

## Genotyping protocol

Project pCAG-Lox-STOP-Lox-Nr3c2 in Rosa

Gt(ROSA)26Sor<sup>tm25(CAG-Nr3c2)Ics</sup>/Ics

(PHENOMIN-ICS reference IR00007177 / Kos7177)

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pCAG-Lox-STOP-lox-Nr3c2 in Rosa - Gt(ROSA)26Sor<sup>tm25(CAG-Nr3c2)Ics</sup>/Ics

This protocol describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **pCAG-Lox-STOP-Lox-Nr3c2 in Rosa** Knockin (KI) project.

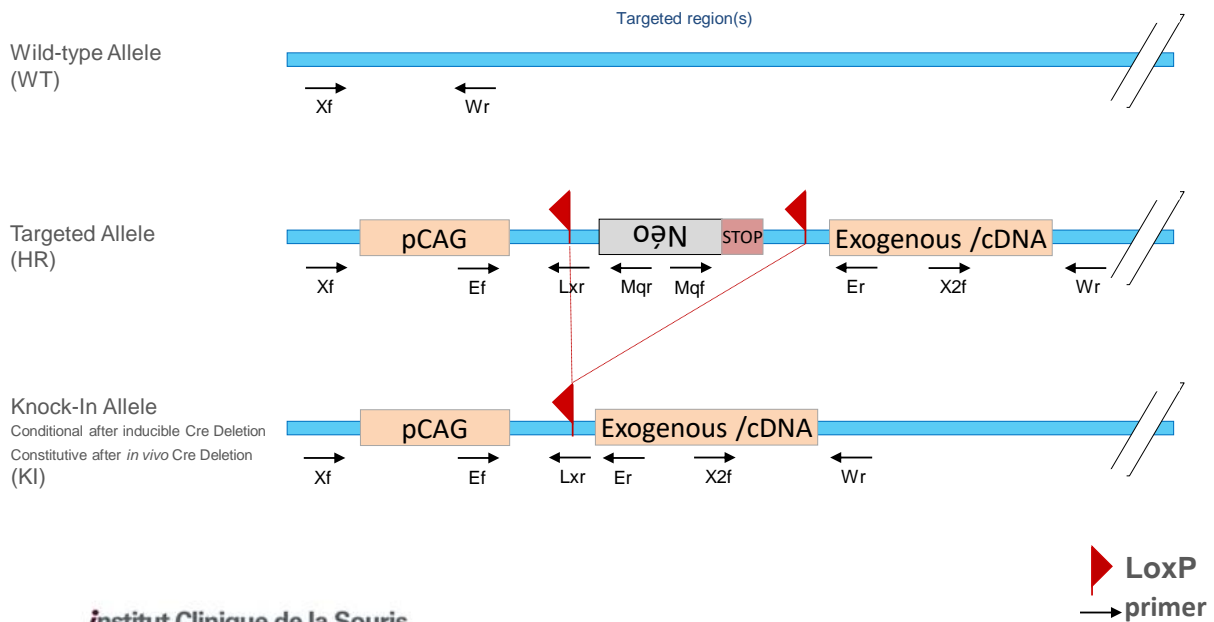
## 1. PCR Genotyping protocol

### 1.1. Genotyping strategy

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



## KI Genotyping strategy



## Sequence of primers used for genotyping:

Position	Sequence
Ef	GGCTCTAGAGCCTCTGCTAACC
Ef <sup>2</sup>	CTAGAGCCTCTGCTAACCATGTTC
Er	GCTCCAAGGTCTGAGACACTTGAC
Er <sup>2</sup>	CAAGAGAAGAACGCTCCAAGGTC
Lxr	CGAGGAATTCTTTGCCAAAATGATGAG
Mq1f	CTGCATTCTAGTTGTGGTTTGTC
Mq1r	CCTTCTTGACGAGTTCTTCTGAGGG
Wr	GTTCAATCCCCTGCAGGACAAC
Xf	GGGGAGGGGAGTGTGCAATAC
X2f	CTTTCACAGGAAGTGAGGCGCG

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

## PCR fragments expected size (bp):

Region analyzed	Position on the primer (see the map above)	Targeted allele (HR)	KI allele	Wild-Type allele
Wild-Type allele specific PCR (5' part of the targeted locus)	Xf / Wr	7935*	5430*	203
Excision of the selection marker	Ef / Er	2755*	250**	---
Excision of the selection marker 2	Ef <sup>2</sup> / Er <sup>2</sup>	2763*	258**	---
5' part of the selection marker	Ef / Mq1r	236	---	---
3' part of the selection marker (with Betaine)	Mq1f / Er <sup>2</sup>	219	---	---
Exogenous/cDNA specific PCR	X2f / Wr	464	464	---
LoxP specific PCR	Ef / Lxr	109	109	---

\*: 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

If mentioned in table "PCR fragments expected size" add 0.5% of Betaine in the reaction mix

Cycling conditions:		
Temp	Time	#Cycles
95°C	4min	1
94°C	30s	
62°C	30s	35
72°C	1min	
72°C	7min	1
14°C	---	

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



## 2. Recommended papers:

### 2.1. Cre and Flp genotyping method

[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.

### 2.1. Tips and tricks for optimizing your PCR genotyping procedures

[Optimizing PCR for mouse genotyping: Recommendations for reliable, rapid, cost effective, robust and adaptable to high-throughput genotyping protocol for any type of mutation.](#)

Jacquot, S, Chartoire, N, Piguet, F, Héroult, Y, Pavlovic, G. (2019).

Current Protocols in Mouse Biology, 9, e65. doi: 10.1002/cpmo.65

Free copy of this paper can be accessed online through this link <http://bit.ly/2sxxWvO>

