



## Genotyping protocol

D16Ertd472e

/ P5374

(ICS internal reference)

This report has been prepared by:

**David Moulaert**  
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

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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](mailto:genotyping@igbmc.fr)  
Web site: <http://www-mci.u-strasbg.fr/>

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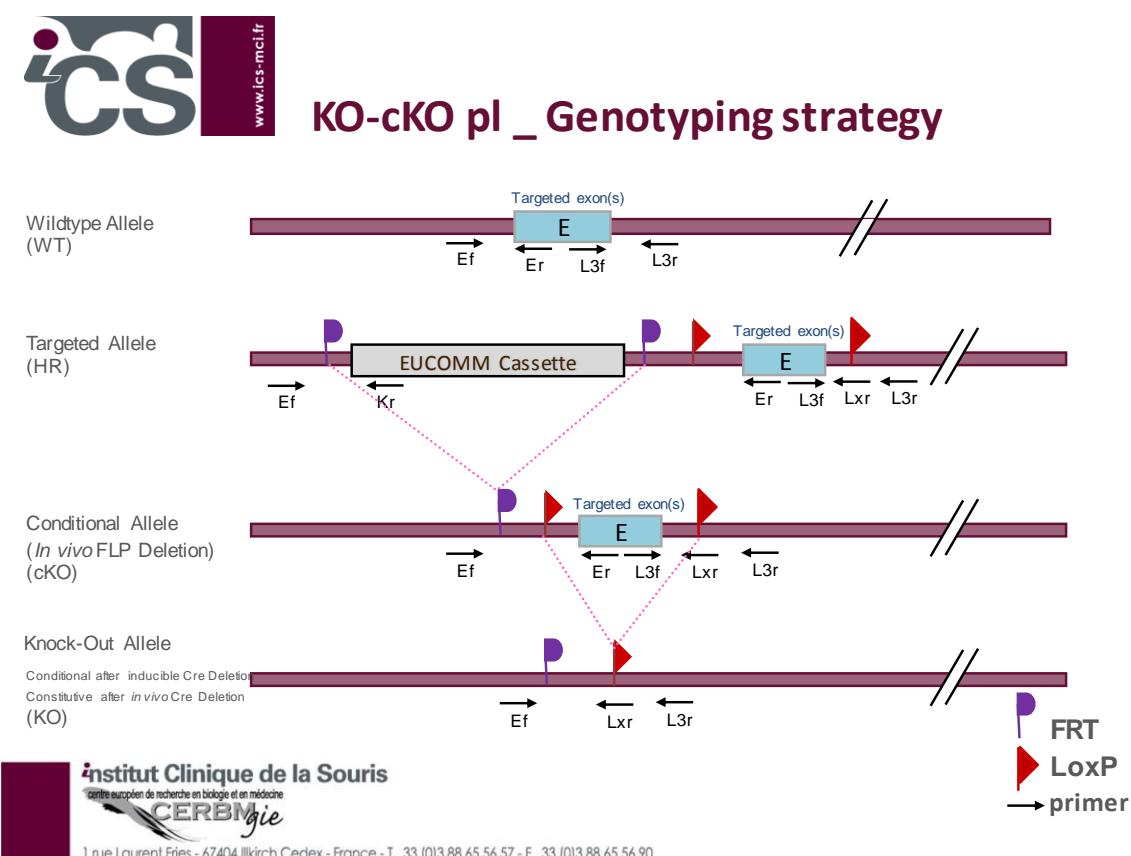
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## 1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **D16Ertd472e** Constitutive Knockout / Conditional Knockout (KO-cKO) project.

### 1.1. Genotyping strategy

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



**Institut Clinique de la Souris**  
 centre européen de recherche en biologie et en médecine  
**CERBM**  
 gie  
 1 rue Laurent Fries - 67404 Illkirch Cedex - France - T. 33 (0)3 88 65 56 57 - F. 33 (0)3 88 65 56 90

## Sequence of primers used for genotyping:

| Position         | Primers | Sequence                     |
|------------------|---------|------------------------------|
| Ef               | 8361    | CAAAGGTTAACATGATCGCTGGGAACC  |
| Ef <sup>2</sup>  | 8362    | CCGGGCTAACAGACTCCAATTGAATC   |
| Er               | 8366    | CACTAGTATGTTGTATGTGTGGAGGCC  |
| Kr               | 3277    | CTCCTACATAGTGGCAGTGTTGG      |
| L3f              | 8364    | CCGTGAAGAATGTGAGTAGTGTGTGACG |
| L3f <sup>2</sup> | 8363    | CTTTTGGGGACGGTTGGTG          |
| L3r              | 8365    | CCAGTGTATAACGGCCAATAGTTCCAG  |
| Lxr              | 3255    | ACTGATGGCGAGCTCAGACCATAAC    |
| Mqf              | 4981    | GGGATCTCATGCTGGAGTTCTCG      |

<sup>2</sup>: for a selected position, a second primer was designed

## PCR fragments expected size (bp):

| Region analyzed                                  | Primers used | Position on the primer (see the map above) | Targeted allele (HR) | conditional allele (KO-cKO) | KO allele | WildType allele |
|--|--------------|--|----------------------|-----------------------------|-----------|-----------------|
| 5' part of the selection marker                  | 8361-3277    | Ef / Kr                                    | 297                  | ---                         | ---       | ---             |
| Presence of the distal loxP                      | 8364-8365    | L3f / L3r                                  | 453                  | 453                         | ---       | 407             |
| Distal loxP specific PCR                         | 8363-3255    | L3f <sup>2</sup> / Lxr                     | 221                  | 221                         | ---       | ---             |
| Excision of the selection marker                 | 8362-8366    | Ef <sup>2</sup> / Er                       | 5834*                | 420                         | ---       | 284             |
| Excision of the floxed exon(s), i.e. knock out 1 | 8361-8365    | Ef / L3r                                   | 7081*                | ---*                        | 475**     | 1485**          |

\*: 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:

- FastStart PCR Master (Roche)
- DNA (50ng/ $\mu$ l)
- 5' primer (100  $\mu$ M)
- 3' primer (100  $\mu$ M)
- Sterile H<sub>2</sub>O

Volume:

- 7.5 $\mu$ l
- 1.5 $\mu$ l
- 0.06 $\mu$ l
- 0.06 $\mu$ l
- up to 15  $\mu$ l

Cycling conditions:

| Temp | Time | #Cycles |
|------|------|---------|
| 95°C | 4min | 1       |
| 94°C | 30s  |         |
| 62°C | 30s  | 34      |
| 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.**

## 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, Héroult Y, Pavlovic G.  
Genesis. 2012 Jun;50(6):482-9. doi:10.1002/dvg.20826. Epub 2012 Mar 20.