



Genotyping protocol
conditional miR-142 in Rosa
Gt(ROSA)26Sor^{tm1(CAG-Mir142)Sbel}/Ics
IR00004618 / K4618
(ICS internal reference)

This report has been prepared by: **Valérie Rousseau**

This report has been validated by: **Sylvie Jacquot, PhD, Head of Genotyping Service**

The first version of this report was generated the: 25 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: mutagenesis@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

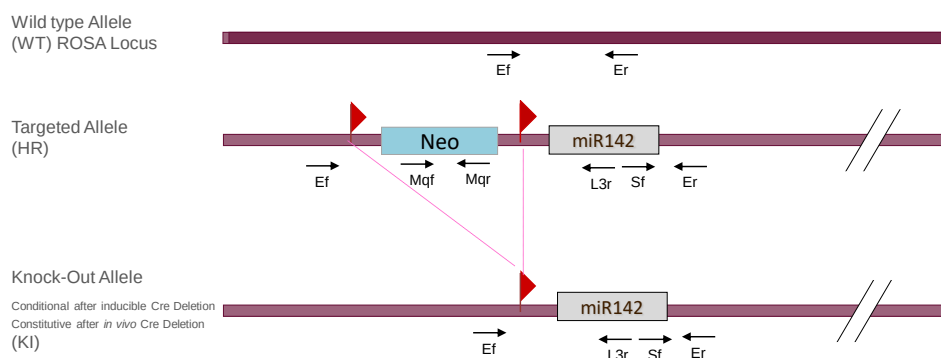
This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **conditional miR-142 in Rosa** Knockin (KI) project.

1.1. Genotyping strategy

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



KI Genotyping strategy



Genotyping protocol

conditional miR-142 in Rosa - Gt(ROSA)26Sor^{tm1(CAG-Mir142)Sbel}/Ics



Sequence of primers used for genotyping:

Position	Primers	Sequence
Ef	7401	AGCACGTTTCCGACTTGAGTTGCC
Er	7402	CACACACCAGGTTAGCCTTTAAGCCT
L3r	7404	CATAAAGTAGGAAACAC
Mqf	4122	GCTATGACTGGGCACAACAGACAATC
Mqr	548	CCAGACTGCCTTGGGAAAAG
Mqr ²	4123	CAAGGTGAGATGACAGGAGATCCTG
Sf	7403	AAGCACTACTAACAGCACTGGAGGGTG

²: for a selected position, a second primer was designed

PCR fragments expected size (bp):

PCR number	Region analyzed	Primers used	Position on the primer (see the map above)	Targeted allele (HR)	KI allele	WT allele
1	WildType allele specific PCR (5' part of the targeted locus)	7401-7402	Ef / Er	3695	1028	386
2	5' part of the selection marker	7401-548	Ef / Mqr	619	---	---
3	internal marker	4122-4123	Mqf / Mqr ²	261	---	---
4	cDNA 3'	7403-7402	Sf / Er	411	411	---
5	Excision of the floxed exon(s), i.e. knock out	7401-7404	Ef / L3r	3330	663**	---

*: 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 ₂ O	up to 15 µl

Cycling conditions:

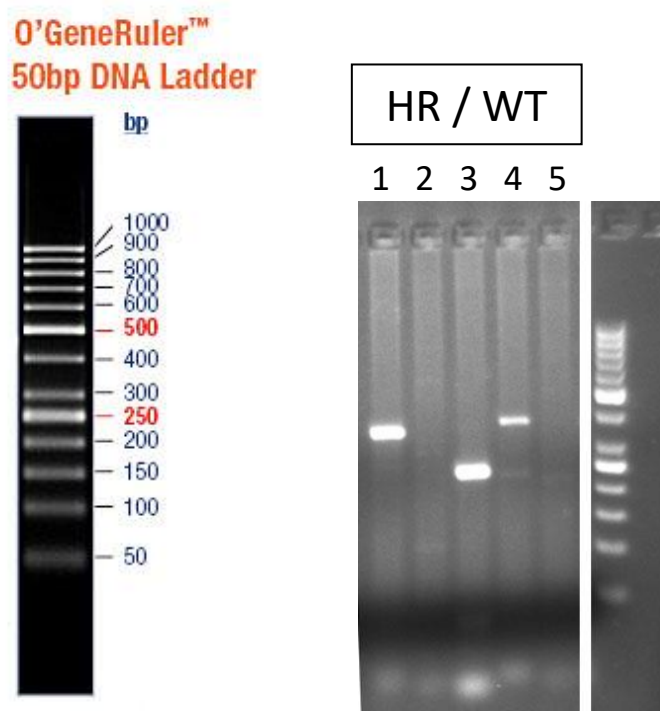
Temp	Time	#Cycles
95°C	4min	1
94°C	30s	34
62°C	30s	
72°C	1min	
72°C	7min	1
20°C	5min	1

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

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.