

C57BL/6JGpt-Tek-P2A-iCre

Strain Name: C57BL/6JGpt-*Tek*^{em1Cin(P2A-iCre)}/Gpt

Strain Type: Knock-in

Strain Number: T055033

Background: C57BL/6JGpt

Description

This mouse strain expresses codon optimized iCre recombinase [1] under the control of the mouse endogenous *Tek* promoter, P2A-iCre was inserted upstream of the stop codon TAG of mouse *Tek* gene by CRISPR/Cas9 technology. When crossed with a strain with loxP site flanked sequence in its genome, Cre-mediated recombination will result in excision of the DNA fragment between the two loxPs in endothelial cells and hematopoietic cells. Recombinase activity was detected in tissues rich in hematopoietic cells and endothelial cells, such as thymus, spleen, kidney, aorta, liver, lung, intestine, etc. Note: mild sperm activity was detected by PCR detection of loxP recombination.

Strategy

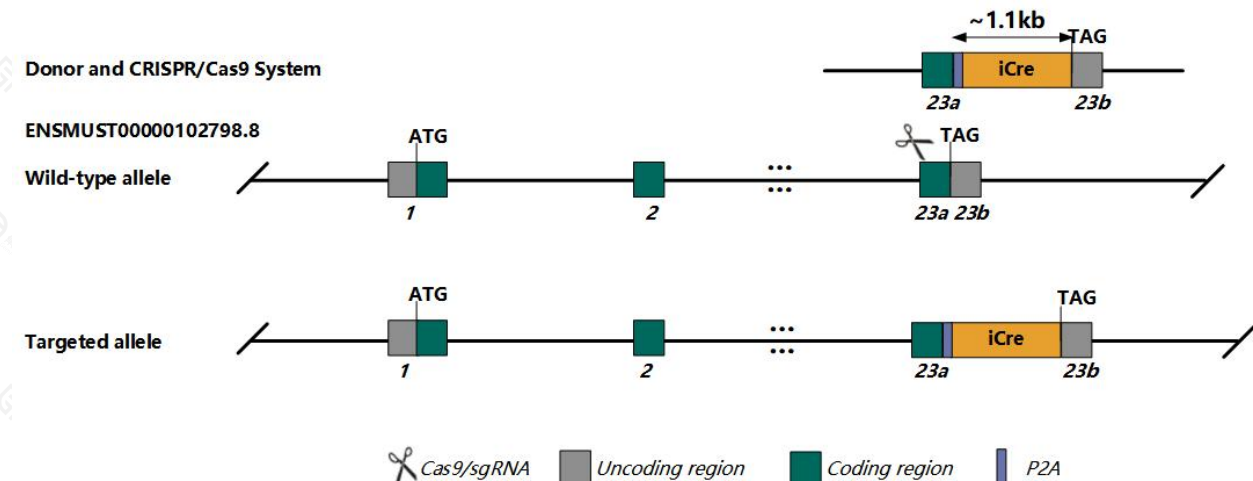


Fig.1 Schematic diagram of C57BL/6JGpt-Tek-P2A-iCre model strategy.

Applications

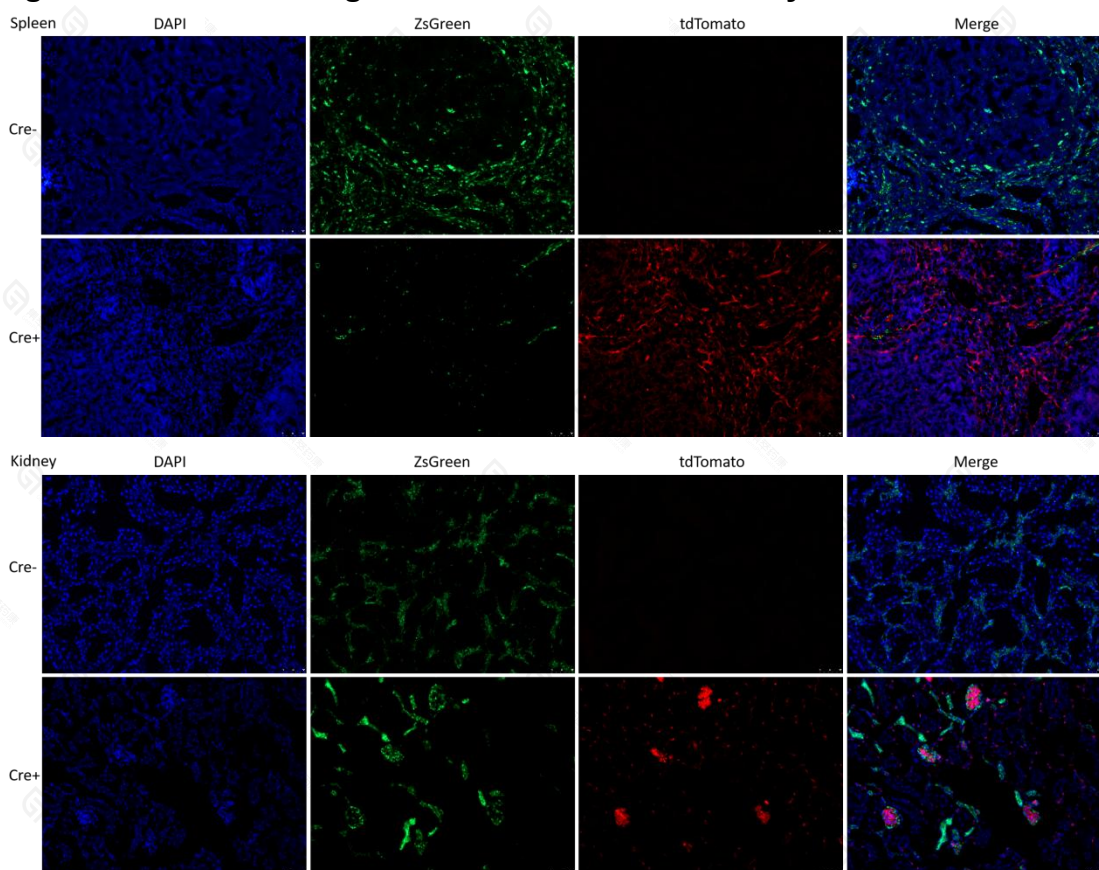
1. Cre tool mice for specific induction of loxP recombination in endothelial cells and hematopoietic cells [2-3].

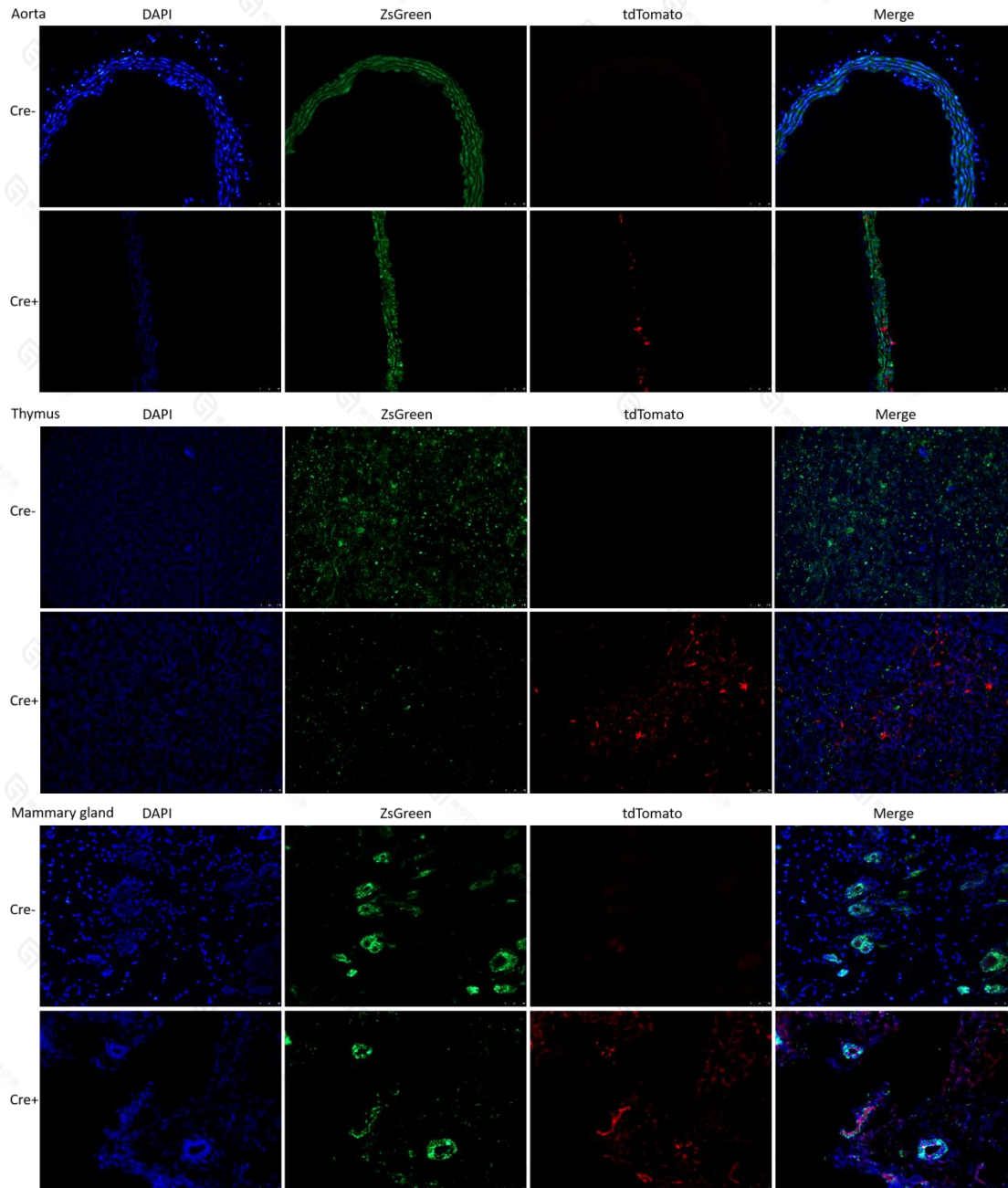
Data support

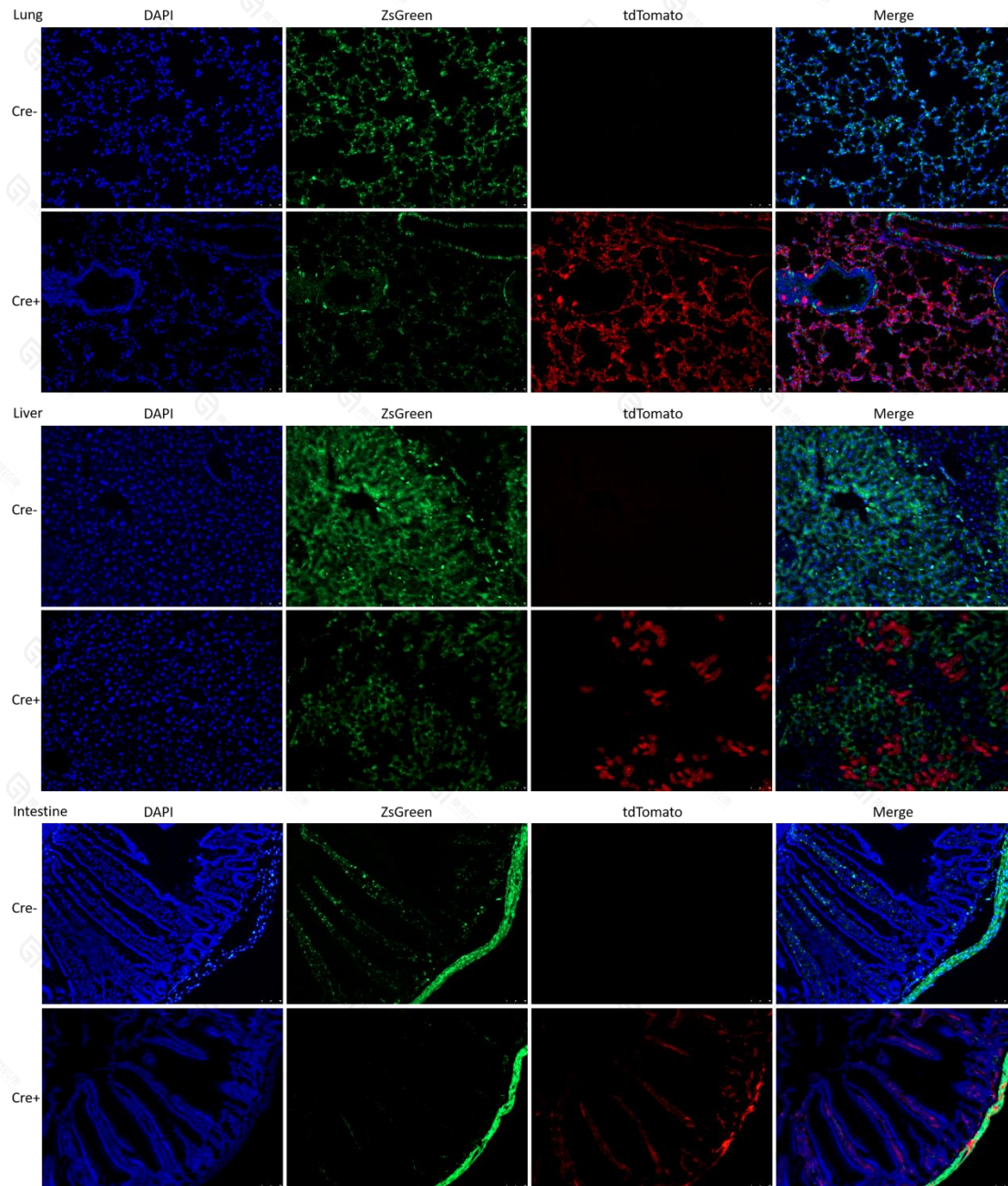
1. Validation methods & notes

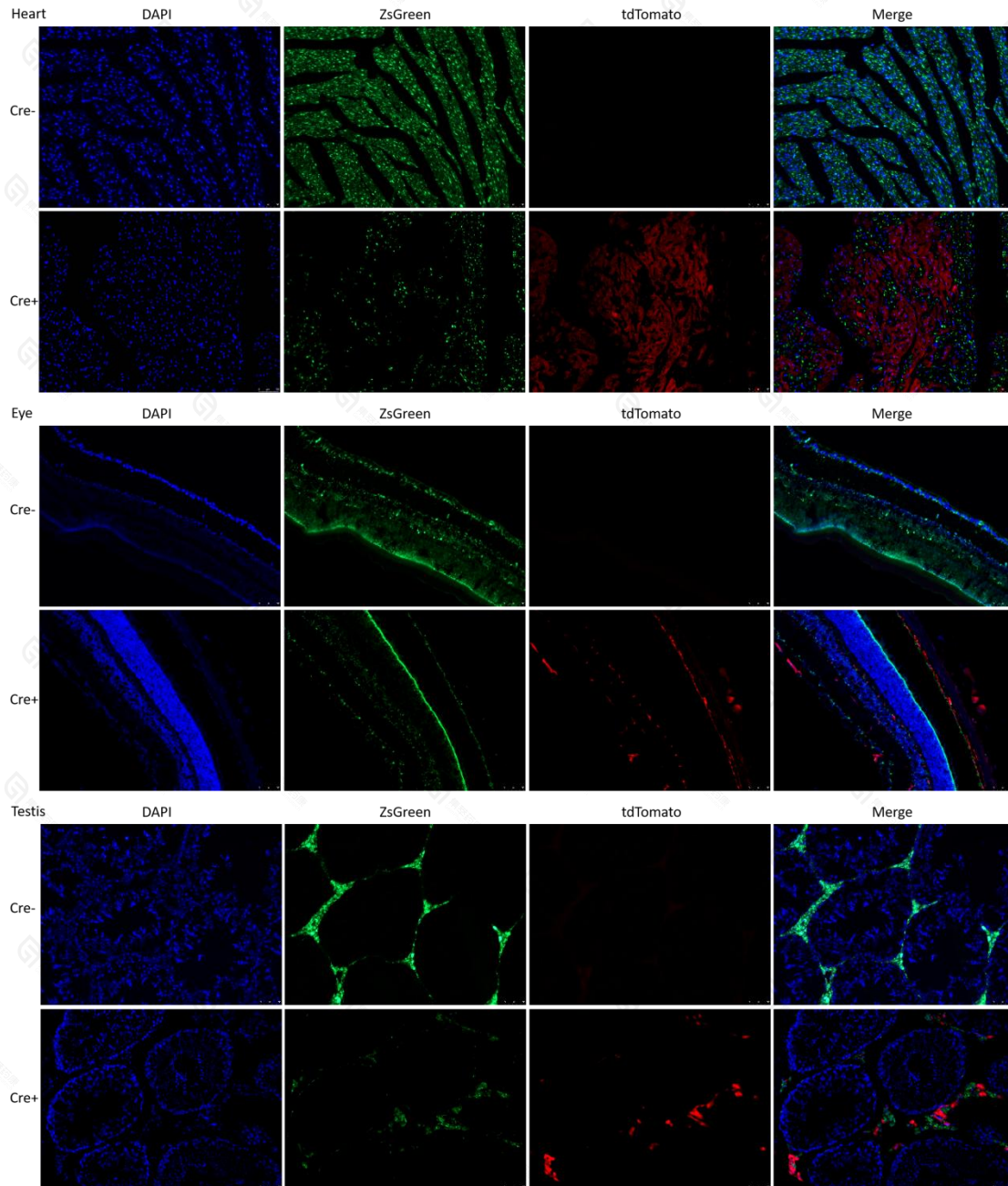
Tek-P2A-iCre mice was crossed with CAG-loxp-ZsGreen-Stop-loxp-tdTomato mice with ubiquitous reporter expression (hereafter referred as CAG-G/R mice), Cre-mediated recombination will lead to excision of ZsGreen and the stop cassette and expression of tdTomato, thus loss of green fluorescence and gain of red fluorescence will indicate Cre activity. Fluorescence imaging of frozen sections were performed to exhibit Cre activity in various tissues and organs. Imaging of sections were performed under a 200x microscopy. Note: these results may only represent the activity of Cre in this strain at the identical stage. Recombinase activity may be different at other stages in your application.

2. Images of tissues and organs with obvious Cre activity









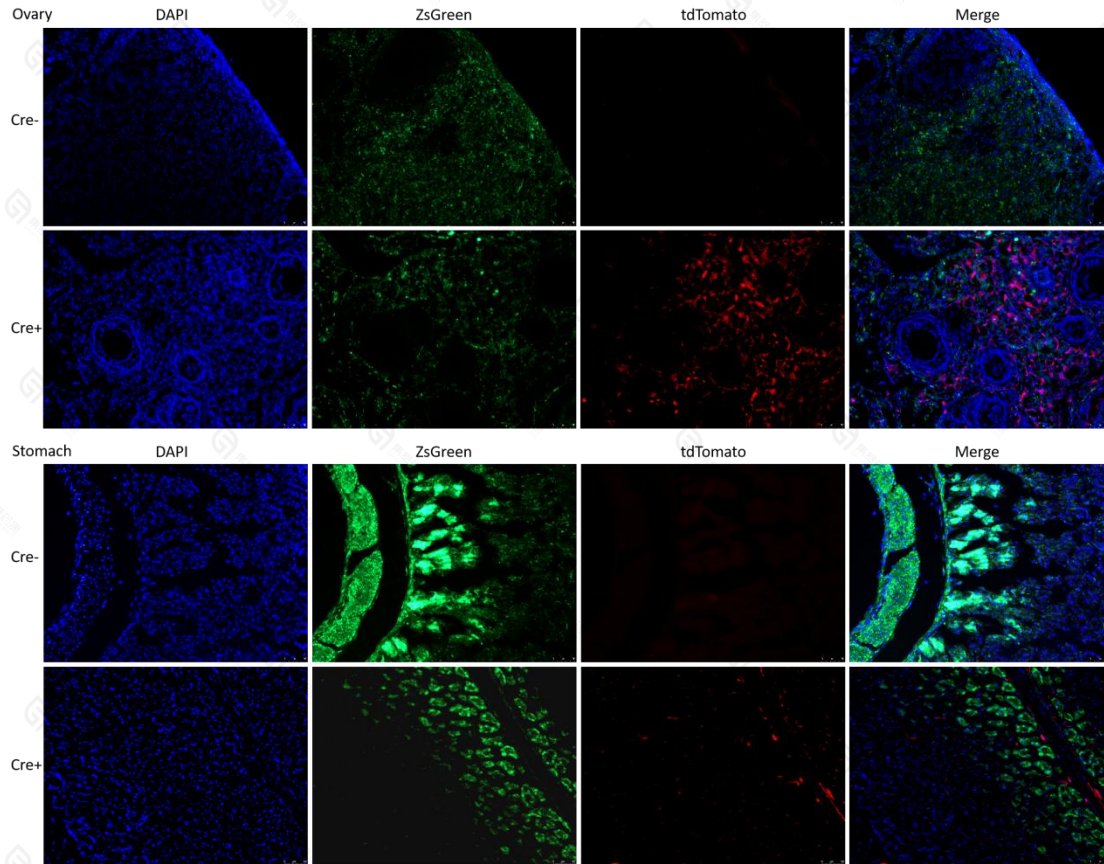
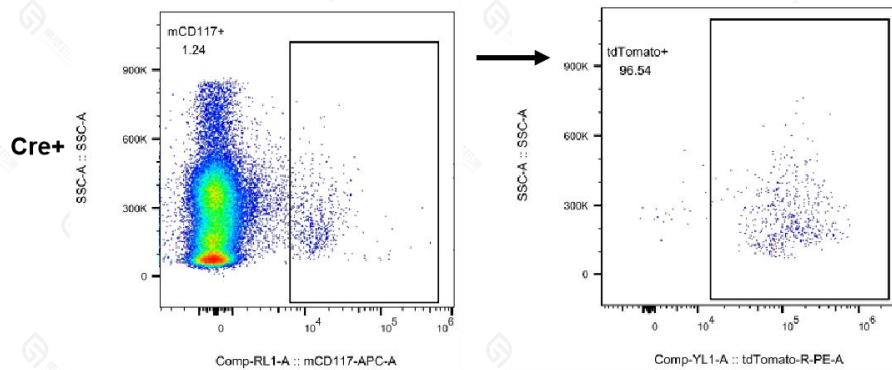


Fig 2. Fluorescence imaging of tissues and organs with obvious Cre activity.
Organ name was indicated in the left top of each subfigure group. Cre-: CAG-G/R single positive individuals; Cre+: Tek-P2A-iCre, CAG-G/R double positive individuals.

3. Flow cytometry analysis of cells with Cre activity

Bone marrow: mCD117+ cells



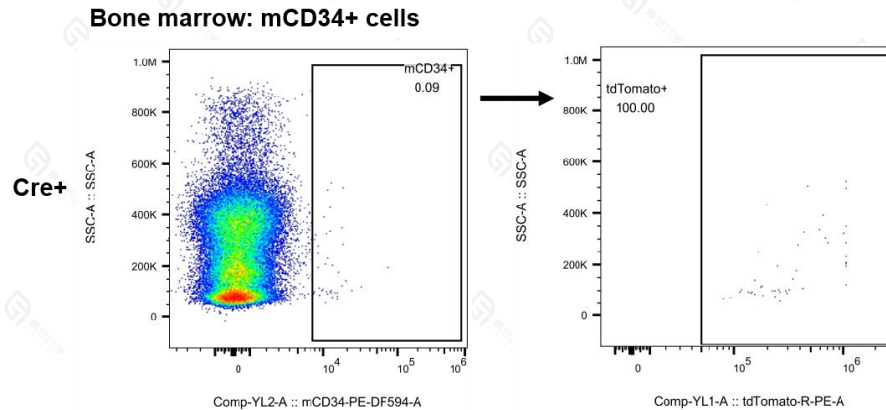


Fig 3. Flow cytometry analysis of cells with Cre activity

Organ name was indicated in the left top of each subfigure group. Cre+: Tek-P2A-iCre, CAG-G/R double positive individuals. Bone marrow-derived cells were harvested from Cre+ mice and analyzed for tdTomato expression with flow cytometry.

Reference

1. Shimshek D R, Kim J, Hübner M R, et al. "Codon-improved Cre recombinase (iCre) expression in the mouse." *genesis* 2002, 32(1): 19-26.
2. Kisanuki YY, Hammer RE, Miyazaki J, et al. Tie2-Cre transgenic mice: a new model for endothelial cell-lineage analysis in vivo. *Dev Biol*, 2001, 230(2): 230-42.
3. Batard P, Sansilvestri P, Scheinecker C, et al. The Tie receptor tyrosine kinase is expressed by human hematopoietic progenitor cells and by a subset of megakaryocytic cells. *Blood*, 1996, 87(6): 2212-20.