

The Gp1ba-Cre transgenic mouse

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Supplemental Table 1

Supplemental Table 1.

The *Pf4-Cre* mouse model was utilized in over 160 studies to date.

Supplemental Table 2

Supplemental Table 2.

Mendelian frequencies of *Gp1ba-Cre* mice.

Genotype	Total number of mice	<i>Expected frequency</i>	<i>Actual frequency</i>
<i>Gp1ba-Cre</i> ^{+/+}	26	25%	20.8%
<i>Gp1ba-Cre</i> ^{+K1}	67	50%	53.6%
<i>Gp1ba-Cre</i> ^{K1/K1}	32	25%	25.6%

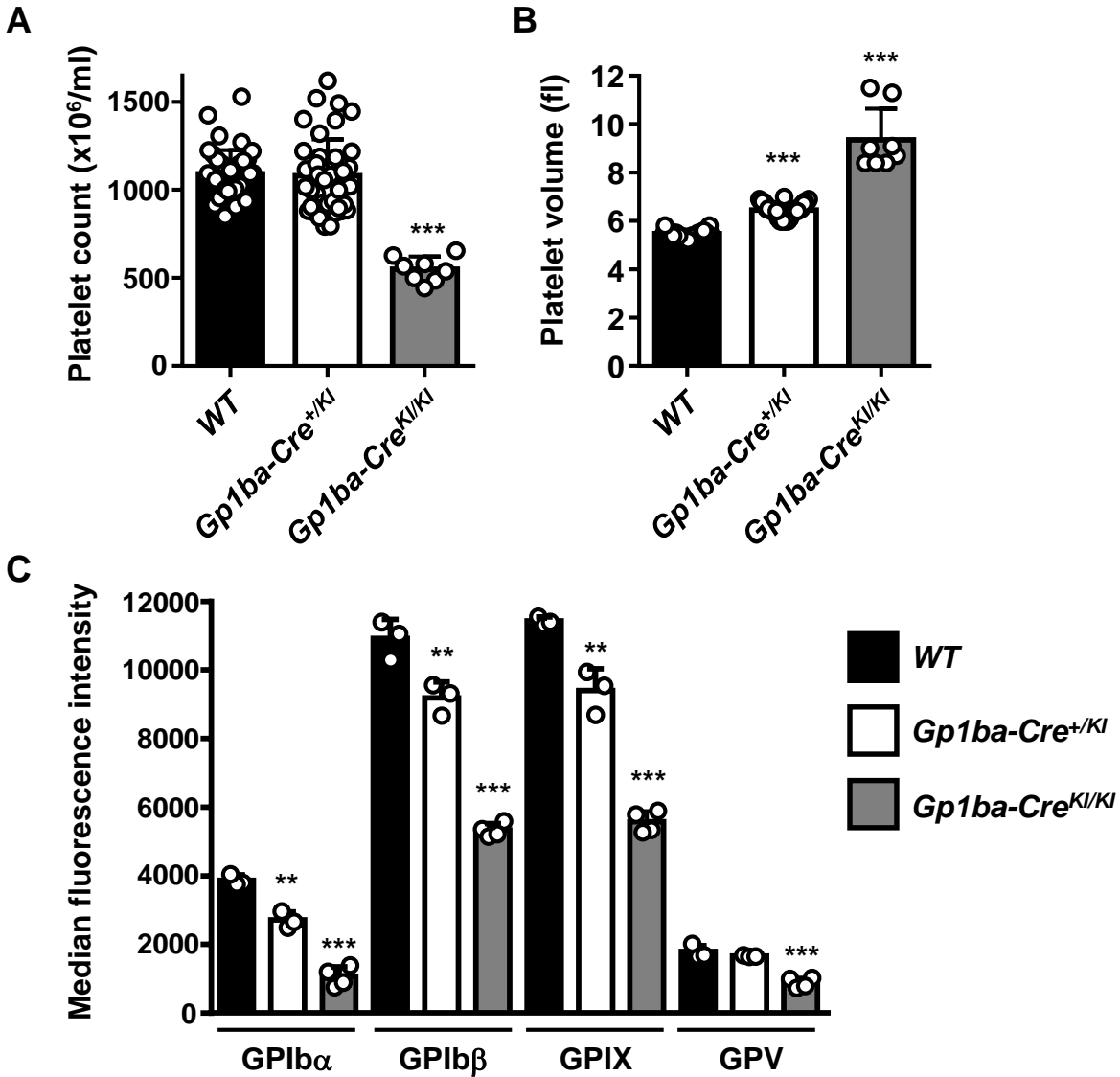
Data was collected from 7 breeding pairs (*Gp1ba-Cre*^{+K1} × *Gp1ba-Cre*^{+K1}).

Supplemental Table 3.
Hematology data of *Gp1ba-Cre* mice.

Hematological parameters	<i>Gp1ba-Cre</i> ^{+/+} (mean ± SD; n = 40)	<i>Gp1ba-Cre</i> ^{+/<i>K</i>} (mean ± SD; n = 42)
PLT (10 ⁶ /mL)	1090 ± 135	1079 ± 208
MPV (fl)	5.5 ± 0.1	6.4 ± 0.3***
RBC (10 ⁶ /μL)	11.4 ± 0.7	11.8 ± 1.0
HCT (%)	32.7 ± 2.8	33.4 ± 2.2
WBC (10 ³ /μL)	8.5 ± 3.1	7.5 ± 2.7
LYM (10 ³ /μL)	11.4 ± 4.2	9.9 ± 3.6
MON (10 ³ /μL)	0.6 ± 0.4	0.8 ± 0.6
NEU (10 ³ /μL)	1.6 ± 0.6	1.4 ± 0.4
EOS (10 ³ /μL)	0.04 ± 0.08	0.03 ± 0.09
BAS (10 ³ /μL)	0.09 ± 0.09	0.07 ± 0.07

PLT, platelets; MPV, mean platelet volume; RBC, red blood cells; HCT, haematocrit, WBC, white blood cells; LYM, lymphocytes; MON, monocytes; NEU, neutrophils; EOS, eosinophils; BAS, basophils. ***P < 0.001, unpaired, two-tailed t-test, mean ± SD.

Supplemental Figure 1



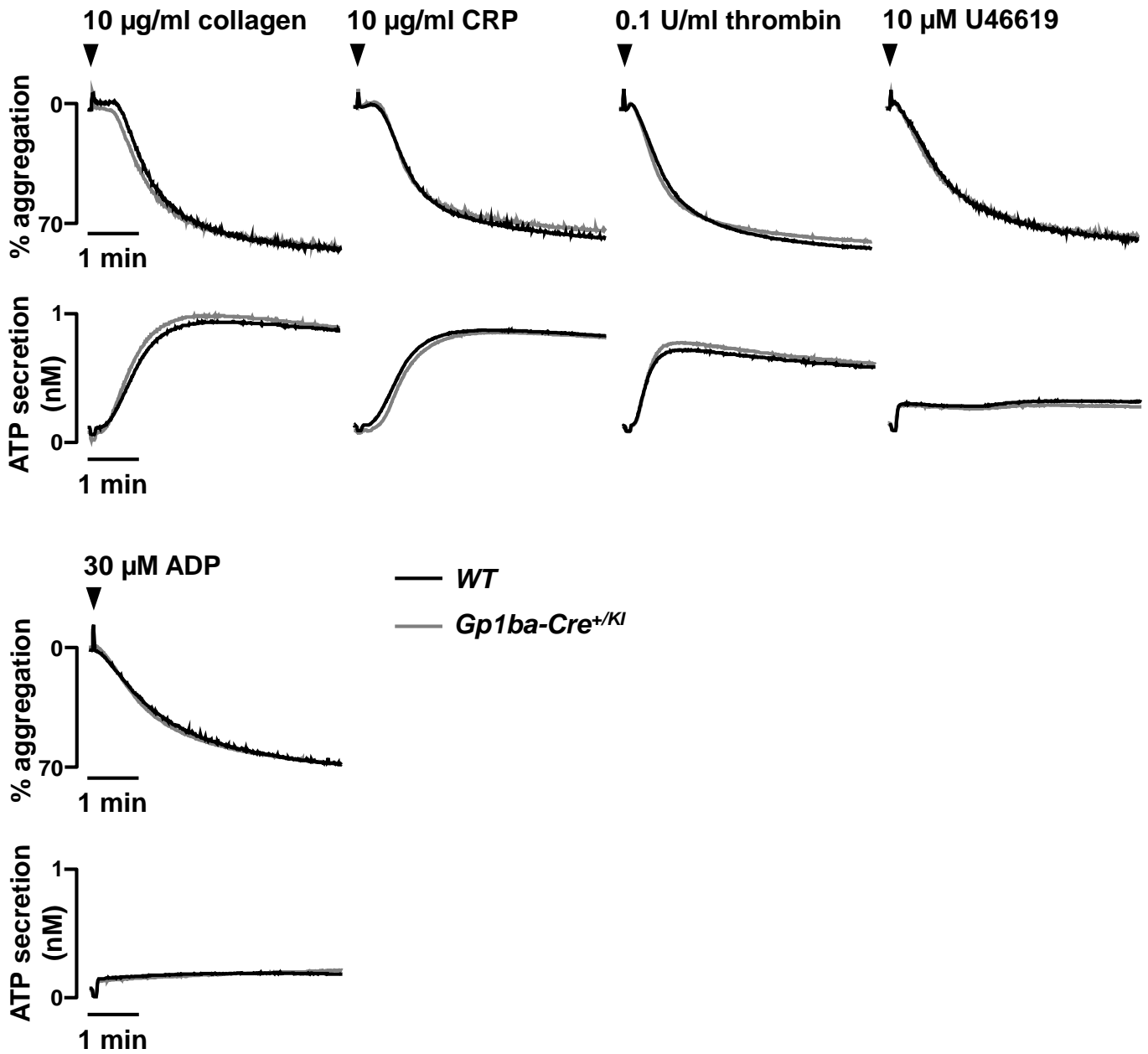
Supplemental Figure 1. Platelet parameters of the *Gp1ba-Cre^{KI/KI}* mouse.

(A) Platelet counts, n = 8-42 mice/genotype. (B) Platelet volumes, n = 8-42 mice/genotype. (C)

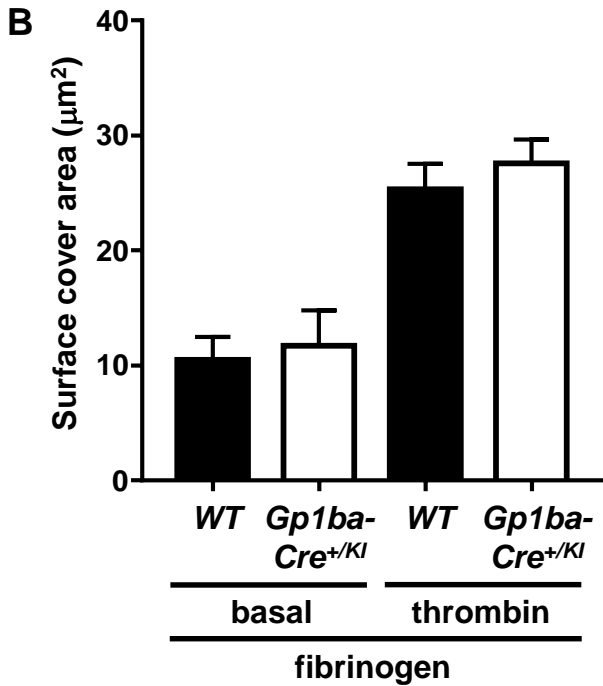
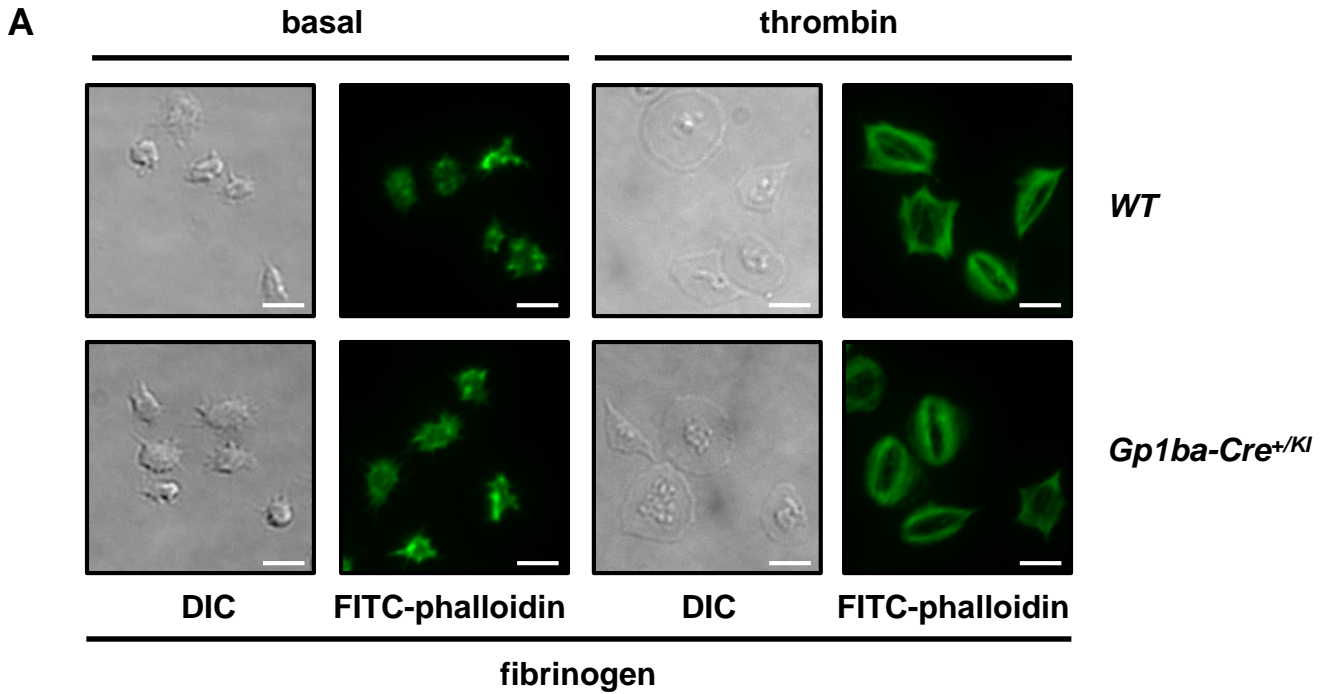
Platelet surface receptor expression of GPIb α , GPIb β , GPIX and GPV were measured by flow cytometry and shown as median fluorescence intensity, n = 3-4 mice/genotype.

Asterisks refer to significant difference compared with WT (** $P < 0.01$, *** $P < 0.001$, 1-way ANOVA with Tukey's test) mean \pm SD

Supplemental Figure 2

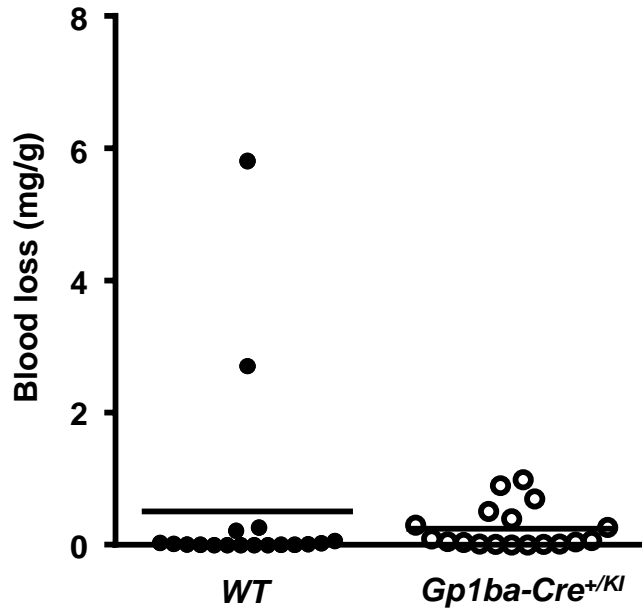


Supplemental Figure 2. Aggregation of *Gp1ba-Cre* platelets in response to higher agonist concentrations. Mean platelet aggregation and secretion traces in response to the indicated agonists, n = 4-8 mice/condition/genotype.



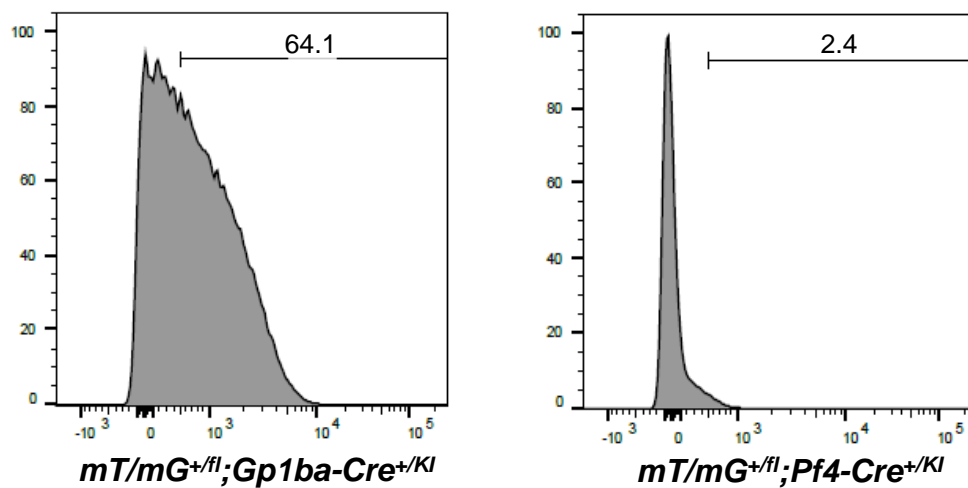
Supplemental Figure 3. Spreading of *Gp1ba-Cre* platelets on fibrinogen. (a) Representative differential interference contrast (DIC) microscopy images of resting (basal) and thrombin-stimulated (0.1 Units/ml) platelets spread on fibrinogen-coated cover-slips (100 µg/ml, 45 minutes, 37°C), scale bar: 5 µm. (b) Mean surface area of individual platelets quantified by ImageJ software, n = 5 mice/condition/genotype. Unpaired, two-tailed t-test, mean ± SD.

Supplemental Figure 4



Supplemental Figure 4. Tail bleeding of the *Gp1ba-Cre* mouse. Hemostatic response was measured in tail bleeding assays by an excision of a 5-mm portion of the tail tip followed by the determination of lost blood/body weight (normalized blood loss), $n = 18$ mice/genotype. Tail bleeding assays were conducted in a blinded manner. Unpaired, two-tailed t-test.

Supplemental Figure 5



Supplemental Figure 5. tdTomato+ fraction of eGFP+ platelets.

n=6