

Genotyping protocol

Synonymous S293S PM in Shh

Shh^{em1.1ics}

(PHENOMIN-ICS reference IR00007498 / Kos7498)

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Synonymous S293S PM in Shh - Shh^{em1.1Ics}

This protocol describes the condition used at the Mouse Clinical Institute (ICS) to genotype the Synonymous S293S point mutation in Shh, Shhem1.1Ics.

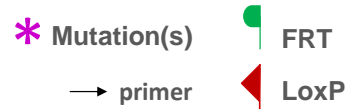
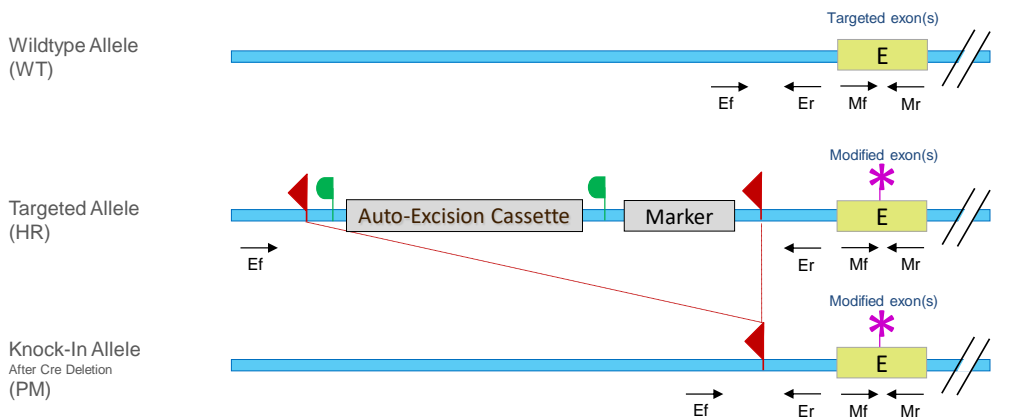
1. PCR Genotyping protocol

1.1. Genotyping strategy

The map below describes the position of the primers used for genotyping for each possible allele.



PM Genotyping strategy



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Sequence of primers used for genotyping:

| Position | Sequence |
|-----------------|------------------------|
| Ef | GGAGAAAAGCTGGGAGTGTCC |
| Ef ² | GCGGAGCAATCTTGGAGGG |
| Er | GGGATAAGGGTGGGAGTAGATC |
| Er ² | CTTTGGGGTGGAGAGACGGG |
| Mf | CGCCAAGAAGGTCTTCTACGTG |
| Mr | CGATGACAGCGTAGCACGAG |

²: for a selected position, a second primer was designed

PCR fragments expected size (bp):

| Region analyzed | Position on the primer (see the map above) | Targeted allele (HR) | PM allele | WildType allele |
|---|---|-------------------------|--------------|--------------------|
| WildType / Mutated alleles (with DMSO) | Mf / Mr | 317 | 317 | 317 |
| Excision selection Marker | Ef / Er | 4421* | 266 | 187 |
| Excision selection Marker (2) | Ef ² / Er ² | 4379* | 224 | 145 |

*: amplicon will not be observed using our genotyping conditions



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This section describes the composition of the mix and cycling conditions used for genotyping.

| Reagents: | Volume: |
|--------------------------------|-------------|
| - FastStart PCR Master (Roche) | 7.5µl |
| - DNA (50ng/µl) | 1.5µl |
| - 5' primer (100 µM) | 0.06µl |
| - 3' primer (100 µM) | 0.06µl |
| - Sterile H ₂ O | up to 15 µl |

If mentioned in table "PCR fragments expected size" add 5% of DMSO in the reaction mix

| Cycling conditions: | | |
|---------------------|------|---------|
| Temp | Time | #Cycles |
| 95°C | 4min | 1 |
| 94°C | 30s | 35 |
| 62°C | 30s | |
| 72°C | 1min | |
| 72°C | 7min | 1 |
| 14°C | --- | --- |

NB: These PCR conditions have been optimized for high-throughput genotyping. Adaptation to small-scale may be required.



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2. Recommended papers:

2.1. Cre and Flp genotyping method

[Highly-efficient, fluorescent, locus directed cre and FlpO deleter mice on a pure C57BL/6N genetic background.](#)

Birling MC, Dierich A, Jacquot S, Héroult Y, Pavlovic G.

Genesis. 2012 Jun;50(6):482-9. doi: 10.1002/dvg.20826. Epub 2012 Mar 20.

2.1. Tips and tricks for optimizing your PCR genotyping procedures

[Optimizing PCR for mouse genotyping: Recommendations for reliable, rapid, cost effective, robust and adaptable to high-throughput genotyping protocol for any type of mutation.](#)

Jacquot, S, Chartoire, N, Pigué, F, Héroult, Y, Pavlovic, G. (2019).

Current Protocols in Mouse Biology, 9, e65. doi: 10.1002/cpmo.65

Free copy of this paper can be accessed online through this link <http://bit.ly/2sxxWvO>

