

Supplementary Table S1. Primers for PCR and Table2. Primer for qPCR

Amplicon	Expected size	Primer pair	Primer name	Sequences 5'→3'
F1A/RFP	537 bp	Forward	F1A(m)-F	gccacctgttaagtggccag
		Reverse	Rfp-R	cgtcagcggggtacagcatc
Cre	562 bp	Forward	CreF	ctaaactagcttcacgtcggtc
		Reverse	CreR	tctgaccagagtcaccttagcg
LacZ	336 bp	Forward	LacZF	actggccgtcggtttacaacg
		Reverse	LacZR	ttctccgtgggaacaaacgg
R26R_ wild type	603 bp	Forward	R26F2	aaagtcgctctgaggttggtat
		Reverse	R523	ggagcggagaaatggatatg
R26R_ transgene	340 bp	Forward	R26F2	aaagtcgctctgaggttggtat
		Reverse	R1295	gcgaagagttgtcctcaacc
GAPDH	500 bp	Forward	Gapdh-F3	aaatggtgaaggtcggtgtga
		Reverse	Gapdh-R3	tcataacggcggttcattcat
PGK neo 4xpA Floxed	~800 bp	Forward	R26F2	aaagtcgctctgaggttggtat
		Reverse	LacZR	ttctccgtgggaacaaacgg

Supplementary Table S2. PCR results of R26F2/LacZR, R26F2/R1295, R26F2/LacZR, R26F2/R1295, R26F2/LacZR and R26F2/R1295 (internal control)

Amplicon	Expected size	Primer pair	Primer name	Sequences 5'→3'
F1A/exon1	499 bp	Forward	F1A-Fw	cccaaagccaagaagccacc
		Reverse	Exon1-Rv	tgtgctggtcgctcctgtccct
Cre	195 bp	Forward	Cre-Fw	gatttcgaccaggttcggtc
		Reverse	Cre-Rv	gctaaccagcggtttcggtc
RFP	333 bp	Forward	RFP-Fw	tgagaatcaaggtggtcgag
		Reverse	RFP-Rv	cgtcagcggggtacagcatc
β-actin	233 bp	Forward	β-actin-Fw	ctaggcaccaggggtgtgatg
		Reverse	β-actin-Rv	gttggccttaggggtcaggg

PCR conditions

[Kapa LongRange Hotstart Reagent ReadyMix \(kk3601\)](#)

Amplicon: F1A/Cre (2217 bp) :

PCR program: 1 cycle of 95 °C for 5 min, 35 cycles of 95 °C for 15 sec, 55 °C for 15

sec, 68 °C for 10 min and final 68°C for 10 min °

[Kapa Taq ReadyMix with dye \(kk1024\)](#)

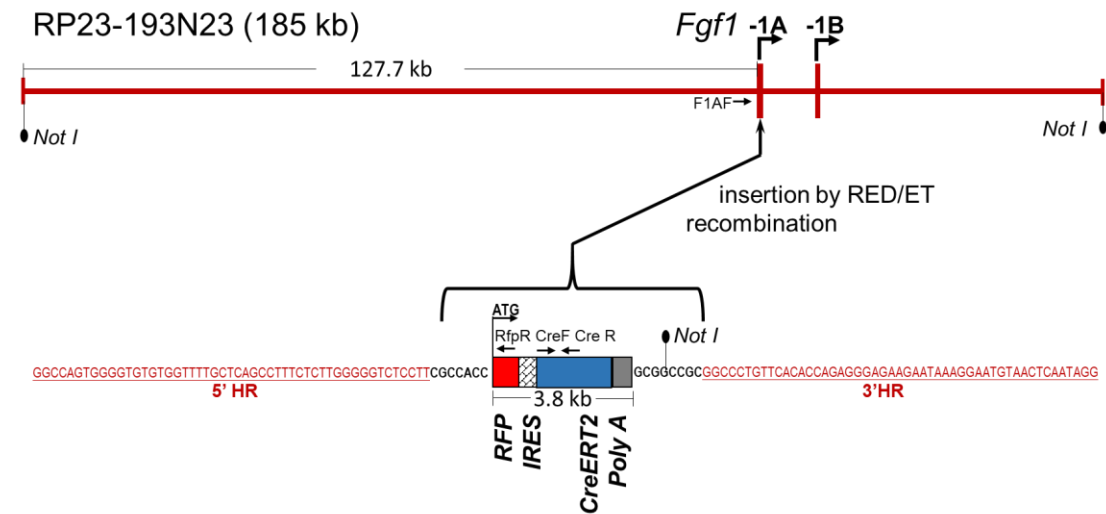
Amplicon: F1A/RFP 、 Cre 、 LacZ 、 RFP : PCR program: 1 cycle of 95 °C for 5 min, 35 cycles of 95 °C for 30sec, 56 °C for 40 sec, 72 °C for 1 min and final 72 °C for 7 min.

Amplicon: actin 、 GAPDH: 1 cycle of 95 °C for 5 min, 20 cycles of 95 °C for 20sec, 55 °C for 30 sec, 72 °C for 50 sec and final 72 °C for 7 min.

qPCR : [KAPA SYBR® FAST qPCR Master Mix \(2X\) Kit](#)

Gene expression detection and data analysis were performed using ABI7500 1.41 version software (ABI). The program is 1cycle of 50 °C for 2min, 1 cycle of 95 °C for 10min, 40 cycles of 95 °C for 15 sec, 60°C for 1min and 1cycle of 95 °C for 15sec 60 °C for 1min 95 °C for 15 sec 60 °C for 15 sec.

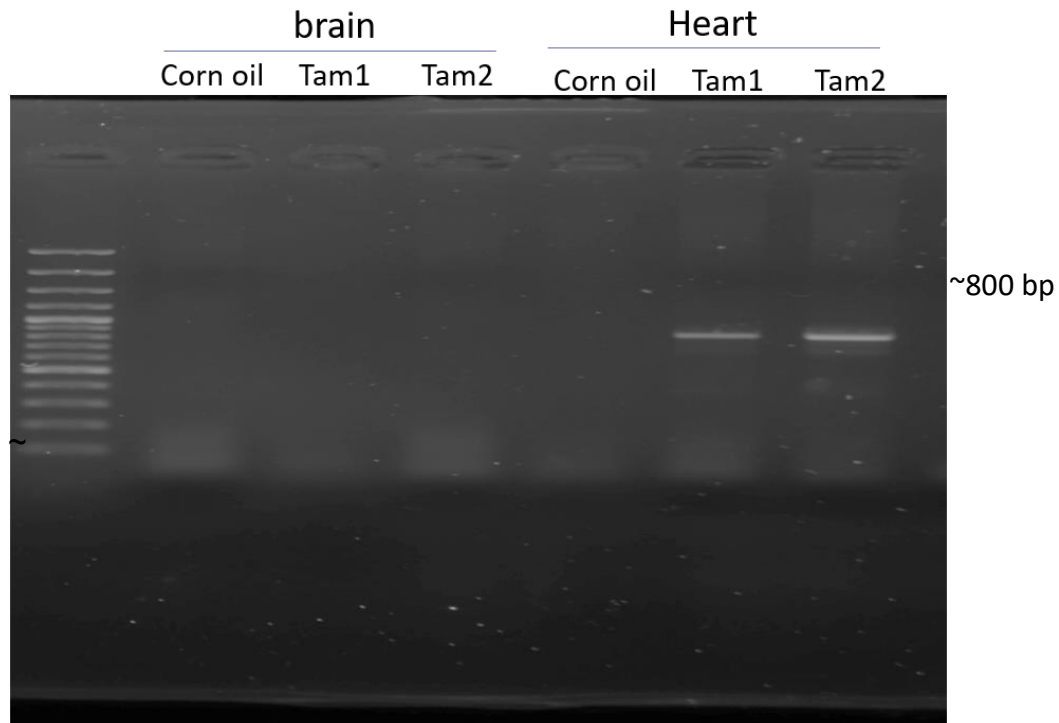
Supplementary Figure S1:



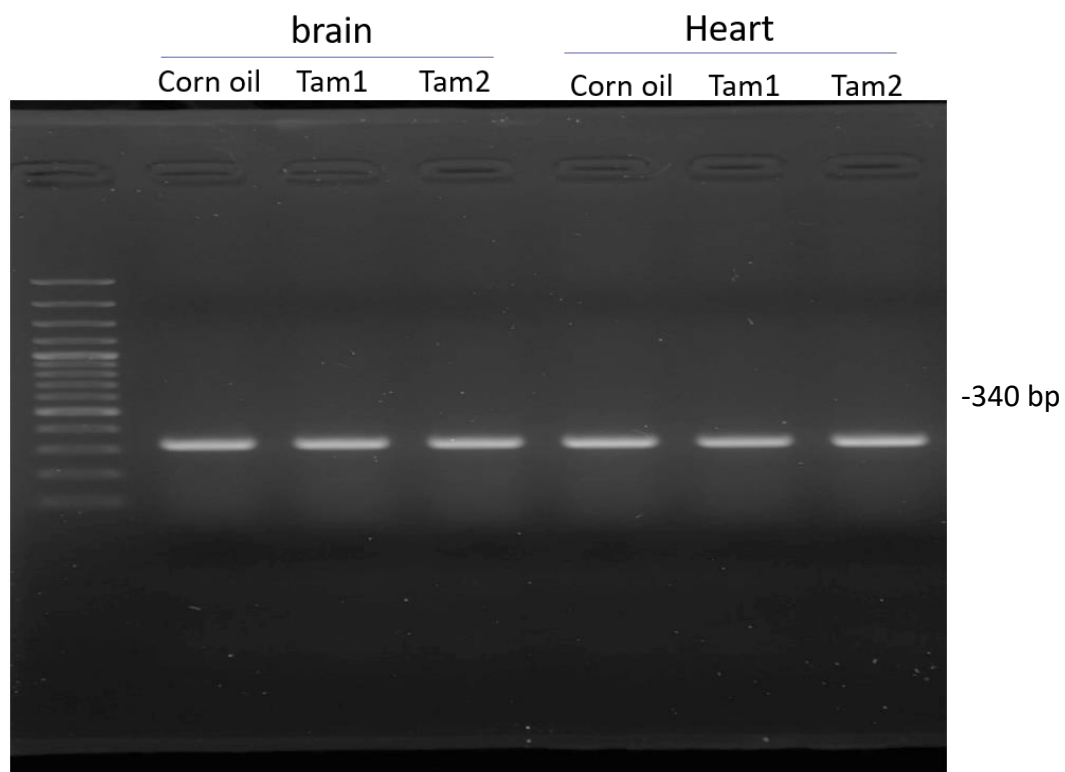
Supplementary Figure S1: Construction map of the mouse *Fgf1A-CreERT^{T2}* transgene. The transgene was constructed using mouse BAC clone RP23-193N23 including *Fgf1A* 5'UTR (127.7 kb), exon 1A, and 1B sequence as the transgene backbone. The bicistronic expression cassette RFP-IRES-CreER^{T2}-polyA containing an upstream Kozak sequence and a downstream *Not I* site was inserted into the exon 1A by RED/ET recombination system. The modified BAC was purified, *Not I* digest, and pulsed-field gel electrophoresis to isolate the transgene including the *Fgf1A* 5'UTR and expression cassette for mouse pronuclear microinjection. The junction sequences flanking the insertion site were also shown. Primers designed for genotyping: F1AF, RfpR, CreF and CreR.

Supplementary Figure S2. PCR results of R26F2/LacZR, R26F2/R1295, R26F2/LacZR, R26F2/R1295, R26F2/LacZR and R26F2/R1295 (internal control)

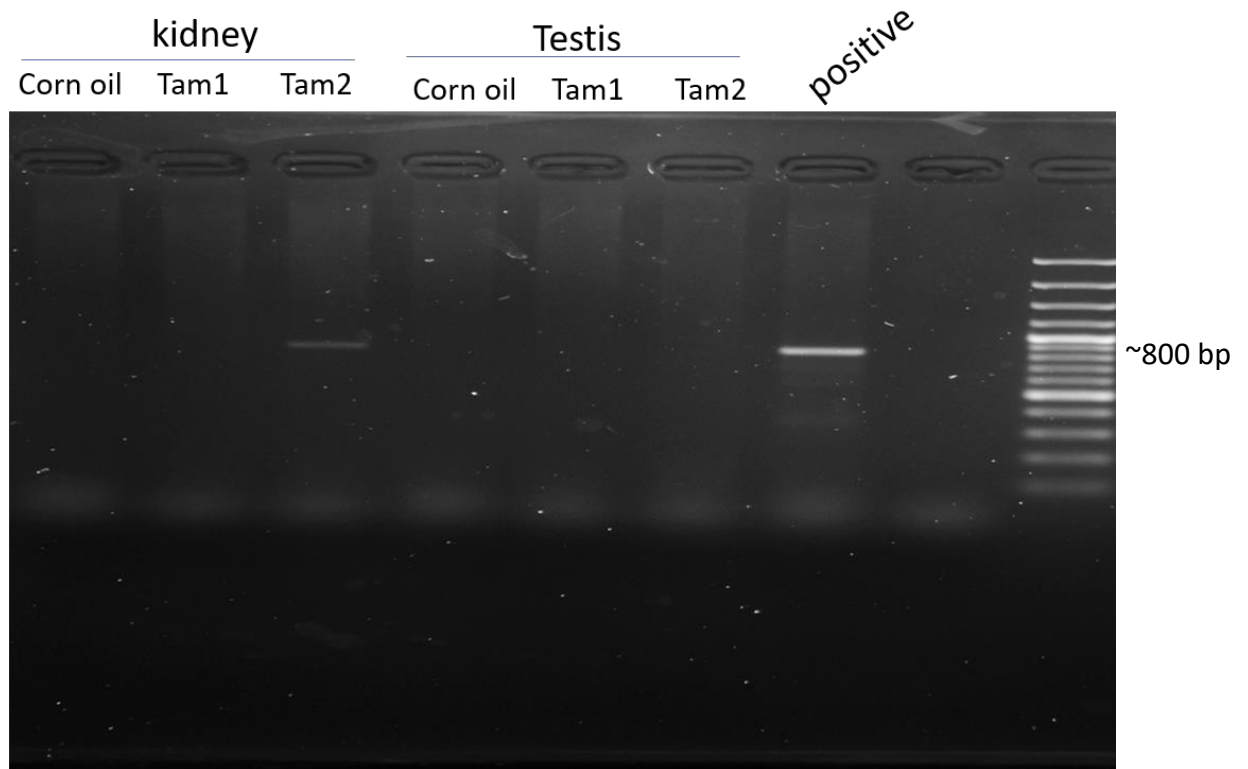
Primer pairs: R26F2/LacZR



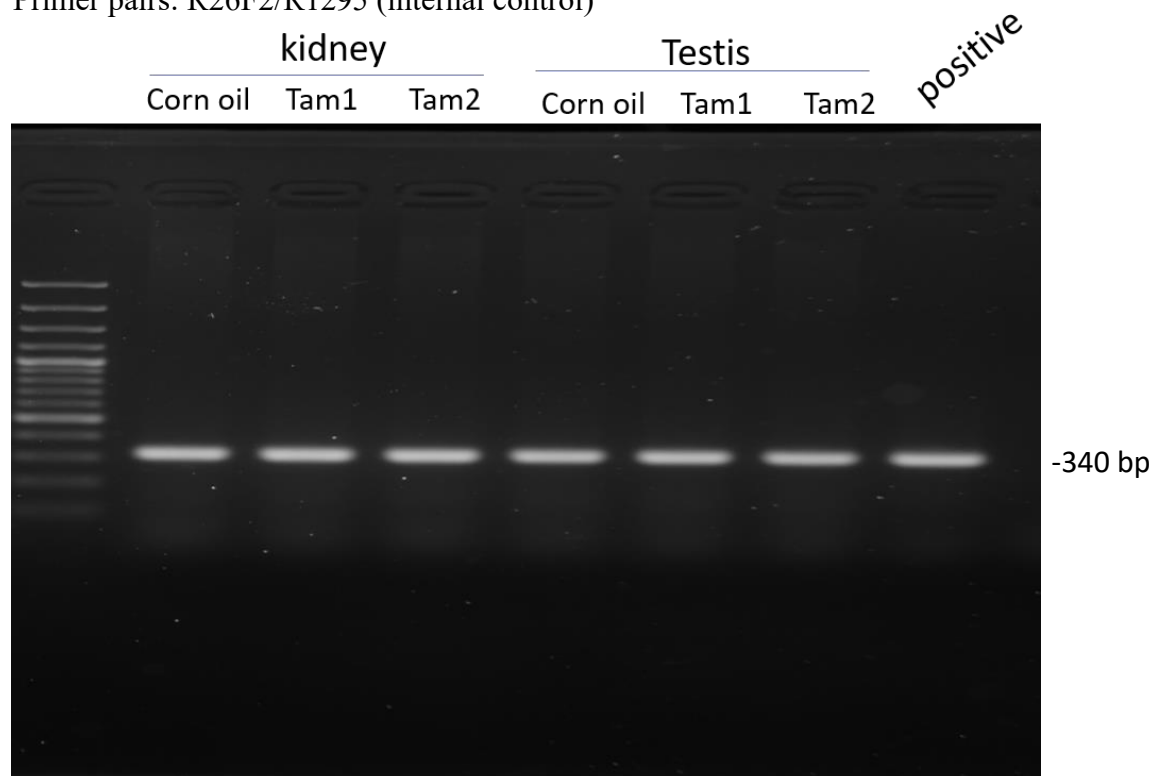
Primer pairs: R26F2/R1295 (internal control)



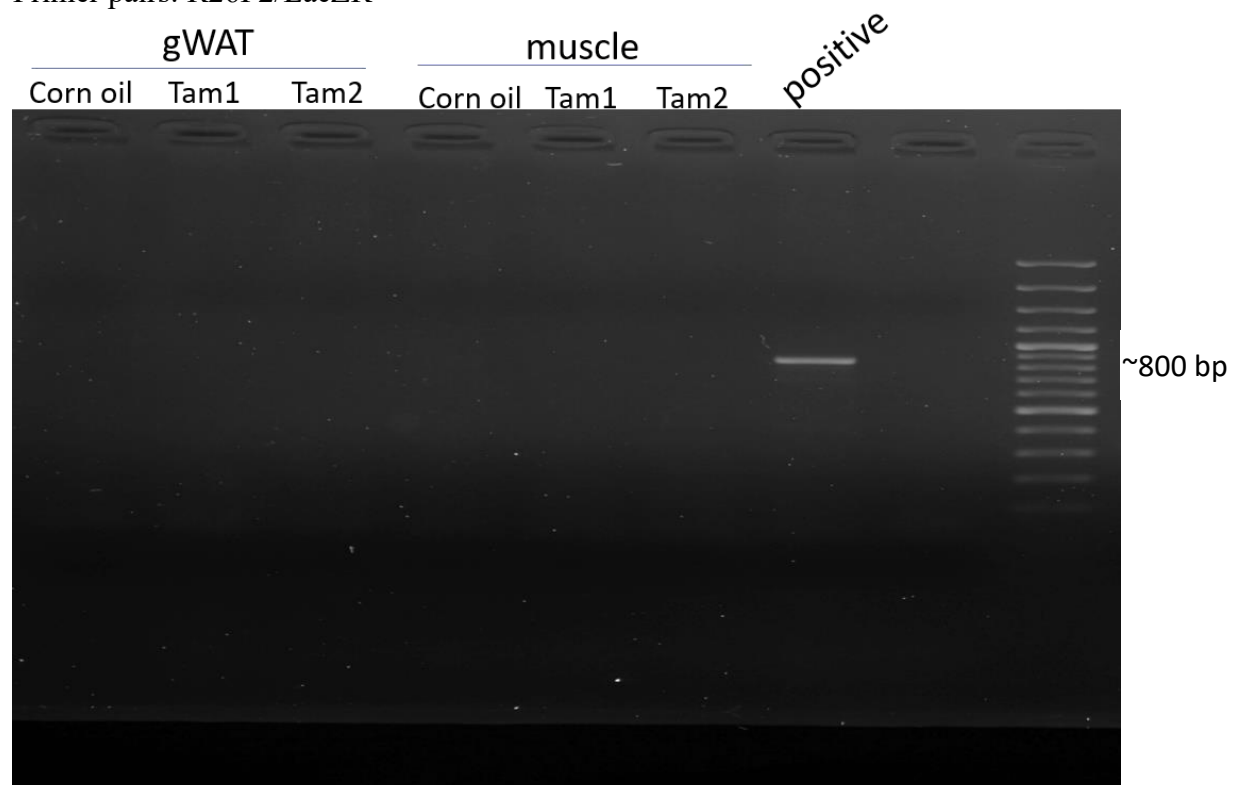
Primer pairs: R26F2/LacZR



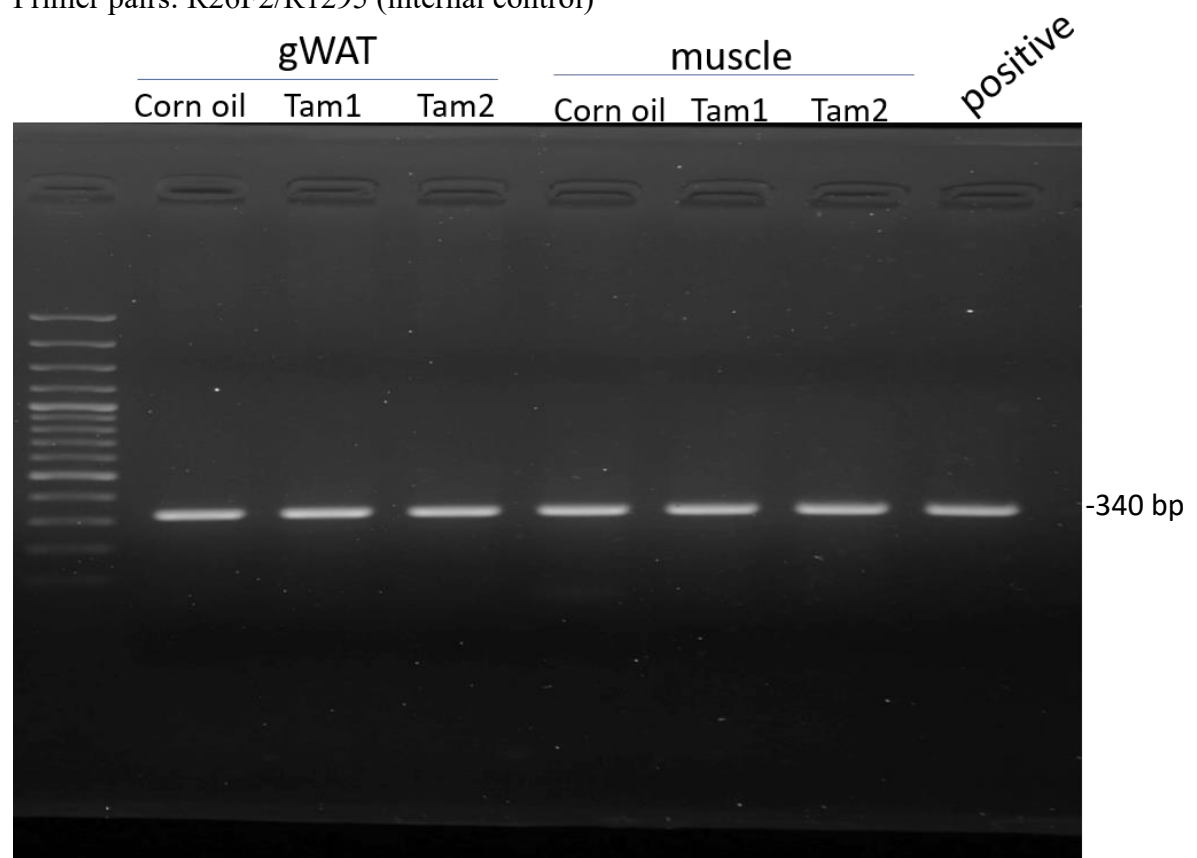
Primer pairs: R26F2/R1295 (internal control)



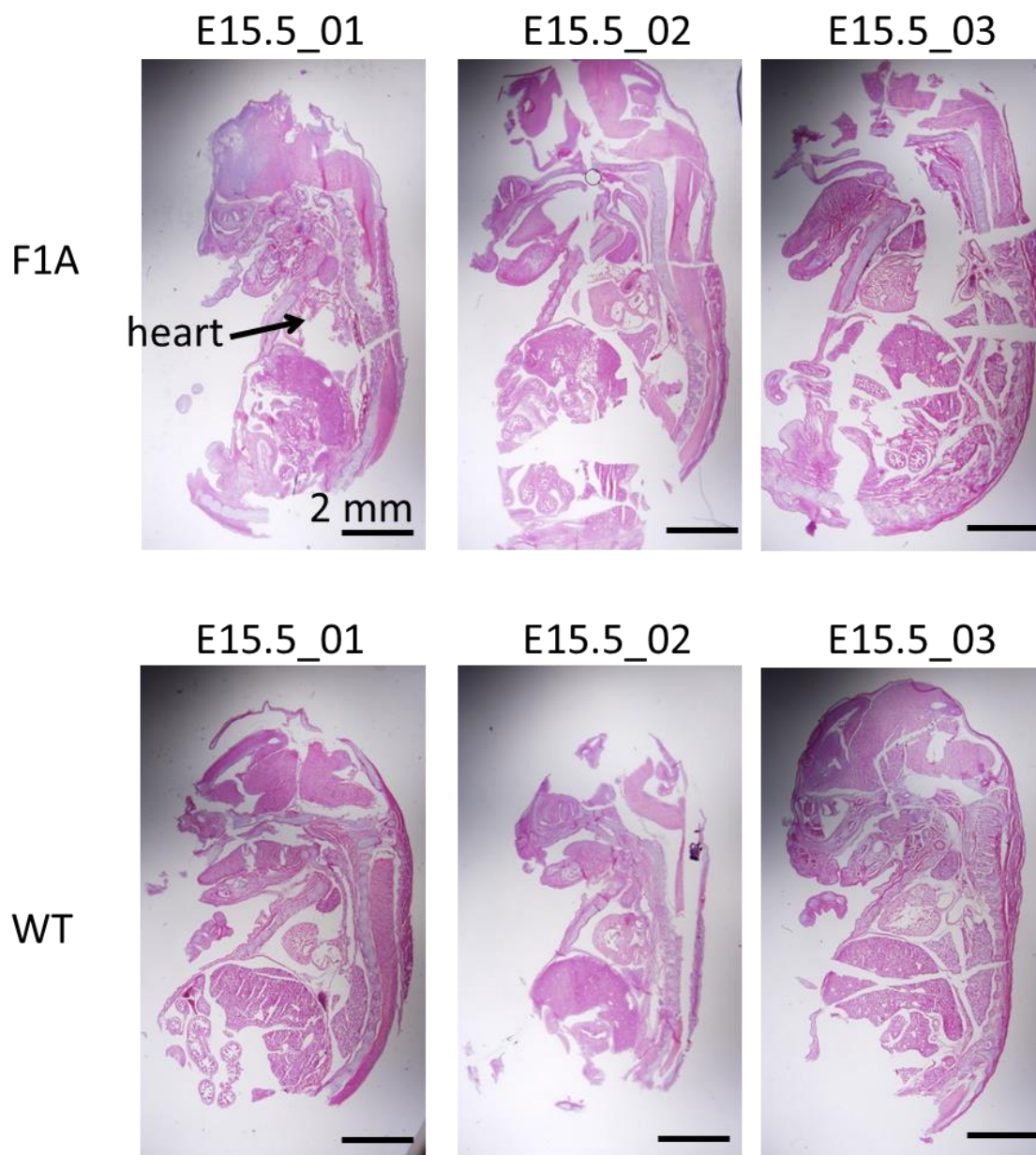
Primer pairs: R26F2/LacZR



Primer pairs: R26F2/R1295 (internal control)

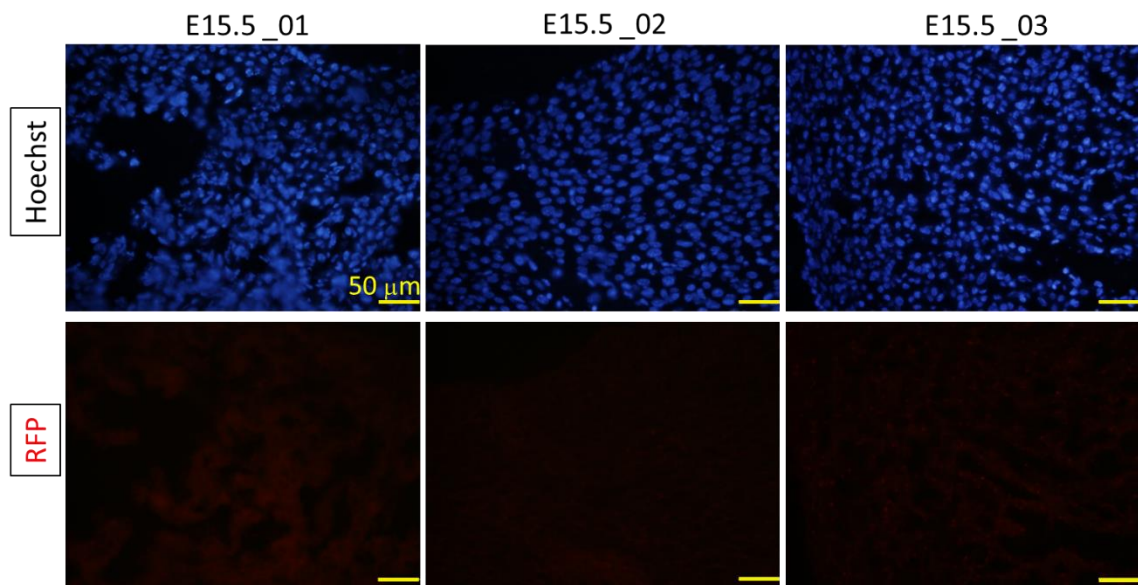


Supplement Figure S3. Immunohistochemical staining in the heart of E18 embryo
A.



B.

F1A



C.

WT

