

Genotyping protocol

General information:

Strain name	tatdn1 floxed
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Primers:

Name	Sequence	Primer type
TatdloxP_Fw	CTACTCTTAAAGACAGCCTTCC	
TatdNeo_Fw	GGTGGATGTGGAATGTGTGC	
TatdNeo_Rv	CTTTATTTCTCAGATGTGTGGGC	

In case more than two primers are introduced, please indicate how they should be combined:

	Forward primer	Reverse primer
wt		
mutant		

Reaction mix:

ddH ₂ O	18,5	μl
PCR Buffer 15mM MgCl ₂	2,5	μl
dNTPs 2,5mM	2,5	μl
Primer 100μM	0,125	μl
Primer 100μM	0,125	μl
Taq Polymerase 5U/μl	0,250	μl
DNA	1	μl
Final volume	25	μl

PCR program:

95	°C	5	min	X35
95	°C	30	sec	
60	°C	30	sec	
72	°C	30	sec	
72	°C	10	min	

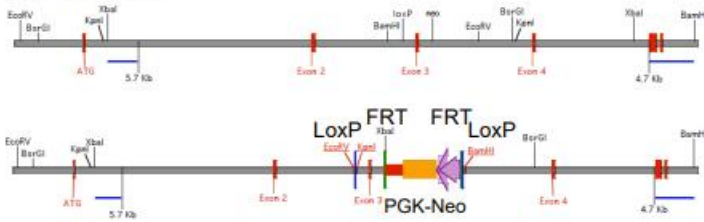
Expected fragment size:

LoxP -/-	270 bp
LoxP +/+	441 bp
Neo +/+	
LoxP -/-	400 bp
Neo -/-	

Comments/Additonal information:

Tatdn1 floxed mice genotyping

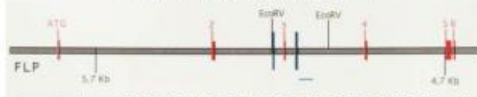
Wild type allele



Targeted allele

Tatd loxP^{+/+} Neo^{+/+} x ROSA FLPe

Conditional allele after FLPe



Tatd loxP^{+/+} Neo^{-/-} x CAG-Cre

Conditional allele after Cre (*Tatd* KO)



Genotyping: FWD1 (5'Tatd) FWD2(PGK) REV (3'Tatd)

Allele		Primer sequence	PCR conditions			PCR product	
			T ^a (°C)	Time	Cycles	Genotype	size (bp)
<i>Tatd</i> <i>loxP</i> <i>neo</i>	Fwd	CTACTCTTAAAGACAGCCTTCC	95	3 min	1	<i>loxP^{+/+}</i> <i>neo^{+/+}</i>	441
	Fwd	GGTGGATGTGGAATGTGTGC	95	30 sec		<i>loxP^{+/+}</i> <i>neo^{-/-}</i>	400
	Rev	CTTTATTTCTCAGATGTGTGGGC	60	30 sec	35	<i>loxP^{-/-}</i>	270
			72	45 sec			
			72	2 min	1		
<i>CAG</i> <i>cre</i>	Fwd	AGGTTTCGTTCACTCATGGA	95	2 min	1	<i>Cre^{+/+}</i>	-
	Rev	TCGACCAGTTTAGTTACCC	95	1 min		<i>Cre^{+/+}</i>	250
			55	1 min	30		
			72	1 min			
			72	10 min	1		
<i>Tatd</i>	Fwd	GGGCTCCAGGTTCAAGTGGAG	95	3 min	1	<i>Tatd^{+/+}</i>	270
	Rev	CTTTATTTCTCAGATGTGTGGGC	95	1 min		<i>Tatd^{-/-}</i>	500
	Fwd	CTACTCTTAAAGACAGCCTTCC	60	1 min	35		
			72	1 min			
			72	10 min	1		
	Fwd	ACGAACCTGGTCGAAATCGTGCG	94	2 min	1	<i>Cre^{-/-}</i>	-
	Rev	CGGTCGATGCAACGAGTGATGAG	94	45 sec		<i>Cre^{+/+}</i>	350

