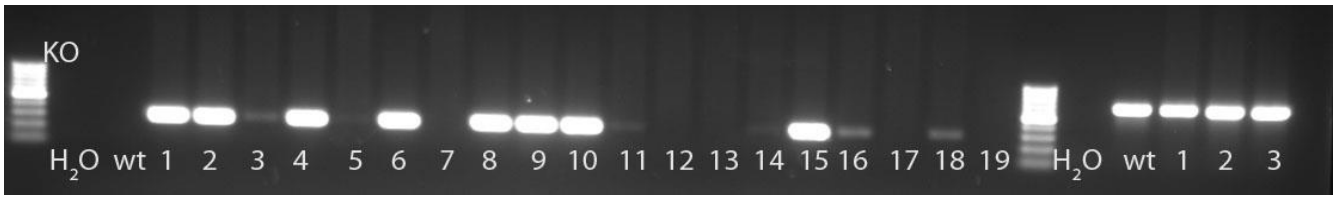
Amplification conditions

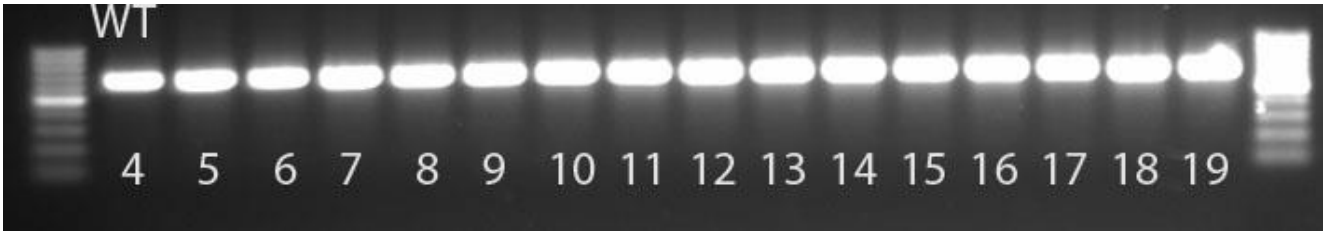
PCR Settings	Temperature (°C)	Time	# of cyles
1 Denaturation (Melting)	95°C	5 min	1
2 Amplification (Melting, Annealing, Polym.) touchdown, decreas annealing 1°C/cycle	94°C 65°C 72°C	30 sec 45 sec 45 sec	10
3 Amplification (Melting, Annealing, Polym.)	94°C 55°C 72°C	30 sec 45 sec 45 sec	30
3 Polymerisation	72°C	10 min	1
4 Cooling	12°C	hold	1

use Touch-Down cycling protocol: first 10 cycles anneal at 65°C, decreasing 1°C per cycle; next 30 cycles aneal at 55°C

These PCR conditions have been optimized for our methods and preparation kits. Adaptions may be required.

Gel Image





Separated by gel electrophoresis on a 2% agarose gel.