



Genotyping protocol

Pgam5

IR00003635 / E264

(ICS internal reference)

This report has been prepared by:

Pauline Cayrou
33 (0)3 88 65 56 55
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](mailto:genotyping@igbmc.fr)

The first version of this report was finalized the: 31 Oct 2013

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

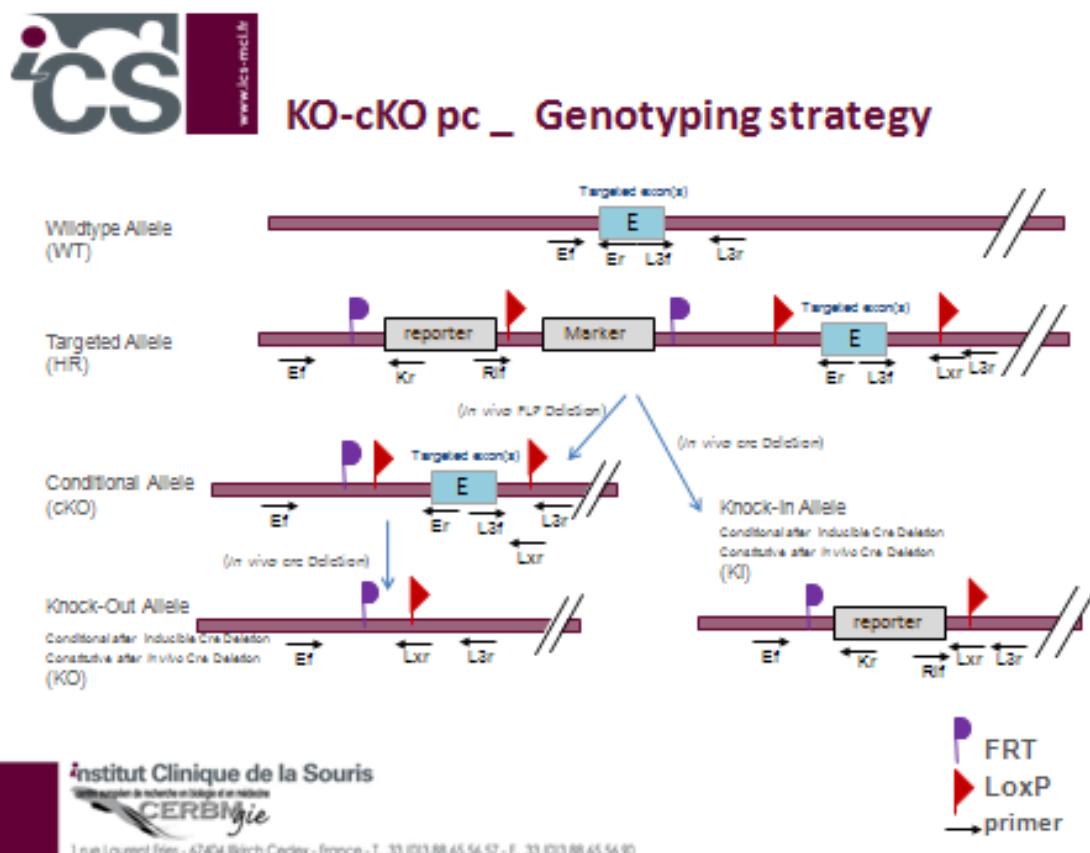
Table of contents	2
1. Genotyping protocol and data.....	2
1.1. Genotyping strategy	2
1.2. PCR protocol	4
1.3. Picture of genotyping with various alleles	5
2. Cre and Flp genotyping method	6

1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Pgam5** 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.



Institut Clinique de la Souris
Réseau national de recherche en biologie et en médecine

CERBInGie

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	6762	CAGTACAGTTCTAGGCTCCG
Er	6763	GTTCTCGGTTGACTGAGAG
Kr	3277	CTCCTACATAGTTGGCAGTGTTGGG
L3f	6764	AGGCTGGATCACTATAAGGC
L3r	6765	CTGGAGACATTGTGACCATC
Lxr	3255	ACTGATGGCGAGCTCAGACCATAAC
Rif	5966	GCACATGGCTGAATATCGACGGT

PCR fragments expected size (bp):

PCR N°	Region analyzed	Primers used	Position on the primer <i>(see the map above)</i>	Targeted allele (HR)	cKO allele	KI allele	WildType allele (WT)
A	5' part of the selection marker	6762-3277	Ef / Kr	346	---	346	---
	Distal loxP specific PCR	6764-3255	L3f / Lxr	396	396	---	---
B	Presence of the distal loxP	6764-6765	L3f / L3r	444	444	---	388
C	Excision of the selection marker	6762-6763	Ef / Er	7338*	434	---	238
D	Cre total excision	5966-3255	Rif / Lxr	3311*	---	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:

- FastStart PCR Master (Roche)
- DNA (50ng/ μ l)
- 5' primer (100 μ M)
- 3' primer (100 μ M)
- Sterile H₂O

Volume:
7.5 μ l
1.5 μ l
0.06 μ l
0.06 μ l
up to 15 μ l

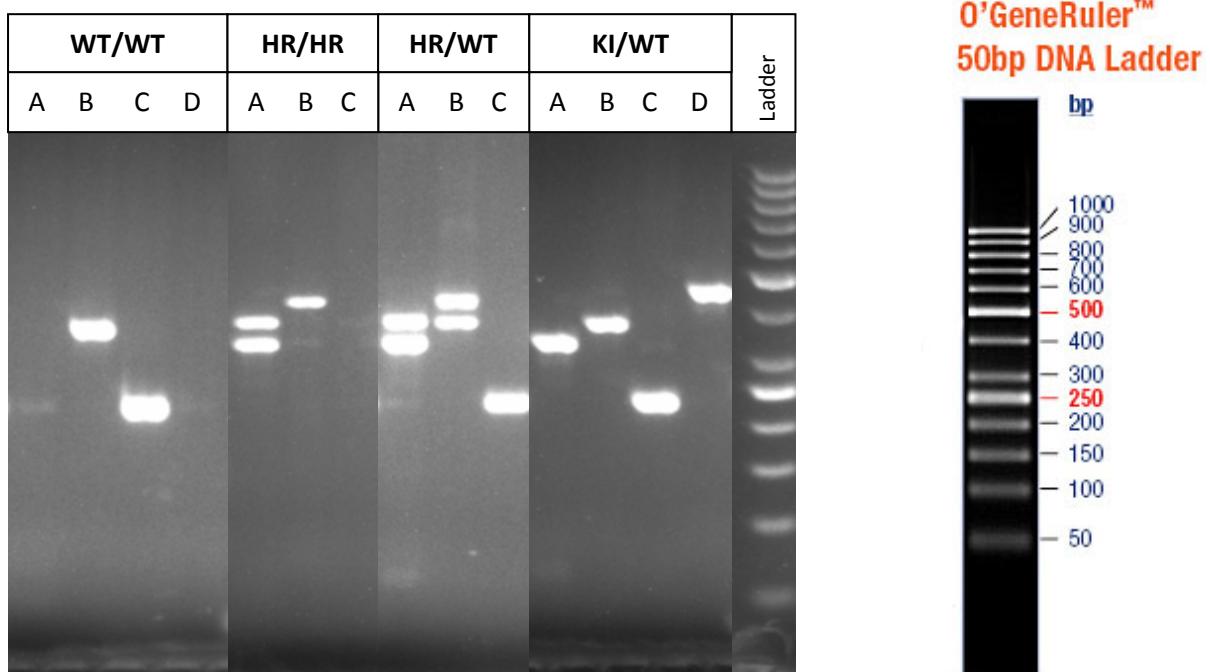
Cycling conditions:		
Temp	Time	#Cycles
95°C	4min	1
94°C	30s	
62°C	30s	34
72°C	1min	
72°C	7min	1
20°C	5min	1

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

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