



Genotyping protocol

Usp25

IR00004144 / P4144

(ICS internal reference)

This report has been prepared by: **David Moolaert**
genotyping@igbmc.fr

This report has been validated by: **Sylvie Jacquot, PhD, Head of Genotyping Service**
33 (0)3 88 65 57 44
genotyping@igbmc.fr

The first version of this report was finalized the: 31 Jul 2015

For any question, please contact:

Institut Clinique de la Souris - ICS - Mouse Clinical Institute
1 rue Laurent Fries, BP 10142
67404 Illkirch Cedex, France
Email: genotyping@igbmc.fr
Web site: <http://www-mci.u-strasbg.fr/>

TABLE OF CONTENTS

Table of contents2

1. Genotyping protocol and data2

 1.1. Genotyping strategy2

 1.2. PCR protocol4

 1.3. Picture of genotyping with various alleles5

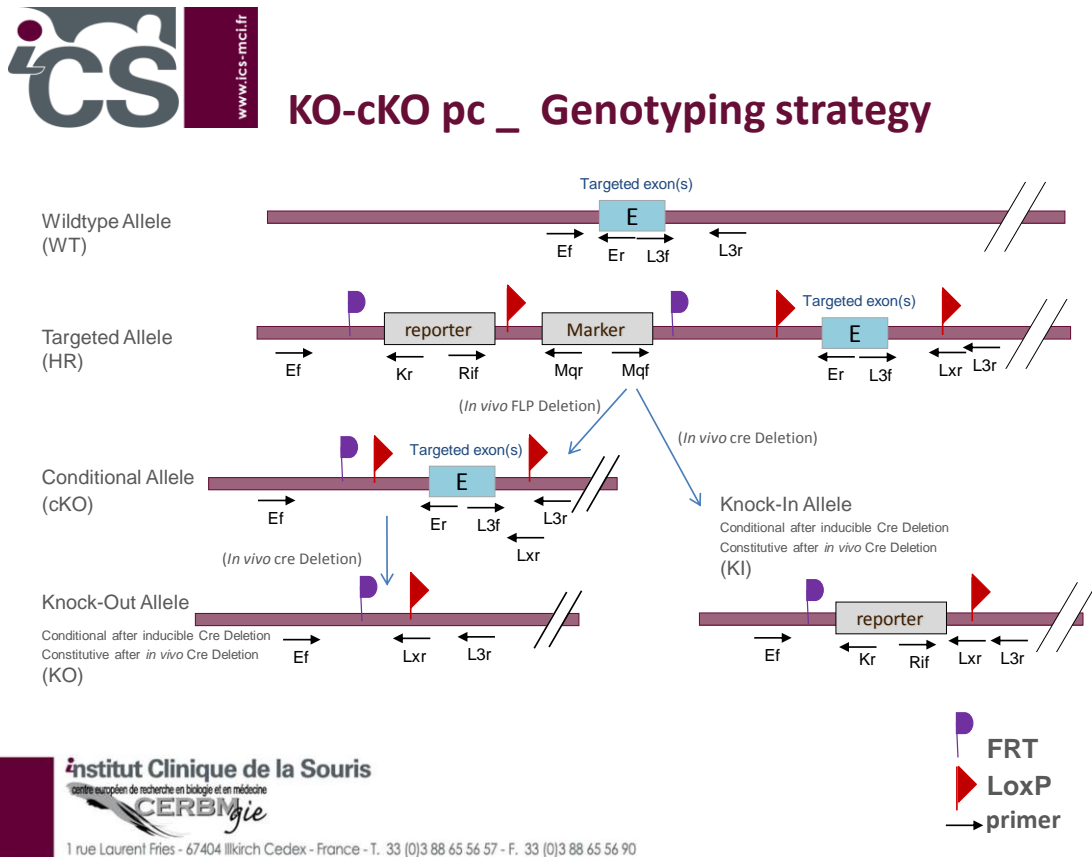
2. Cre and Flp genotyping method6

1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Usp25** Constitutive Knockout / Conditional Knockout (KO-cKO x Cre) project.

1.1. Genotyping strategy

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



Sequence of primers used for genotyping:

| Position | Primers | Sequence |
|----------|---------|---------------------------|
| Ef | 7153 | CGCGCACACACACTCCTATGCT |
| Er | 7152 | AATGCCCTGTTTGACTCCGCCA |
| Kr | 3277 | CTCCTACATAGTTGGCAGTGTGGG |
| L3f | 7150 | TCTGAAGAATGTGGCAACACCTGC |
| L3r | 7151 | CGGCACTCCAAAGCTGAAGTAGCAG |
| Lxr | 3255 | ACTGATGGCGAGCTCAGACCATAAC |
| Ri1f | 5966 | GCACATGGCTGAATATCGACGGT |

PCR fragments expected size (bp):

| Region analyzed | Primers used | Position on the primer (see the map above) | Targeted allele (HR) | cKO allele (cKO) | KI allele | WildType allele (WT) |
|----------------------------------|--------------|---|----------------------|------------------|-----------|----------------------|
| 5' part of the selection marker | 7153-3277 | Ef / Kr | 231 | --- | 231 | --- |
| Presence of the distal loxP | 7150-7151 | L3f / L3r | 369 | 369 | --- | 352 |
| Distal loxP specific PCR | 7150-3255 | L3f / Lxr | 283 | 283 | --- | --- |
| Excision of the selection marker | 7153-7152 | Ef / Er | 7592* | 688 | --- | 478 |
| Cre total excision | 5966-3255 | Ri1f / Lxr | 5112* | --- | 471 | --- |

*: this PCR product will not be observed using our PCR genotyping conditions (see description below)

** : this PCR is only verified if mice are generated

---: no Amplicon should be obtained

1.2. PCR protocol

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 |

Cycling conditions:

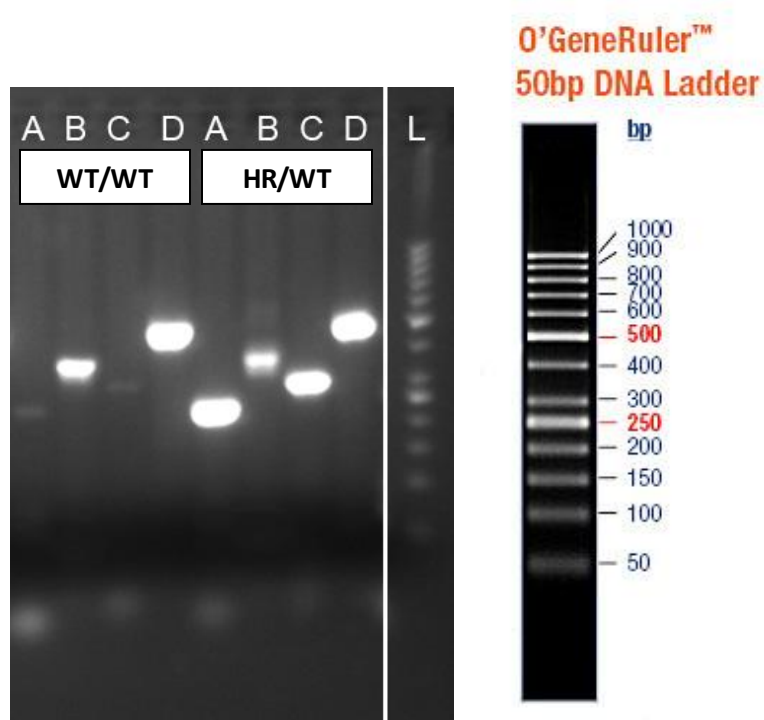
| Temp | Time | #Cycles |
|------|------|---------|
| 95°C | 4min | 1 |
| 94°C | 30s | 34 |
| 62°C | 30s | |
| 72°C | 1min | |
| 72°C | 7min | 1 |
| 20°C | 5min | 1 |

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

1.3. Picture of genotyping with various alleles

Analysis of PCR products pattern was done by gel electrophoresis 2% agarose (SB buffer).

Representative genotyping picture



- A: 5' part of the selection marker
- B: Presence of the distal loxP
- C: Distal loxP specific PCR
- D: Excision of the selection marker
- L: O'GeneRuler 50bp DNA Ladder

2. Cre and Flp genotyping method

You will find the genotyping protocol in the publication:

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

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

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