

Genotyping protocol

Project conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa

Gt(ROSA)26Sor^{tm30.1(CAG-C4b,-mScarlet)Ics}/Ics

(PHENOMIN-ICS reference IR00007413 / Kos7413)

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 finalized the: 14 Dec 2020

The last update of this report was done the: 1 July 2021

For any question, please contact:

PHENOMIN-ICS

Email: mutagenesis@igbmc.fr

Web site: <http://www.ics-mci.fr/>



conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa - Gt(ROSA)26Sor^{tm30.1(CAG-C4b,-mScarlett)lcs}/lcs

Table of contents

1. Genotyping protocol and data	3
1.1. Genotyping strategy	3
1.2. PCR protocol	5
2. Cre and Flp genotyping method	5



conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa - Gt(ROSA)26Sor^{tm30.1(CAG-C4b,-mScarlett)Ics}/Ics

1. Genotyping protocol and data

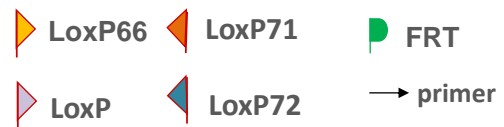
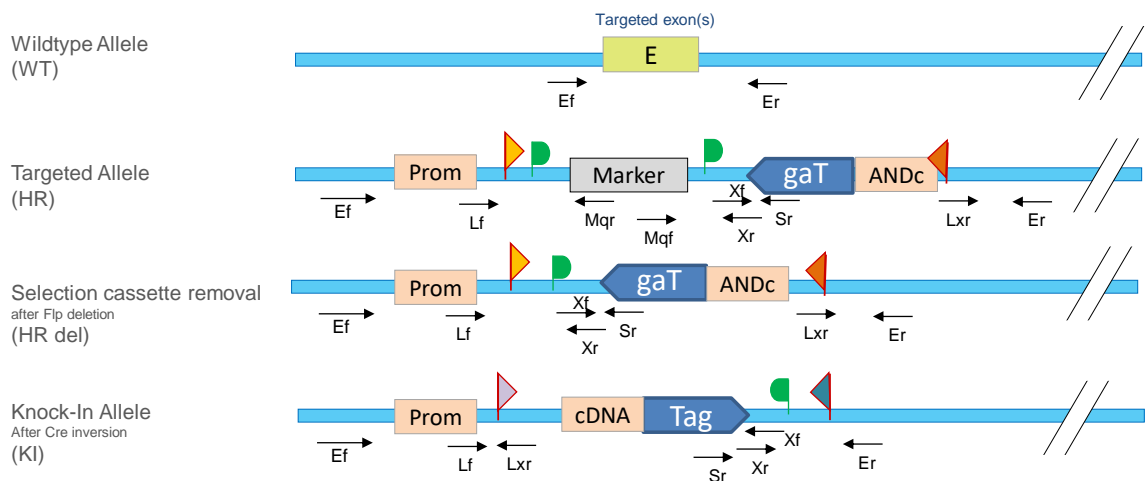
This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa** Conditional Knock In (Flex strategy) (KI-Flex) project.

1.1. Genotyping strategy

The map below describes the position of the primers used for genotyping for each possible allele. HR del allele corresponds to tm30.1



KI_FLEX Genotyping strategy



conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa - Gt(ROSA)26Sor^{tm30.1(CAG-C4b,-mScarlett)lcs}/Ics

Sequence of primers used for genotyping:

Position	Sequence
Ef	GCACTTGCTCTCCCAAAGTCGC
Er	CCGAGGCGGATCACAAGC
Lf	GGCAACGTGCTGGTTA
Lxr	GGCCGAATTCATCGATATAACTTCG
Sr	GGCATGGACGAGCTGTACAAG
Mq1f	AGGGCCAGCTCATTCTCCCACTC
Mq1r	TGCTAAAGCGCATGCTCCAGACTGC
Xf	CATTGATGAGTTTGGACAAACAC
Xr	CTGCATTCTAGTTGTGGTTTGTC

PCR fragments expected size (bp):

Region analyzed	Position on the primer (see the map above)	Targeted allele (HR)	HR del allele	KI allele	WildType allele
WildType allele specific PCR (3' part of the targeted locus)	Ef / Er	10354*	8501*	8535*	449
Excision of the selection marker	Lf / Sr	2272*	419	---	---
5' part of the selection marker	Lf / Mq1r	190	---	---	---
3' part of the selection marker	Mq1f / Xr	199	---	---	---
Exogenous/cDNA specific PCR	Xf / Sr	261	261	261	---
LoxP specific PCR	Lf / Lxr	---	---	100	---

*: this PCR product will not be observed using our PCR genotyping conditions (see description below)

---: no Amplicon should be obtained



conditional overexpression of 5X HA-C4b-T2A-mScarlett in Rosa - Gt(ROSA)26Sor^{tm30.1(CAG-C4b,-mScarlett)lcs}/Ics

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	
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.

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.

