



## Genotyping protocol

Fkbp9

IR00003188 / E217

(ICS internal reference)

This report has been prepared by: **Nathalie Chartoire**  
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: 06 Mar 2014

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

    1.3. Picture of genotyping with various alleles .....5

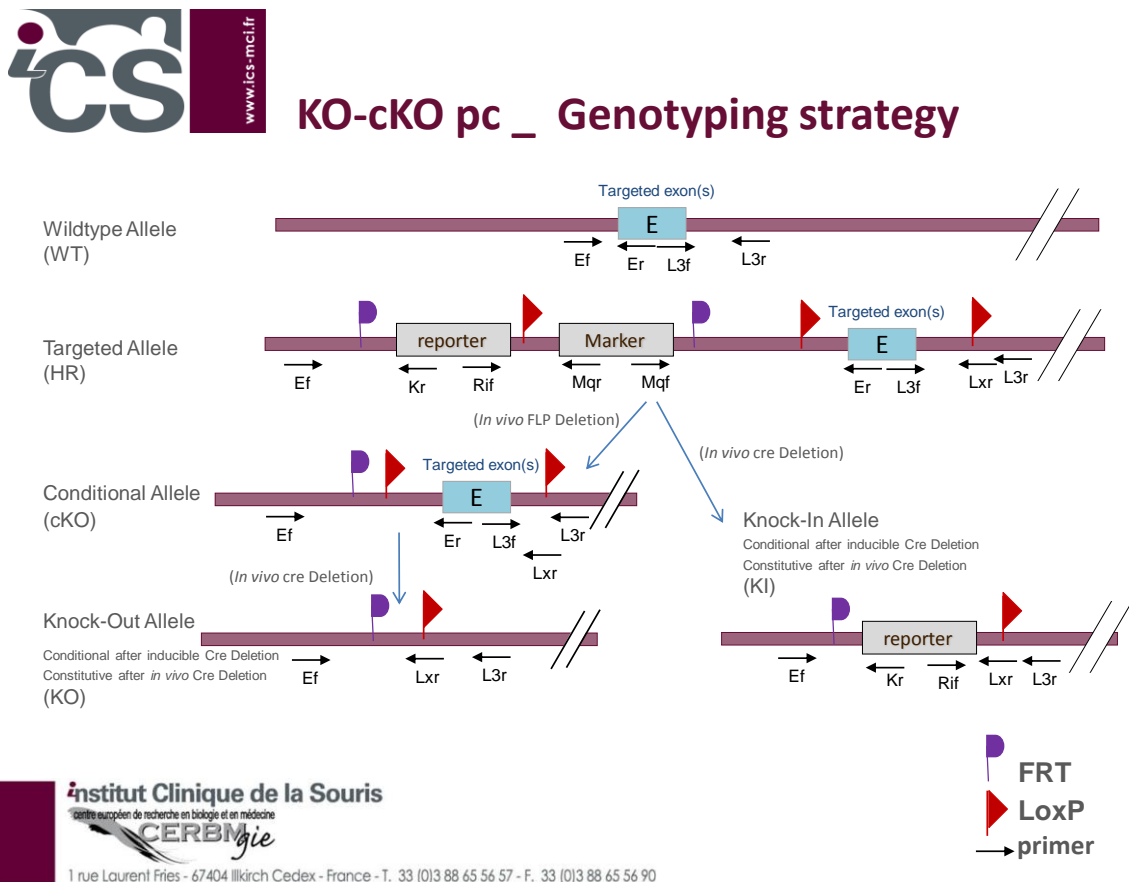
**2. Cre and Flp genotyping method** .....6

### 1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Fkbp9** Constitutive Knockout / Conditional Knockout (KO-cKOxcre) 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	5939	GTGGAAGCTGTAGTGTGTCACAATGTG
Ef <sup>2</sup>	5938	CACCCATATACTTCTCCATCCTGG
Er	5942	CTATAGGATCTCCGTGACACAGGG
Kr	3210	CCTGTCCCTCTCACCTTCTACC
L3f	5940	GCCCAACACCCATAGCTAGCATTTTC
L3r	5941	CTATCAATGACCTACGTGCCTAAGTGG
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 (KO-cKO)	KO allele	WildType allele
5' part of the selection marker	5939-3210	Ef / Kr	351	---	351	---
Presence of the distal loxP	5940-5941	L3f / L3r	321	321	---	279
Distal loxP specific PCR	5940-3255	L3f / Lxr	188	188	---	---
Excision of the selection marker	5938-5942	Ef2 / Er3	7297*	393	---	262
Cre total excision	5966-3255	Ri1f / Lxr	---	---	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:

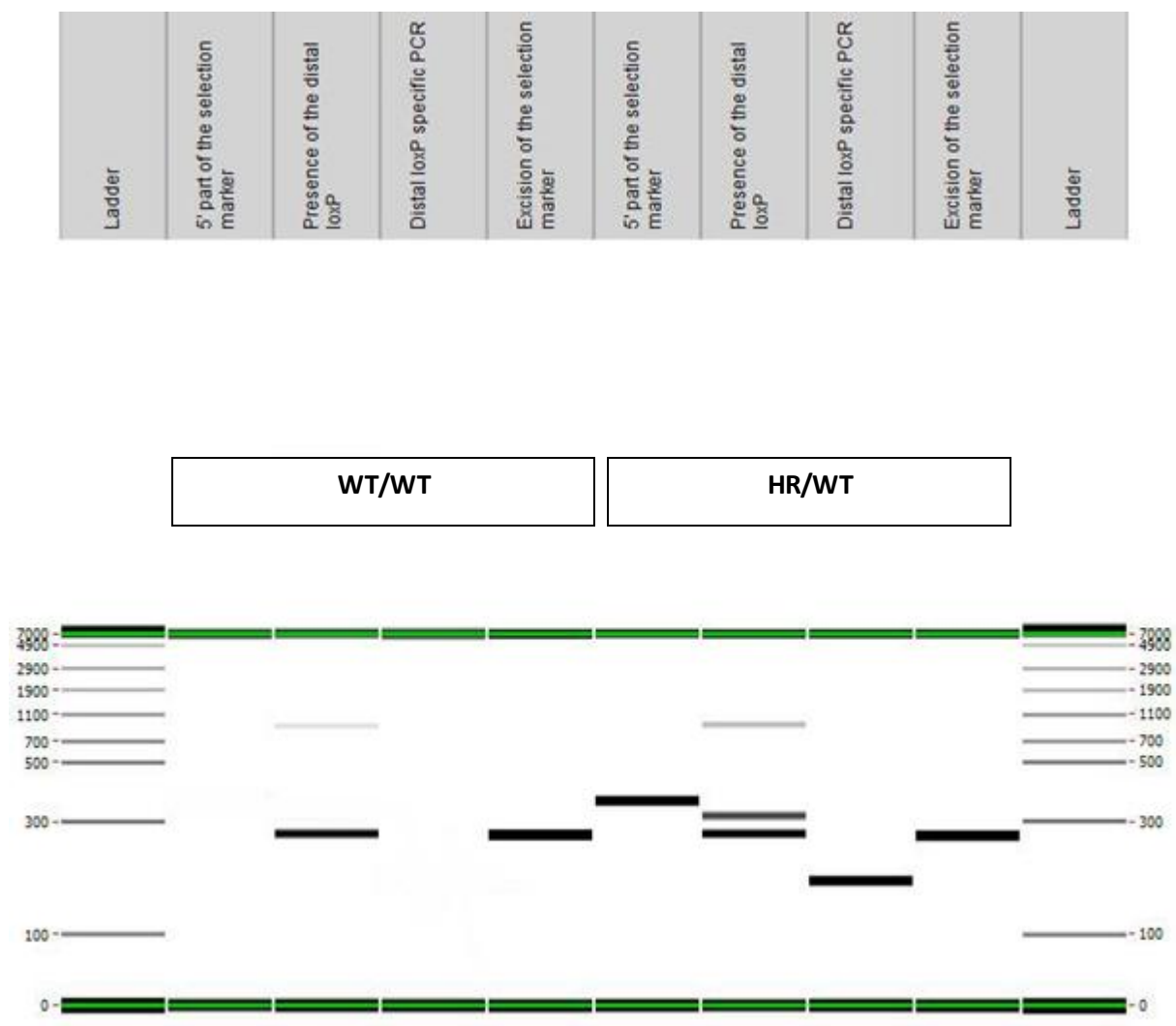
Temp	Time	#Cycles
95°C	4min	1
94°C	30s	
62°C	30s	34
72°C	1min	
72°C	7min	1
20°C	5min	1

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

### 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



## 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.