DOI: 10.1111/ith.15864

# ORIGINAL ARTICLE

# A gain-of-function filamin A mutation in mouse platelets induces thrombus instability

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#### **Funding information**

Fondation pour la Recherche Médicale, Grant/Award Number: LPC20170637458; INSERM

## Abstract

**Background:** Filaminopathies A are rare disorders affecting the brain, intestine, or skeleton, characterized by dominant X-linked filamin A (*FLNA*) gene mutations. Macrothrombocytopenia with functionally defective platelets is frequent. We have described a filaminopathy A male patient, exhibiting a C-terminal frame-shift FLNa mutation (Berrou et al., Arterioscler Thromb Vasc Biol. 2017;37:1087–1097). Contrasting with female patients, this male patient exhibited gain of platelet functions, including increased platelet aggregation, integrin  $\alpha$ Ilb $\beta$ 3 activation, and secretion at low agonist concentration, raising the issue of thrombosis risk.

**Objectives:** Our goal is to assess the thrombotic potential of the patient FLNa mutation in an *in vivo* model.

Methods: We have established a mutant FInA knock-in mouse model.

**Results:** The mutant FlnA mouse platelets phenocopied patient platelets, showing normal platelet count, lower expression level of mutant FlnA, and gain of platelet functions: increased platelet aggregation, secretion, and  $\alpha$ Ilb $\beta$ 3 activation, as well as increased spreading and clot retraction. Surprisingly, mutant FlnA mice exhibited a normal bleeding time, but with increased re-bleeding (77%) compared to wild type (WT) FlnA mice (27%), reflecting hemostatic plug instability. Again, in an *in vivo* thrombosis model, the occlusion time was not altered by the FlnA mutation, but arteriolar embolies were increased (7-fold more frequent in mutant FlnA mice versus WT mice), confirming thrombus instability.

**Conclusions:** This study shows that the FInA mutation found in the male patient induced gain of platelet functions *in vitro*, but thrombus instability *in vivo*. Implications for the role of FLNa in physiology of thrombus formation are discussed.

Manuscript handled by: Katsue Suzuki-Inoue

Final decision: Andreas Greinacher, 22 August 2022

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# 1 | INTRODUCTION

Filamin(s) (FLN), a family of three members (FLNa, FLNb, and FLNc), are essential components of the cell cytoskeleton by stabilizing the actin network. FLN(s) are large dimeric proteins composed of N-terminal actin-binding domains followed by 24 immunoglobulin (Ig)-like repeats interrupted by two hinge regions and a C-terminal region that ensures the dimerization.<sup>1,2</sup> FLNa and FLNb are ubiquitously expressed whereas FLNc is restricted to smooth and striated muscles.<sup>3,4</sup> FLNa, which is the most abundant isoform, is encoded by the FLNA gene located on chromosome Xq28, and is predominantly expressed in platelets. Today more than 90 FLNa partners have been described.<sup>5</sup> Most of them bind to Ig repeats 16–23 of FLNa. These interactions are essential for a large variety of cellular functions such as migration, adhesion, and spreading. Deletion of FLNa in mouse platelets, as shown with FlnA<sup>loxP</sup> GATA1-Cre mice, induces larger platelets, macrothrombocytopenia, and increased tail-bleeding.<sup>6</sup> In platelets, FLNa interacts with glycoprotein (GP) $lb\alpha$ through the Ig repeat 17 and ensures the mechanical stabilization of the plasma membrane under high shear.<sup>7</sup> The GPlb $\alpha$ -FLNa interaction is also essential for the trafficking of GPIb to the plasma membrane.<sup>8</sup> Another essential partner of FLNa in platelets is the integrin  $\alpha$ IIb $\beta$ 3. FLNa by its Ig repeat 21 binds to  $\beta$ 3 tail (amino acids 747–755) and regulates the activation of  $\alpha$ Ilb $\beta$ 3 integrin.<sup>9</sup> It has been proposed that constitutively bound FLNa to  $\beta$ 3 in resting platelets maintains the integrin in an inactive state, possibly by clasping  $\alpha$ IIb and  $\beta$ 3 tails together.<sup>10</sup> but definitive experimental evidence is still lacking. Likewise, it has been proposed that the activation of  $\alpha$ IIb $\beta$ 3 integrin requires the dissociation of FLNa from  $\beta$ 3 allowing talin binding, but hard evidence is missing as well. Recently, our group showed clearly that the FLNa- $\beta$ 3 interaction exerts a negative regulation on talin- $\beta$ 3 interaction after activation, using a megakaryocytic cell line (HEL).<sup>11</sup>

The tyrosine kinase Syk involved in the GPVI signaling is another partner of FLNa (Ig repeat 5) highlighted in *Flna<sup>-/-</sup>* mice.<sup>6</sup> This interaction, essential in platelet activation under arterial shear, allows Syk recruitment to GPVI.

Recently, new partners have been described in platelets. The adaptor protein PACSIN2, which interacts with FLNa Ig repeat 20, regulates membrane tubulation in megakaryocytes and platelets.<sup>12</sup> STIM1, which regulates store-operated calcium entry (SOCE) and Ca<sup>2+</sup> influx, interacts with FLNa.<sup>13</sup> Finally, the Rho GTPase, RhoA, which binds constitutively to FLNa via the C-terminal Ig repeat 24, has recently been reported to be dependent on  $\alpha$ IIb $\beta$ 3. This interaction modulates proplatelet formation in megakaryocytes<sup>14</sup> and outside-in signaling in platelets.<sup>11</sup> How the interactions of FLNa with its partners integrate cellular signaling and cellular functions remains an open question. Investigating *FLNA* gene variants is of great help. Today, *FLNA* mutations are associated with a wide spectrum of 10 distinct genetic syndromes called filaminopathies A affecting numerous organs including the brain, skeleton, heart, and skin.<sup>15</sup> Periventricular

### Essentials

- Filaminopathies A are multi-organ and platelet ailments due to X-linked filamin A gene mutations.
- We assessed the thrombosis ability of a patient gain-offunction filamin A (FLNa) mutation in a mouse model.
- Mouse platelets showed gain of platelet functions, but paradoxical thrombus instability *in vivo*.
- Thrombus instability suggests FLNa is involved in the strength of platelet-platelet interaction.

nodular heterotopia (PVNH), the most frequent syndrome, is characterized by a defect of neuron migration.<sup>16,17</sup> Most of PVNH, observed in heterozygous females exhibiting loss-of-function mutations, ablate FLNa expression.<sup>17-19</sup> In males, filaminopathy A is generally associated with perinatal lethality; only rare males survivie.<sup>20,21</sup> The PVNH phenotype can be associated with Ehlers-Danlos syndrome, with skeletal dysplasia, congenital intestinal pseudo-obstruction, and familial cardiac valvular dystrophy.<sup>22-24</sup> PVNH is often associated with macrothrombocytopenia and bleeding tendency.<sup>17</sup> We have previously shown in female patients with PVNH that have low platelet count characterized by giant and normal platelets was associated with a reduced FLNa level whereas normal platelet count was associated with a normal FLNa content.<sup>14,19</sup> In PVNH, depending on the level of FLNa, platelet functions were often impaired, causing bleeding tendency.<sup>19,25</sup> Moreover, we have shown that the absence of FLNa protein in megakaryocytes leads to their inability to produce platelets.<sup>14</sup> More recently, we have studied the rare case of a male filaminopathy A patient with a novel mutant FLNa with a 100 amino acid-long C-terminal extension.<sup>20</sup> Surprisingly, platelet functions were significantly enhanced, contrary to functional inhibition in heterozygous female patients. This was the first observation in humans of the role of FLNa in the downregulation of  $\alpha$ IIb $\beta$ 3 activation, a key feature in platelet physiology and pathology. However, a number of questions remain unanswered. The aim of this study was to establish an "in vivo" mutated FLNa knock-in mouse model and to address the question of the potential role of this FLNa mutation in thrombosis.

### 2 | MATERIAL AND METHODS

### 2.1 | Reagents

Fibrillar collagen (equine type I) and adenosine diphosphate (ADP) were obtained from Kordia. Monoclonal antibody directed against GPVI was generously provided by Dr. M. Jandrot-Perrus (Paris, France). Apyrase grade VII, bovine thrombin, rhodamine 6G, and prostaglandin

E1 were from Sigma-Aldrich. Fibrinogen was obtained from HYPHEN BioMed SAS. Convulxin was purchased from Cryopep. D-Phe-Pro-Arg chlormethylketone dihydrochloride (PPACK) was from Calbiochem-VWR. Adenosine triphosphate (ATP) determination kit, Alexa Fluor 488-labeled phalloidin, and Alexa Fluor secondary antibodies were from Molecular Probes. The rat monoclonal antibody against the activated form of mouse αIIbβ3 (JON/A-Phycoerythrin PE), fluorescein isothiocyanate (FITC)-labeled rat anti-mouse GPVI monoclonal antibody (mAb; JAQ1), FITC-labeled rat anti-mouse CD62P (P-selectin) mAb (Wug.E9), and FITC-labeled rat anti-mouse GPIb amAb (Xia.B2) were from emfret ANALYTICS. FITC-labeled rat anti-mouse integrin allb chain (CD41) mAb (MWReg30) was purchased from BD Biosciences. A rabbit polyclonal antibody was raised against the extended C-terminal FLNa sequence of a patient by GenScript, from a mix of two synthetic FLNa peptides (APPSSPQPRPRPPWC and DGGPVHNHPLVLFSSQEEC) coupled to the carrier protein keyhole limpet hemocyanin, followed by affinity purification of the IgG fraction. The mAb specific for N-terminal FLNa and the polyclonal antibody specific for phosphorylated linker for activation of T cells (LAT) were obtained from Millipore. The polyclonal antibodies against phosphorylated Syk and protein kinases C (PKCs) substrates were from Cell Signaling Technology. Peroxidase-conjugated AffiniPure secondary antibodies were obtained from Jackson ImmunoResearch Laboratories, Inc.

### 2.2 | Mouse strains

The mutant FlnA knock-in mouse strain was generated by genOway. A construct was built (Figure 1A) by polymerase chain reaction (PCR), consisting of mouse FInA exon 45 added to a polyadenylation site. and flanked with 3' and 5' loxP sites, followed by a neomycin cassette and a second mouse exon 45 where the STOP codon and the mouse 300 bp long 3' downstream untranslated sequence were swapped for the mutated STOP (ct deletion, see Figure S1A in supporting information) followed by the human 300 bp 3' untranslated sequence of the patient human FLNA gene. The human 3' sequence was used instead of the mouse 3' sequence because of the very poor homology between the two sequences, in particular at the peptide sequence level (see Figure S1). The construct was intended to be recombined after Cre expression with deletion of the wild-type (WT) exon 45, leaving only the mutant human sequence-extended exon 45 (Figure 1A). This cassette was recombined at the site of the WT exon 45 of the chromosome X-linked mouse FlnA gene by CRISPR-Cas9 technology on mouse embryonic stem cells. After injection into blastocysts, chimeric animals were selected for germ-line transmission. Absence of leakage of recombinant FInA was checked by real-time PCR. FIna<sup>loxP</sup> females were then selected and crossed with PF4-Cre male mice, to generate mice expressing platelet FInA with the extended human C-terminal sequence. Only hemizygous males were studied further.

All experimental procedures were carried out in accordance with the European legislation concerning the use of laboratory animals and approved by the ethical committee CEEA26 under the number Apafis #13732–2018092708484303.

### 2.3 | Isolation of mouse platelets

Mice were anesthetized by intraperitoneal injection of xylazin (10 mg/kg) and ketamine (100 mg/kg), then whole blood was collected by cardiac puncture into  $80\mu$ M PPACK and 10% ACD/C buffer (124 mM sodium citrate, 130 mM citric acid, 110 mM dextrose, pH 6.5). Isolated platelets were obtained as previously described.<sup>26</sup>

## 2.4 | Tail bleeding time

Bleeding time of anesthetized mice (10 weeks old) was determined by cutting off the tip of the tail (3 mm of distal mouse tail) and immediate immersion of the tail into 37°C isotonic saline (0.9% NaCl). Bleeding time was defined as the first cessation of bleeding for at least 60s, to ensure that bleeding did not start again, otherwise it was considered rebleeding.

### 2.5 | Hematologic analysis

Blood counts were determined with an automatic cell counter (Scil Vet abc Plus, Horiba Medical).

### 2.6 | Flow cytometry

Washed platelets (3  $\times$  10<sup>8</sup>/ml) were stimulated with a range of activators (convulxin, ADP, or thrombin). After incubation for 10 min without stirring, platelets were diluted at 10<sup>8</sup>/ml and incubated with the appropriate fluorophore-conjugated antibodies for 20min at room temperature. The samples were then directly analyzed with an Accuri C6<sup>+</sup> flow cytometer (BD Biosciences).

### 2.7 | Platelet aggregation

Platelet aggregation was monitored by measuring light transmission through the stirred suspension of washed platelets ( $3 \times 10^8$ /ml) at  $37^{\circ}$ C using a Chronolog aggregometer. Platelet aggregation was triggered by adding fibrillar collagen, convulxin, ADP, or thrombin. Representative traces for aggregation were obtained from at least three independent experiments.

# 2.8 | Measurement of platelet dense granule secretion

Dense granule secretion was evaluated by measuring ATP release with the ATP determination kit at the end of aggregation as previously described.<sup>26</sup> Briefly, platelet aggregation was stopped by adding ice-cold ethylenediaminetetraacetic acid (EDTA; 16 mM) before centrifugation (12000 g; 1 min). Supernatants were then



FIGURE 1 Schematic representation of the mutant humanized FInA cloning strategy and FInA expression in platelets of male mice. AThe region of the mouse *Flna* gene targeted for homologous recombination is shown. *Flna WT* includes exons from #38 to 45 (hashed boxes). Exon 45 shows the relative position of the stop codon and the non-coding 3' sequence of the exon is drawn as a white box. *Flna<sup>loxP</sup>* shows the *FLNA* gene after recombination of the exon38--45 region with a construct including mouse exon 45 with an upstream lox P site (blue arrowhead), and followed by the neomycin ("neo") negative selection gene and a second exon 45 harboring the downstream loxP site (blue arrowhead), the human stop mutation (red .) followed by the 3' 300 bp human sequence (black box), which includes an in-frame second stop codon, shown on the drawing. *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> shows the same exon 38-45 region after expression of the Cre recombinase in the platelet–megakaryocyte lineage: deletion of the loxP–loxP fragment (shown by the red lines and the "Cre recombination" mention), leads to swapping of the wild-type exon 45 for the mutated one. B, *PF4-Cre*<sup>+</sup> males were crossed with *Flna<sup>loxP</sup>* females. *PF4-Cre* genomic insertion in *Flna<sup>loxP</sup>* male offsprings was investigated by genomic polymerase chain reaction and agarose gel electrophoresis. C, Percentage of male over female mice during breeding. D, In a population of 200 mice, distribution of males and females and of *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> versus *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> and *Flna<sup>loxP</sup>PF4-Cre*<sup>-</sup> males as assessed by western blotting. The graph represents the means  $\pm$  standard error of the mean of four independent *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> male mice and four independent *Flna<sup>loxP</sup>PF4-Cre*<sup>-</sup> male mice, \*\**p* <.001 (unpaired Student t-test)

incubated with recombinant firefly luciferase (0.25 mg/ml) and its substrate, D-luciferin (0.5 mM), according to the manufacturer's instructions. Light emission was assessed with a luminometer

(Fluoroskan Ascent FL; Thermo LabSystems). Dense granule secretion was expressed as the amount of ATP released (in picomoles) from  $3 \times 10^8$  platelets.

### 2.9 | Western blotting

Washed platelets (300 µl;  $3 \times 10^8$ /ml) were activated for 5 min at 37°C by various concentrations of Cvx (0 to 400 pM) in unstirred conditions and in the presence of apyrase (5 U/ml) and indomethacin (5 µM). Then, platelets were lysed in denaturing buffer (50mM Tris, 100mM NaCl, 50mM NaF, 5 mM EDTA, 40mM β-glycerophosphate, 100µM phenylarsine oxide, 1% sodium dodecyl sulfate [SDS], 5 µg/ml leupeptin, 10 µg/ml aprotinin, pH 7.4). Proteins were separated by SDS-polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes. Membranes were incubated with primary antibodies followed by horseradish-peroxidase-labeled secondary antibodies. Immunoreactive bands were visualized with enhanced chemiluminescence detection reagents (Perbio Science) using a G:BOX Chemi XT16 Image System and quantified using Gene Tools version 4.03.05.0 (Syngene).

### 2.10 | Spreading

Platelet spreading was performed on fibrinogen matrix ( $100 \mu g/ml$ ). Platelets were incubated at 37°C for 30min, fixed with 4% paraformaldehyde, permeabilized, and stained by incubation for 30min with Alexa Fluor<sup>488</sup>–labeled phalloidin (0.3  $\mu$ M). Then platelets were visualized under an epifluorescence microscope (Nikon, Eclipse 600). Cell surfaces were analyzed using Fiji software (rsb.info.nih.gov/ij).

### 2.11 | Clot retraction

Clot retraction studies were performed at 37°C in aggregometer tubes. First, a 6% (w/v) polyacrylamide cushion was polymerized at the bottom of the tube to avoid clot adherence to the glass bottom. Then, washed platelets (300 $\mu$ l at 3 $\times$ 10<sup>8</sup> platelets/ml) was added to fibrinogen 500 $\mu$ g/ml, thrombin 2 U/ml, and Ca<sup>2+</sup> 2 mM in the tubes. Clot retraction was monitored for 1 h.

# 2.12 | *In vitro* thrombus formation under flow conditions

Thrombus formation was evaluated in a whole blood perfusion assay on a fibrillar collagen matrix  $(50\,\mu g/ml)$  under arterial shear conditions (shear rate of  $1500\,s^{-1}$ ). Thrombus formation was quantified by assessment of the mean fluorescence intensity (MFI) of adherent thrombi.

### 2.13 | Ferric chloride-induced thrombosis model

Ferric chloride (FeCl<sub>3</sub>) injury was induced in 4- to 5-week-old mice, as previously described.<sup>26</sup> To facilitate visualization of thrombus

formation, platelets from anesthetized mice were fluorescently labeled *in vivo* by injection of rhodamine 6G (3.3 mg/kg) into the retro-orbital plexus. After topical deposition on the mesenteric vessels of an FeCl<sub>3</sub> solution (10%), thrombus growth was monitored in real time with an inverted epifluorescent microscope (×10; Nikon Eclipse TE2000-U). During the observation period, the number of full detachments of thrombi from the vessel wall at the injury site was recorded and counted as embolization events.

### 2.14 | Statistical analysis

Statistical significance was evaluated with unpaired Student's *t*tests or one-way analysis of variance followed by a post hoc test, as indicated in the figure legends. All analyses were performed with GraphPad Prism software (version 6, GraphPad Inc.).

### 3 | RESULTS

# 3.1 | Decreased expression level of mutated FlnA in platelets

Because the patient we reported previously is a male, thus hemizygous for the mutation of the X-linked FLNA gene, the effect of the FInA mutant was investigated on male mice expressing PF4-Cre or not (Figure 1B). Only mutated FInA was detected in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets whereas only WT FlnA was detected in Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets. Note that the population of male mice was significantly lower than that of female mice (male mice: 37.4% + 5.4%vs. female mice:  $63.7\% \pm 5.4\%$ ; p = .003; Figure 1C) and that this difference was the result of a lower number of Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> male mice (30.6%  $\pm$  6.6%; p = 000.1) compared to Flna<sup>loxP</sup> PF4-Cre<sup>-</sup> male mice (Figure 1D). During this study, we never noticed dead bodies of newborn mice suggesting strongly embryonic lethality. Western blotting using an antibody specific for the extended C-terminal tail of the patient FLNa showed first that mutated FlnA was only detected in platelets of Flna<sup>loxP</sup>PF4Cre<sup>+</sup> mice (Figure 1E). In contrast, western blotting using an anti-whole FLNa antibody showed that both Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> and Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> mice expressed FInA in platelets but at a lower level in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets  $(49.5\% \pm 3.9\%)$  compared to Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets (100%; Figure 1E, F). In contrast, the expression level of FlnB was similar in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets compared to Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets (Figure S2 in supporting information).

Moreover, analysis of the hematologic parameters showed a normal platelet count ( $Flna^{loxP}PF4$ - $Cre^-$ :  $844\pm32\times10^9$ /L vs.  $Flna^{loxP}PF4$ - $Cre^+$ :  $880\pm44\times10^9$ /L; n = 6) and normal platelet volume ( $Flna^{loxP}PF4$ - $Cre^-$ :  $5.55\pm0.07$  fL vs.  $Flna^{loxP}PF4$ - $Cre^+$ :  $5.50\pm0.07$  fL; n = 6; Table 1). Moreover, no difference was observed in the expression level of the major receptors ( $\alpha$ IIb $\beta$ 3, GPVI, and GPIb $\alpha$ ) analyzed by flow cytometry and of granule content (von Willebrand factor) measured by western blotting (Figure S3 in supporting information).

### TABLE 1 Hematological parameters

	FIna <sup>loxP</sup> PF4-Cre <sup>−</sup>	FIna <sup>loxP</sup> PF4- Cre <sup>+</sup>
Platelets (×10 <sup>9</sup> /L)	844 ±32	$880 \pm 44$
WBC (×10 <sup>9</sup> /L)	$8.1 \pm 1.0$	9.4 ± 0.4
RBC (×10 <sup>12</sup> /L)	7.7 ±0.8	9.4 ± 0.6
Hematocrit (%)	49.1 ± 1.5	47.2 ± 1.1
MPV (fL)	$5.55 \pm 0.07$	$5.50 \pm 0.07$

Abbreviations: MPV, mean platelet volume; RBC, red blood cells; WBC, white blood cells.

Altogether, these results show that this mouse platelet model with an extended human sequence at the FInA C-terminal tail specifically expressed in platelets reproduces the results obtained with the patient's platelets<sup>20</sup> with a normal platelet count and a lower level of mutated FLNa (patient: 30%, mice: 50%).

# 3.2 | Unstable hemostasis in mutated FInA mice

We next investigated the role of mutated FInA in hemostasis by comparing tail bleeding times in *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> and *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice *in vivo*. Similar bleeding times were observed in *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> and *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice (*Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup>: 51.9  $\pm$ 6.3 s vs. *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup>: 54.4  $\pm$ 6.7 s; Figure 2A). In contrast, rebleeding was observed in a large percentage of *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice (77%) compared to *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> mice (27%; Figure 2B) consistent with an instability of the thrombus in *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice.

# 3.3 | Mutated FInA increases platelet aggregation induced by thrombin and GPVI agonists

The next step was to investigate the impact of the FInA mutation on platelet function in vitro. First, platelet aggregation and dense granule secretion measured by the quantification of the ATP release was higher in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets compared to Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets in conditions of low concentration of thrombin (40 mU/ml; aggregation: 77.1% ± 2.6% for Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets vs.  $63.1\% \pm 3.3\%$  for Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets, p = .007; secretion: 20.1  $\pm$  0.5 pmoles for Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets vs. 14.4  $\pm$  0.2 pmoles for Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets, p < .001 n = 5; Figure 3A-C). In contrast, in conditions of unstirred platelets after thrombin activation, no difference between *Flna<sup>loxP</sup>PF4-Cre*<sup>-</sup> platelets and *Flna*<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets was observed for  $\alpha$ IIb $\beta$ 3 integrin activation and  $\alpha$  granule secretion, measured by flow cytometry with a rat mAb JON/A(PE) specific for the activated conformation of the mouse integrin and a FITC-labeled rat anti-mouse CD62P mAb for Pselectin expression (Figure 3D,E) suggesting that outside-in signaling and/or shear were essential for the enhanced effect observed in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets after thrombin activation. Finally, aggregation and  $\alpha$ Ilb $\beta$ 3 activation induced by another agonist of G proteincoupled receptor such as ADP (5 to  $20 \mu M$ ) were normal (Figure S4 in



**FIGURE 2** Mouse tail bleeding time. A, Tail bleeding was assessed in *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> (mutated FlnA) and *Flna<sup>loxP</sup>PF4-Cre*<sup>-</sup> mice (wild-type FlnA). B, Quantification of rebleedings is expressed as mean  $\pm$  standard error of the mean of 11 *Flna<sup>loxP</sup>PF4-Cre*<sup>-</sup> male mice and 13 *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> male mice

supporting information). These results show that mutated FInA may affect platelet activation induced by G-coupled receptors.

Then the role of mutated FInA was investigated in platelet activation induced by GPVI, the collagen receptor. First, in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets, aggregation induced by different concentrations of agonists (Cvx: 100 to 400 pM; Figure 4A,B) and collagen (2 to 4 µg/ml; Figure 4F,G) was significantly increased but only with agonists at low concentration (Cvx: 100 pM; p = .02 and collagen 2 and 3  $\mu$ g/ml; p = .002p = .006, respectively). In parallel, ATP secretion from dense granules in conditions of platelet aggregation was also significantly increased with Cvx (100 pM) reaching 18.8 pmoles for Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets versus 3.0 pmoles for Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets (p < .0001) and collagen (2  $\mu$ g/ml: Flna<sup>loxP</sup>PF4-Cre<sup>+</sup>: 10.4 ±0.2 pmoles vs. Flna<sup>loxP</sup>PF4-Cre<sup>-</sup>: 9.2 ±0.3 pmoles, p = .02; 3 µg/ml: Flna<sup>loxP</sup>PF4-Cre<sup>+</sup>: 11.5 ±0.3 pmoles vs. Flna<sup>loxP</sup>PF4-Cre<sup>-</sup>: 9.5  $\pm$ 0.3 pmoles, p = .002; Figure 4C-H). Note the plateau of ATP secretion was obtained at 200 pM Cvx for both Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> and Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets (Figure 4C). Furthermore, the activation of αllbβ3 measured with mAb JON/A in flow cytometry showed a significant increase of JON/A binding (p = .02) in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets at Cvx 400 pM (Flna<sup>loxP</sup>PF4-Cre<sup>+</sup>: 2157.5 ±49.8 MFI vs. Flna<sup>loxP</sup>PF4-Cre<sup>-</sup>: 1519.1 ±199.7 MFI; Figure 4D). Finally, the quantification of P-selectin exposure induced by various Cvx concentrations (0 to 400 pM) in unstirred conditions confirmed the gain of function of mutated FInA (Figure 4E). Note that no activation of  $\alpha$ IIb $\beta$ 3 was shown when platelets were activated by collagen in contrast to Cvx. The unstirring conditions of flow cytometry are not compatible with fibrillar collagen activation.

We then assessed GPVI signaling induced by various concentrations of Cvx (0 to 400 pM) in unstirred conditions and in the presence of apyrase and indomethacin. The phosphorylation of two proteins (Syk and LAT) essential in GPVI signaling was tested by western blotting. Indeed, the phosphorylation of Syk was shown to be dependent on its interaction with FInA.<sup>6</sup> In *FIna<sup>loxP</sup>PF4-Cre*<sup>+</sup> platelets, Syk phosphorylation was significantly decreased after activation by 200



FIGURE 3 Aggregation, secretion, and allbβ3 activation induced by thrombin. A, B, Aggregation of washed Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> (Ctrl) and Flng<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets was induced by thrombin (30, 40, and 80 mU/ml). C, Dense granule secretion measured after 3 min of platelet aggregation was quantified by measuring adenosine triphosphate release. D, E, Quantification of P-selectin exposure and  $\alpha$ IIb $\beta$ 3 activation in conditions of unstirred platelets was assessed by flow cytometry using an anti-P-selectin antibody and the JON/A monoclonal antibody specific for the activated form of  $\alpha$ IIb $\beta$ 3. The graph represents the means  $\pm$  standard error of the mean of three independent determinations. \*p < .05, \*\*\*p < .001 (one-way analysis of variance with Sidak's post hoc test for multiple comparisons)

pM (p = .007) and 400 pM Cvx (p < .0001; Figure 5A), as well as LAT phosphorylation after activation by 400 pM Cvx (p < .001; Figure 5B). These decreases in Syk and LAT phosphorylation reproduce the results obtained with the patient's platelets, and are probably the conseguence of the low platelet level of FInA (50% and 30%, respectively).

Finally, we decided to test the activation of PKCs, previously shown to be involved in secretion,<sup>27,28</sup> by measuring phosphorylated PKC substrates induced by Cvx (0 to 400 pM) in unstirred conditions (Figure 5C). The intensity of phosphorylated bands was significantly increased by 37% (p = .004) at 400 pM Cvx in Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets compared to Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> platelets indicating that mutated FInA increased the PKC activity. The results indicate that mutated FInA does not affect proximal GPVI signaling but increases PKC activities required for platelet secretion.

#### 3.4 **Mutated FInA induces increased** outside-in signaling

We next addressed the guestion of the impact of mutated FInA on outside-in signaling. First, platelet spreading on fibrinogen matrix was examined. Platelet area of Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> mice was slightly but significantly increased ( $127\% \pm 12\%$ ; p = .03) compared to platelet area of *Flna<sup>loxP</sup>PF4-Cre<sup>-</sup>* mice (100%; Figure 6A). Then, we examined the role of mutated FInA on clot retraction (Figure 6B-C). Clot retraction of Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelets was significantly accelerated comparatively to  $Flna^{loxP}PF4$ -Cre<sup>-</sup> platelets, reaching 40.3 ±4.1% (p = .04) versus 25.7%  $\pm$  3.9% for  $Flna^{loxP}$  PF4-Cre<sup>-</sup> platelets after 10 min, consistent with mutated FInA upregulating clot retraction and outside-in signaling in the absence of shear.

FIGURE 4 Aggregation, secretion, and allbβ3 activation induced by convulxin or collagen. A, B, F, G, Platelet aggregations of washed platelets were initiated by various doses of Cvx (100 to 400 pM) and collagen (2 to 4 µg/ml). C, H, Dense granule secretion measured after 3 min of platelet aggregation was quantified by assessing adenosine triphosphate release. D, E, Quantification of P-selectin exposure and allbβ3 activation in conditions of unstirred platelets was assessed by flow cytometry using an anti-P-selectin antibody and the JON/A monoclonal antibody specific for the activated form of allbb3. The graph represents the means ± standard error of the mean of four independent determinations. \*p <.05, \*\*p <.01, \*\*\*p <.001 (one-way analysis of variance with Sidak's post hoc test for multiple comparisons)



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# 3.5 | Mutated FInA induces thrombus instability *in vivo*

We next investigated the effect of mutated FInA by assessing thrombus formation *in vitro* and *in vivo*. *In vitro*, thrombus formation was assessed on a collagen matrix at arterial shear rate  $(1500s^{-1})$ . Thrombus size was similar in *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice and in *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> mice (Figure 7A). Then *in vivo* thrombosis induced by ferric chloride on exposed mesenteric vessels was assessed. The initial thrombus was formed much more rapidly in venules of



FIGURE 5 Cell signaling induced by convulxin. Western blotting of phosphorylated Syk (A), LAT (B), and PKC substrates (C) of platelets activated for 5 min by various doses of Cvx (100 to 400 pM) in the presence of apyrase (5 U/ml) and indomethacin (5  $\mu$ M). Phosphorylation levels were normalized in relation to the expression level of total Syk (A), total LAT (B) of 14–3-3 $\zeta$  (C), and then compared with each other to the phosphorylation of *Flna<sup>loxP</sup>PF4-Cre<sup>-</sup>* (Ctrl) at 400 pM Cvx which was set to 100%. The graphs represent the means of phosphorylation ± standard error of the mean of at least three independent determinations. \*\*p <.01, \*\*\*p <.001 (one-way analysis of variance with Sidak's post hoc test for multiple comparisons)



FIGURE 6 Outside-in signaling of washed platelets. A, Platelet spreading was performed on a fibrinogen matrix (100 µg/ml). Platelets were incubated at 37°C for 30min and then stained with Alexa Fluor488-labeled phalloidin (0.3 µM). Platelets were visualized under an epifluorescence microscope (Nikon, Eclipse 600). Cell surfaces were analyzed using the Fiji software. The graph represents the means  $\pm$  standard error of the mean (SEM) of four independent determinations. \*p <.05, (unpaired Student *t*-test). B, Clot retraction of washed platelets was initiated by adding fibrinogen (500 µg/ml) and thrombin (0.5 U/ml). Photographs were taken at different times and (C) the area of clot was measured using the Fiji software. The graph represents the means  $\pm$  SEM of four independent determinations. \*p <.05, \*\*p <.01 (unpaired Student *t*-test)

*Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice (5.4 ± 1.8 min; *p* = .0013) than in venules of *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> mice (14.6 ± 1.2 min; Figure 7B). In contrast, no difference was found in arterioles. Surprisingly, if the occlusion times were similar between *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> and *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> mice in arterioles and in venules (Figure 7C), thrombus instability was observed in arterioles of all *Flna*<sup>loxP</sup>*PF4-Cre*<sup>+</sup> mice (100%) compared to only 14% of *Flna*<sup>loxP</sup>*PF4-Cre*<sup>-</sup> mice (Figure 7D). These results indicate that mutated FlnA affects *in vivo* thrombus stability but not thrombus size and occlusion.

# 4 | DISCUSSION

In this study, we have shown that the expression in platelets of FInA mutated in the C-terminal tail leads *in vitro* to a gain-of-function counteracted by thrombus instability *in vivo*. The gain-of-functions was translated by an increased platelet aggregation induced by low

concentrations of agonists, increased spreading, and clot retraction. In contrast, if the time of the first thrombus formation *in vivo* was shorter in *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> mice, unexpectedly, thrombosis *in vivo* was characterized by emboli formation in arterioles.

We found that the expression level of mutant FlnA (49% of total FlnA) in mice reproduces the expression level of mutant FLNa (30%) previously described in a male patient with the same mutation and exhibiting a PVNH and a congenital intestinal pseudo-obstruction.<sup>20</sup> Moreover, normal platelet count and normal platelet volume, both described in the patient<sup>20</sup> and in mice, indicate that this mouse platelet model with the extended sequence in the C-terminal tail present in platelets reproduces the results obtained with the male patient's platelets.<sup>20</sup>

Platelet aggregation and  $\alpha$ IIb $\beta$ 3 activation induced by low concentrations of thrombin, collagen, and Cvx were upregulated in *Flna<sup>loxP</sup>PF4-Cre*<sup>+</sup> platelets. In the patient, a gain of platelet aggregation was also observed with low concentrations of Cvx (aggregation and  $\alpha$ IIb $\beta$ 3 activation induced by thrombin and collagen as



FIGURE 7 In vivo thrombus instability of Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> mice. The impact of the FInA mutation on thrombus formation was investigated in vitro. A, A wholeblood perfusion assay was carried out on collagen matrix (50µg/ml) at an arterial shear rate (1500 s<sup>-1</sup>). Thrombus formation was evaluated by fluorescence microscopy and the thrombus size was quantified by measurement of the mean fluorescence intensity (MFI) of thrombi. B-D. In vivo thrombosis was assessed in exposed mesenteric vessels (venules and arterioles) after FeCl<sub>2</sub>-induced injury. Thrombus formation of fluorescently labeled platelets was monitored by intravital videomicroscopy. The graphs represent (B) the time point of the formation of the first thrombus of 30 µm, (C) the occlusion time defined as the stopping blood flow for at least 1 min, and (D) the number of emboli shedding from thrombi in venules throughout the procedure. The graph represents the means + standard error of the mean of eight FIna<sup>loxP</sup>PF4-Cre<sup>+</sup> and seven Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> mice. Statistical difference between Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> mice and Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> mice was evaluated by the unpaired Student *t*-test (\*\*p < .01)

well as  $\alpha$ IIb $\beta$ 3 activation induced by Cvx were not performed in the patient).<sup>20</sup> Again, and as found in the patient, the first steps of GPVI signaling (Syk and LAT phosphorylations) were not at the origin of increased platelet GPVI signaling. The phosphorylation of Syk and LAT was decreased and probably the result of a lower level of FInA (49%) and consequently lower interaction of mutated FInA with Syk required for GPVI signaling.<sup>6</sup> In contrast, in the advanced steps of GPVI signaling, PKC activities measured by the phosphorylation of PKC substrates were upregulated by mutated FlnA. This upregulation could explain the increased Flna<sup>loxP</sup>PF4-Cre<sup>+</sup> platelet secretion. A possible mechanistic explanation could be that, because the interaction of FLNa with PKCs and notably with PKC $\alpha$ , may be direct,<sup>5,29</sup> and because PKCs are mostly cytosolic, the FLNa/PKCα interaction increases PKC translocation to the plasma membrane, upon activation. Further studies will be required to explore this hypothesis.

Surprisingly, in contrast to the patient, platelet aggregation and  $\alpha$ Ilb $\beta$ 3 activation induced by ADP were not upregulated in mice. If the platelet cascade is genetically well conserved between humans and mice, some differences have been reported.<sup>30</sup> For example, purinergic receptor signaling regulated by the ADP receptor  $P2Y_{12}$  is clearly higher in mice compared to humans.<sup>30,31</sup> This difference could be the cause of a difference observed in ADP-induced platelet signaling between FLNa mutated human versus mouse platelets.

Whereas FInA full ablation in mice resulted in a diminution in the level of GPIb-IX-V complex,<sup>6</sup> a level of 49% of mutated FInA, as seen in our model, did not affect the expression level of GPIb $\alpha$ . This is most likely due to the partial diminution of FInA, but we cannot rule out a possible gain-of-function effect of the mutation on GPIb-IX-V trafficking.

Finally, spreading on fibrinogen and clot retraction were also upregulated in  $Flna^{loxP}PF4$ - $Cre^+$  mice. These upregulations could be the result of the FlnA mutation per se and/or of the different levels of FlnA in  $Flna^{loxP}PF4$ - $Cre^-$  mice (100%) versus  $Flna^{loxP}PF4$ - $Cre^+$ mice (49%). The only way to answer this question is to design a new mouse model expressing the same level of FlnA in both  $Flna^{loxP}PF4$ - $Cre^-$  and  $Flna^{loxP}PF4$ - $Cre^+$  mice.

In vivo results are highly contrasted with *in* vitro results. A rebleeding was observed in a large percentage of  $Flna^{loxP}PF4$ - $Cre^+$ mice (77%) compared to  $Flna^{loxP}PF4$ - $Cre^-$  mice (27%) indicating thrombus instability in mutated FlnA mice. Moreover, the *in* vivo thrombosis model induced by ferric chloride clearly showed the presence of embolies in arterioles (not in venules) of all FIna<sup>loxP</sup>PF4-Cre<sup>+</sup> mice (100%) compared to only 14% of Flna<sup>loxP</sup>PF4-Cre<sup>-</sup> mice. The apparent discrepancy between a gain-of-function in vitro and thrombus instability in vivo is interesting. A defect of platelet force generation, cytoskeletal reorganization, or fibrin formation are indeed hypotheses to study and require a full study in another paper.  $\alpha$ IIb $\beta$ 3 activation is regulated by talin, kindlin-3, and FLNa interactions with  $\beta$ 3 integrin where talin and kindlin-3 synergistically coactivate  $\alpha$ IIb $\beta$ 3<sup>32-34</sup>: talin modulating  $\alpha$ IIb $\beta$ 3 affinity and kindlin-3 modulating clustering and avidity.<sup>33</sup> In our thrombosis model, a defect in  $\alpha$ Ilb $\beta$ 3 clustering required for cytoskeleton reorganization could be the cause of the thrombus instability. Nonexclusive with the first hypothesis, cytoskeleton remodeling per se in response to mechanical forces could be affected by mutated FLNa. FLNa, described as an actin-crosslinking protein, is known to interact with 90 binding partners.<sup>35</sup> In addition to the dimerization domain,<sup>36</sup> the Ig repeats 24 and 23 of FLNa interact with different partners involved in cytoskeleton remodeling, notably the Rho GTPases (Ral1, Cdc42, Rac, and Rho)<sup>37,38</sup> and the GTPase FilGAP, specific for Rac. Dissociation occurs when FLNa-actin networks are subjected to mechanical shear strain.<sup>35</sup> In our model, we do not know the incidence of the extended (100 amino acids) C-terminal region of mutated FLNa on the dimerization and on the interaction partners with Ig repeat 24 and 23 domains and with the extended C-terminal region. Further experiments (proteomic analyses) will be required to identify new partners.

Finally, does the thrombus instability in knock-in mice correspond to clinical conditions for the patient? For obvious ethical reasons, the patient was not explored with *in vivo* tests. However, he underwent several surgery interventions, but for which the patient was unaware of any hemostatic outcomes. However, one must note that first, there may have been some bleeding, but not mentioned to the patient, if not dramatic; second, human surgery and mouse tail bleeding time may not be compared, because they are conducted in different conditions (different anesthetics, challenge of the wound [mouse] or not [surgery], use of local procoagulation during surgery, and so on).

In conclusion, this study shows evidence for thrombus instability of mutated FInA and raises a possible cardiovascular risk in the patient with this *FLNA* mutation.

#### AUTHOR CONTRIBUTIONS

F.A., A.K., L.L., J.S., C.S., and C.R., performed experiments; F.A., A.K., and M.B. designed experiments; F.A., A.K., H.R., J.-P.R., and M.B wrote the manuscript.

### FUNDING INFORMATION

This work was supported by INSERM and by a grant from Fondation pour la Recherche Médicale (LPC20170637458).

### CONFLICTS OF INTERESTS

The authors declare no competing financial interests.

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### SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

How to cite this article: Adam F, Kauskot A, Lamrani L, et al. A gain-of-function filamin A mutation in mouse platelets induces thrombus instability. *J Thromb Haemost*. 2022;20: 2666-2678. doi: 10.1111/jth.15864