

EMMA ID: 04425

Gene: *Abca3*

Common name: *Abca3tm1Holz*

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 | Abca3_wtF | Abca3_R | 708 |
| Mutant | Neo_F | Abca3_R | 350 |

Primer sequences

| Primer Name | Sequence 5' --> 3' |
|-------------|------------------------------|
| Abca3_wtF | CACAGCCTACTACCAGCAAACAGGA |
| Abca3_R | GAAAGCCCATCCTAAAGTATCAGCC |
| Neo_F | CTGAAGAGCTTGGCGGCGAATGGGCTG/ |

PCR setup (LongAMP Taq)

| Component | Volume (µl) 1x |
|-------------------------|----------------|
| DNA (~ 50-100 ng) | 4 |
| 100% DMSO | 0,4 |
| PCR-Buffer (5x) | 4 |
| DNTP mix (10 mM) | 0,5 |
| Primer 1 (10 pmol/µl) | 1 |
| Primer 2 (10 pmol/µl) | 1 |
| Taq Polymerase (5 U/µl) | 0,3 |
| H ₂ O* | 13,7 |
| Final volume | 20 |

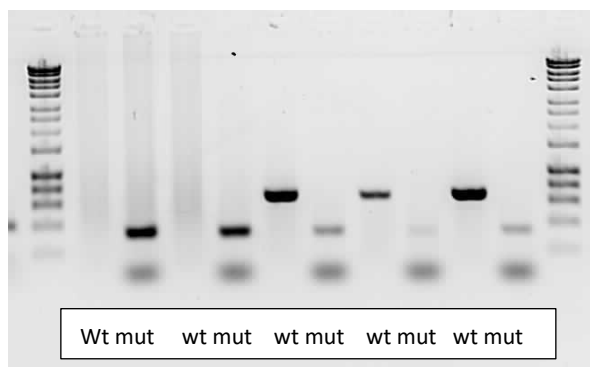
* The amount of H₂O is adjusted with the number of primer.

Amplification conditions

| PCR Settings | Temperature (°C) | Time | # of cycles |
|--|--------------------|--------|-------------|
| 1 Denaturation (Melting) | 94°C | 3 min | 1 |
| 2 Amplification (Melting, Annealing, Polym.) | 94°C | 30 sec | 39 |
| | 68-58 (↓1°C/Cycle) | 20 sec | |
| | 65°C | 60 sec | |
| 3 Polymerisation | 65°C | 10 min | 1 |
| 4 Cooling | 4°C | hold | 1 |

use Touch-Down cycling protocol: first 10 cycles anneal at 68°C, decreasing 1°C per cycle, next 30 cycles anneal at 58°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.