



## Genotyping protocol

Cd276

IR00005544 / P5544

(ICS internal reference)

This report has been prepared by: **Christelle Roth**  
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 generated the: 14 Mar 2016

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/>

## TABLE OF CONTENTS

**Table of contents** ..... 2

**1. Genotyping protocol and data**..... 2

    1.1. Genotyping strategy ..... 2

    1.2. PCR protocol ..... 4

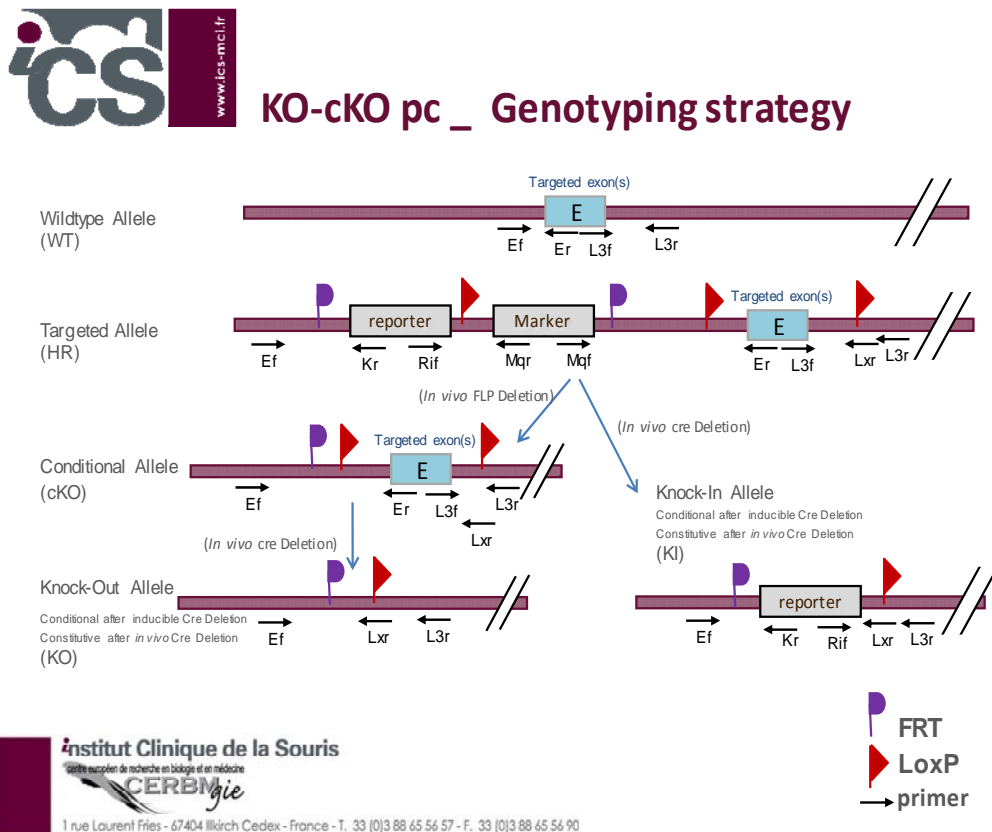
**2. Cre and Flp genotyping method** ..... 5

### 1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Cd276** 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	8962	GCTGCAGAGGTCCTATTCCTCCTTAATG
Ef <sup>2</sup>	8963	GGTCTCTGTGATTCTGACTCAGGAGGC
Er	8966	CTGGGATAGGCCTGCCCTGGAC
Kr	3209	CCAACAGCTTCCCCACAACGG
L3f	8964	CCTCTCCTCTAGGCCCTAACTGGAATATC
L3r	8965	GAGATGCACCTTCAAGGCTCCTGG
Lxr	3255	ACTGATGGCGAGCTCAGACCATAAC
Ri1f	5966	GCACATGGCTGAATATCGACGGT

<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 (cKO)	KI allele	WildType allele
5' part of the selection marker (with DMSO)	8962-3209 (with 5% DMSO)	Ef / Kr	325	---	---	---
Presence of the distal loxP (with Betaine)	8964-8965 (with 0.5% Betaine)	L3f / L3r	321	321	---	342
Distal loxP specific PCR	8964-3255	L3f / Lxr	214	214	---	---
Excision of the selection marker (with DMSO)	8963-8966 (with 5% DMSO)	Ef <sup>2</sup> / Er	7288*	384	---	232
Cre total excision	5966-3255	Ri1f / Lxr	3401*	---	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 <sub>2</sub> 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.**

## 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.