



Xiap (IR00002785 / E114 ICS internal reference) mouse line genotyping protocol

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For any question, please contact:

Mouse Clinical Institute – Institut Clinique de la Souris (ICS)

ICS genotyping service

1 rue Laurent Fries, BP 10142

67404 Illkirch Cedex France

Email: genotyping@igbmc.u-strasbg.fr

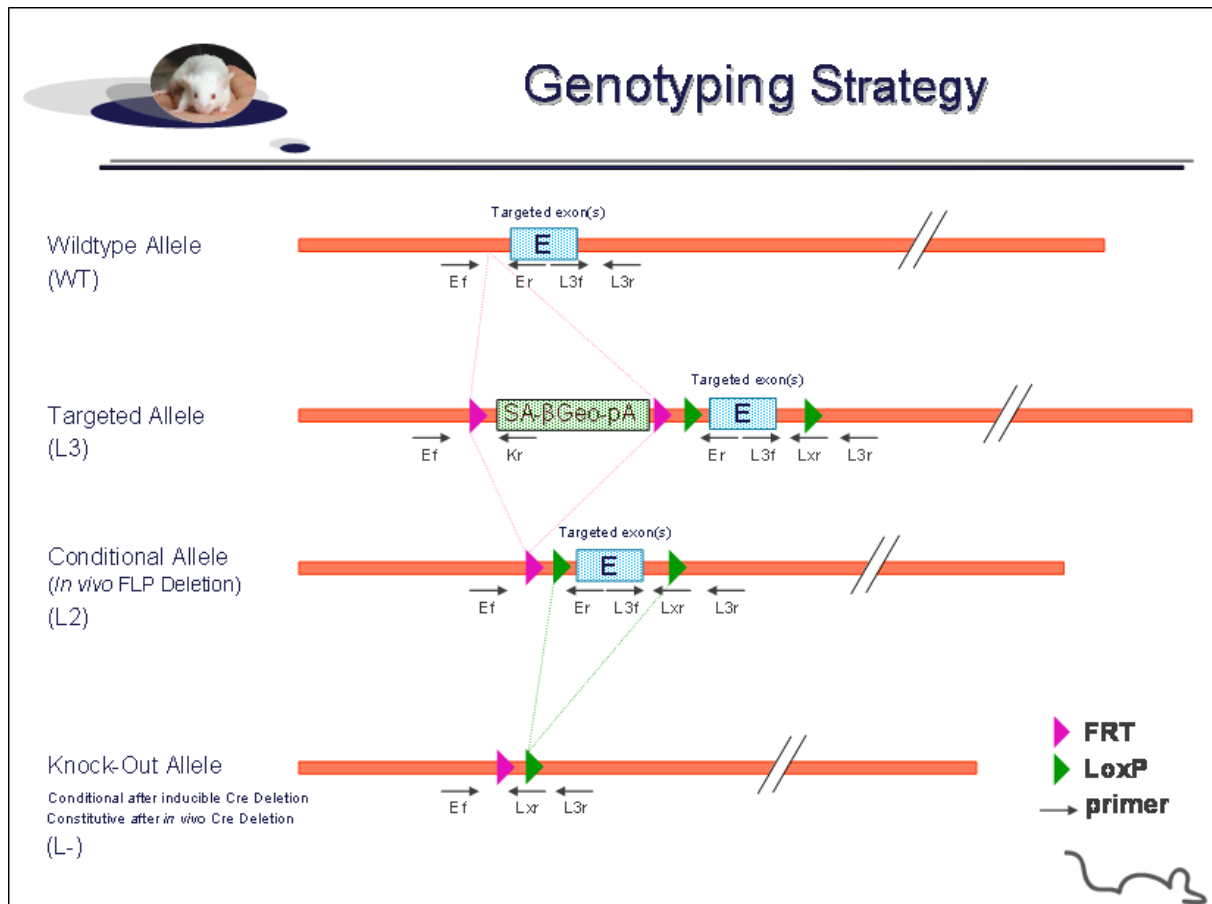
This protocol has been validated by Karim Essabri.

1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Xiap** Constitutive Knockout / Conditional Knockout (KO-cKO) 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	5250	GGCACAGACTATAGCAGAGTAGCC
Ef	5251	ATAGCAGTACTTTGTGGCTCCTCAG
Er	5252	CCATGCAATGGACAAACAAGT
Kr	3278	GGGCAAGAACATAAAGTGACCCCTCC
L3f	5253	GATAGTTAATGGAA TTGAATGCTGG
L3f	5254	TGCAAATACCTA TTGGATGAGAAGG
L3r	5255	CAGCTTGGTCTATG TGGGTCC
L3r	5256	CTGAATCACCAGCCAAA GAGC
Lxr	3254	TTATCATTAA TTGCGTTGCGCCA TC



Genotyping protocol

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PCR fragments expected size (bp):

Region analyzed	Primers used	Position on the primer (see the map above)	Targeted allele (KO allele) (L3)	cKO allele (L2)	KO allele (L-)	WildType allele (WT)
5' part of the selection marker	5251-3278	Ef / Kr	461	---	---	---
Presence of the distal loxP	5254-5255	L3f / L3r	392	392	---	392
Distal loxP specific PCR	5253-3254	L3f / Lxr	361	361	---	---
Excision of the selection marker	5250-5252	Ef / Er	7675*	772	---	637
Excision(s) of the floxed exon(s), i.e. knock out	5251-5256	Ef / L3r	8160*	1256*	542**	1122*

* 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/μl)
- 5' primer (100 μM)
- 3' primer (100 μM)
- Sterile H₂O

Volume:

- 7.5μl
- 1.5μl
- 0.06μl
- 0.06μl
- up to 15 μl

Cycling conditions:

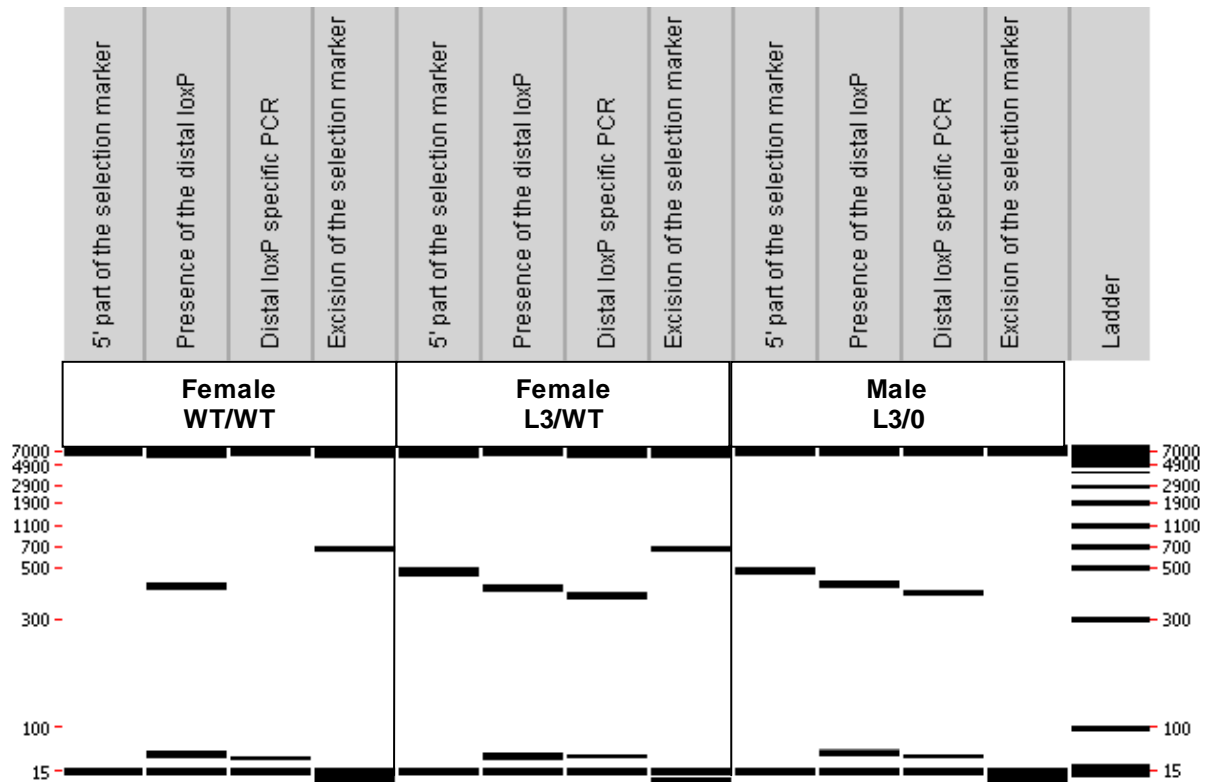
Temp	Time	#Cycles
95°C	4min	1
94°C	30s	34
62°C	30s	
72°C	1min	
72°C	7min	1
20°C	5 min	1

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 not done by gel electrophoresis but using LabChip® 90 microfluidic apparatus. PCR products were run on the HT DNA 5K LabChip® 90 Assay Kit.

Representative genotyping picture



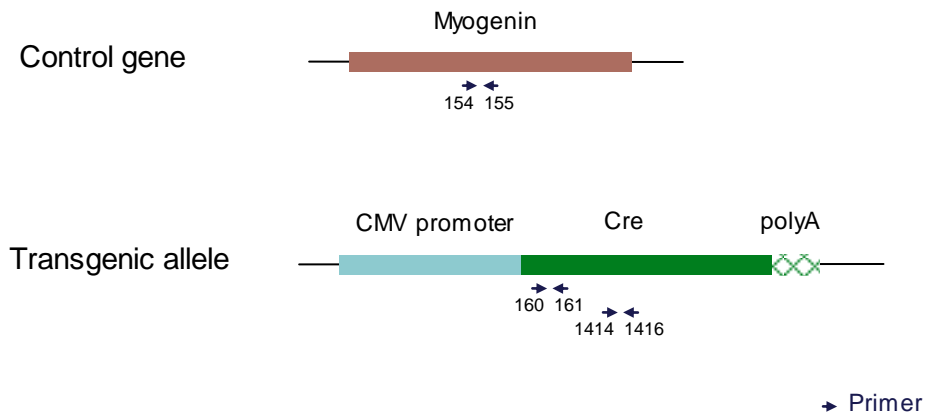
Note that as this technology is more sensitive than gel analysis, non specific signals and/or primer dimers may be visible on the picture.

2. Cre and Flp genotyping method

The protocol used to segregate the cre and/or flp transgene is indicated below.

2.1. Cre genotyping

Schematic representation of the genotyping strategy



PCR fragments expected size (bp):

Sequence of primers used for genotyping

Primers	Sequence
154	ACTCCCTTACGTCCATCGTG
155	ACCCAGCCTGACAGACAATC
160	GAACCTGATGGACATGTTTCAGG
161	AGTGCGTTCGAAACGCTAGAGCCTGT
1414	CGTACTGACGGTGGGAGAAAT
1416	CCCGGCAAAACAGGTAGTTA

Primer pair	160-161	1414-1416	154-155
Region analyzed	5' part of Cre transgene	Middle of Cre transgene	Myogenin control gene
Control gene	/	/	99
Tg allele	345	165	/

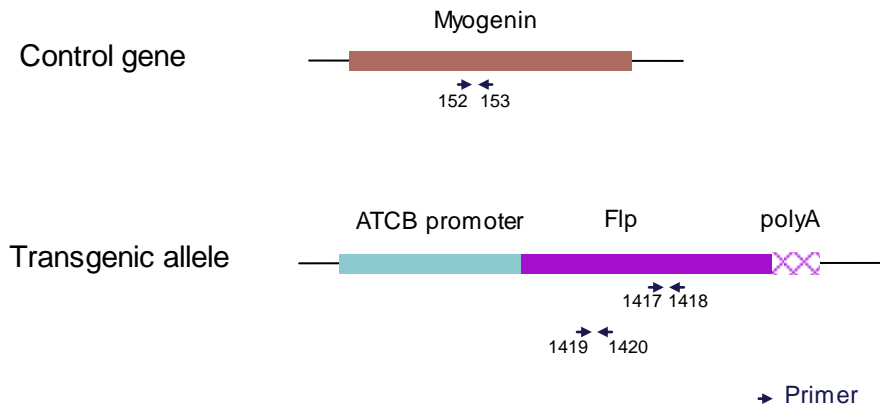
Cycling conditions:

Temp	Time	#Cycles
95°C	3min	1
95°C	10s	35
62°C	20s	
72°C	20s	
95°C	5s	1 (melting curve generation)
62°C	30s	
72°C	72s	
37°C	30s	1
4°C	∞	

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

2.2. Flp genotyping

Schematic representation of the genotyping strategy



Sequence of primers used for genotyping

Primers	Sequence
152	TTACGTCCATCGTGGACAGC
153	TGGGCTGGGTGTTAGCCTTA
1417	TTCTTTAGCGCAAGGGGTAG
1418	GCTCCAATTTCCCACAACAT
1419	TGGGAAATTGGAGCGATAAG
1420	CTGCCACTCCTCAATTGGAT

PCR fragments expected size (bp):

Primer pair	1417-1418	1419-1420	152-153
Region analyzed	Middle part of Flp transgene	5' of Flp transgene	Myogenin control gene
Control gene	/	/	245
Tg allele	299	175	/

PCR protocol and cycling conditions are identical to those described in chapter 1.2