The factor XIII V34L polymorphism accelerates thrombin activation of factor XIII and affects cross-linked fibrin structure

Robert A. S. Ariëns, Helen Philippou, Chandrasekaran Nagaswami, John W. Weisel, David A. Lane, and Peter J. Grant

Factor XIII on activation by thrombin cross-links fibrin. A common polymorphism Val to Leu at position 34 in the FXIII A subunit is under investigation as a risk determinant of thrombosis. Because Val34Leu is close to the thrombin cleavage site, the hypothesis that it would alter the function of FXIII was tested. Analysis of FXIII subunit proteolysis by thrombin using sodium dodecyl sulfate-polyacrylamide gel electrophoresis and high-performance liquid chromatography showed that FXIII 34Leu was cleaved by thrombin more rapidly and by lower doses than 34Val. Mass spectrometry of isolated activation peptides confirmed the predicted single methyl group difference demonstrated that the thrombin cleavage site is unaltered by Val34Leu. Kinetic analysis of activation peptide release demonstrated that the catalytic efficiency (kcat/Km) of thrombin was 0.5 for FXIII 34Leu and 0.2 (μmol/L)−1 × sec−1 for 34Val. Presence of fibrin increased the catalytic efficiency to 4.8 and 2.2 (μmol/L)−1 × sec−1, respectively. Although the 34Leu peptide was released at a similar rate as fibrinopeptide A, the 34Val peptide was released more slowly than fibrinopeptide A but more quickly than fibrinopeptide B generation. Cross-linking of γ- and α-chains appeared earlier when fibrin was incubated with FXIII 34Leu than with 34Val. Fully activated 34Leu and 34Val FXIII showed similar cross-linking activity. Analysis of fibrin clots prepared using plasma from FXIII 34Leu subjects by turbidity and permeability measurements showed reduced fiber mass/length ratio and porosity compared to 34Val. The structural differences were confirmed by electron microscopy. These results demonstrate that Val34Leu accelerates activation of FXIII by thrombin and consequently affects the structure of the cross-linked fibrin clot. (Blood. 2000;96:988-995)

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Introduction

On cleavage of fibrinopeptides A and B by thrombin, fibrin spontaneously polymerizes into a network of multimeric strands, initially held together by noncovalent interaction. Blood coagulation factor XIII (FXIII) is activated by thrombin, and activated FXIII (FXIIIa) covalently cross-links the fibrin clot to increase resistance to chemical, mechanical, and proteolytic insults. FXIII is a tetrameric protransglutaminase consisting of 2 A subunits, which contain the active site, and 2 B subunits, which serve a carrier function for the A subunit in plasma.1-2 FXIII is also found in platelets as an A-subunit dimer.2 Platelets do not release FXIII on activation, but lysis of platelets entrapped in the blood clot may increase local concentrations of FXIII.3 FXIIIa catalyzes the introduction of γ-glutamyl-ε-lysine peptide bonds between fibrin γ- and α-chains. Other substrates of FXIIIa are α2-antiplasmin,4 von Willebrand factor,5 thrombospondin,6 and fibronectin.7 Cross-linking of these substrates into the clot further contributes to the mechanical strength, viscosity, and resistance to proteolysis of fibrin. Activation of FXIII involves thrombin-induced cleavage of the peptide bond between Arg37 and Gly38 of the A subunit,2,9 resulting in the release of an amino-terminal activation peptide. In a second step, calcium induces dissociation of the A-subunit dimer from the B subunit.10,11 Both steps are essential for the activation of FXIII; the activation peptide release induces a conformational change in the A subunit and enables the dissociation of the subunits in the presence of calcium to unmask the catalytic site.

A common polymorphism with an allele frequency of around 25% has been identified in the FXIII A subunit (Val34Leu), 3 amino acids from the thrombin activation site.12 Recent studies have reported that the prevalence of the Leu encoding allele is lower in patients with myocardial infarction,13,14 deep vein thrombosis,15,16 and cerebral infarction17 when compared with matched control groups. These clinical studies suggest that this polymorphism may be a risk determinant of thrombosis in both the arterial and venous systems. Paradoxically, ex vivo and in vitro studies have suggested that possession of the Leu allele leads to increased cross-linking rates by FXIIIa.18-20 The mechanisms by which this occurred remained unclear. In view of the close proximity of the Val34Leu polymorphism to the activation site, we tested the hypothesis that this polymorphism alters the activation rate of FXIII by thrombin and affects fibrin structure.

Materials and methods

Blood sampling and processing

Venous blood was obtained from individuals with the homozygous FXIII 34Val, homozygous 34Leu, and heterozygous genotypes after an overnight

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fast. Blood was collected in 0.1 mol/L trisodium citrate. 9 parts of blood to 1 part of trisodium citrate. Within 1 hour after collection, the samples were centrifuged at 2560g for 20 minutes at room temperature to obtain platelet-poor plasma, frozen in aliquots in liquid nitrogen, and stored at −40°C until analysis.

Determination of the FXIII Val34Leu genotype

Genomic DNA was extracted with the BACC3 DNA extraction kit (Nucleon Biosciences Ltd, Glasgow, Scotland) from 10 mL venous blood that was anticoagulated with 1.6 mg/mL EDTA and from 5 mL of buffy coat obtained from the regional blood transfusion center. The FXIII Val34Leu genotype was determined by polymerase chain reaction and single-stranded conformational polymorphism analysis as previously described.13

Purification of the FXIII Val34Leu variants

Buffy coats from 34 outdated donations of human platelet-poor plasma were obtained from the regional blood transfusion center; genomic DNA was extracted and genotyped for the Val34Leu polymorphism. There were 19 homozygous FXIII 34Val samples, 1 homozygous 34Leu sample, and 14 heterozygous samples. FXIII 34Val was purified from a pool of 10 of the homozygous plasma donations (vol = 2.170 L) and homozygous FXIII 34Leu from a single plasma donation (vol = 0.205 L). In addition, FXIII was purified from the plasma of 40 outdated, mixed (unknown) genotype donations (vol = 11.895 L). Purification of FXIII was performed using a method adapted from previous publications.21,22 In brief, plasma was subjected to repeated precipitations with ammonium sulfate: 20% saturation at room temperature pH 7.0, 16% saturation at 4°C pH 5.4, 16% saturation at 4°C pH 7.0, and 36% saturation at 4°C pH 7.5. EDTA was added at 1 mmol/L to all the buffers used to prevent inopportune activation of FXIII. The final precipitate was resuspended in 1/1000 plasma volume of 0.05 mol/L Tris-HCl pH 7.5, 1 mmol/L EDTA, dialyzed against the same buffer and further purified by gel filtration on a Sephrose 6B column (2.6 × 40 cm), equilibrated and developed with 0.05 mol/L Tris-HCl pH 7.5, 1 mmol/L EDTA. Peak fractions containing both FXIII A and FXIII B subunit were pooled, concentrated on aquagel (Calbiochem Corp, La Jolla, CA) and extensively dialyzed against 0.05 mol/L Tris-HCl pH 7.5, 1 mmol/L EDTA. Purity and activity of the preparations were tested with sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), in-house FXIII A- and B-subunit sandwich enzyme-linked immunosorbent assays,23 and 5-(b-tinamido) pentylamine pentamethylene incorporation assay.23,24 Concentration of the preparations was measured by absorbance at 280 nm using an extinction coefficient of E1%1cm = 1.38.22

Cross-linking assay for FXIII

The FXIIIA-specific cross-linking activity was determined with a microtiter assay using fibrinogen and 5-(b-tinamido) pentylamine as substrates, as previously described.23,24 The assay is based on the incorporation of 5-(b-tinamido) pentylamine by FXIIIA into microtiter plates coated with fibrinogen, following activation with thrombin. The amount of cross-linked 5-(b-tinamido) pentylamine is detected by measuring phosphatase activity after incubation with a streptavidin-alkaline phosphatase conjugate. FXIIIA activity was measured before and after preincubation with human a-thrombin (Sigma Chemical, St Louis, MO) in plasma samples from subjects homozygous for FXIII 34Val, FXIII 34Leu, heterozygous, and in pooled normal plasma obtained from 47 healthy donors. Plasma samples (150 µL) were preincubated with 150 µL a-thrombin at a final concentration of 5 U/mL in microtiter plates, after which FXIIIA activity was measured with the pentylamine-incorporation assay in a separate plate. Samples were diluted 1/10 in Tromethamine-buffered saline (TBS) (40 mol/L Tris-HCl, 140 mol/L NaCl, 0.02% (w/v) NaN3, pH 8.3) containing 0.3 mg/mL of the synthetic peptide Gly-Pro-Arg-Pro-Ala-amide (Sigma) to prevent inopportune polymerization of fibrin in the preincubation mixture, which would interfere with the following subsampling procedure. After 30 minutes of preincubation, aliquots of 20 µL were subsampled into the pentylamine-incorporation plate containing 80 µL of the reaction mixture of 1.25 mol/mL 5-(b-tinamido) pentylamine, 0.63 mol/mL diethyloleitol, and 0.12 mol/mL CaCl2 in TBS. The pentylamine-incorporation assay was further performed as described.23

FXIII proteolysis studies using SDS-PAGE

The SDS-PAGE procedure was performed using a MiniProtein 3 (Biorad, Hercules, CA) electrophoresis unit. Gels were cast at a polyacrylamide concentration of 8% (bis/acrylamide ratio of 1:37.5) in 1.5 mol/L Tris-HCl, pH 8.8, and run at 150 V for 80 minutes. Gels were stained with Coomassie blue (2.5 g/L in 45% methanol, 10% acetic acid) for 30 minutes at room temperature and destained for 1 to 2 hours in 45% methanol, 10% acetic acid with 3 changes of destaining solution. Dose response of FXIII proteolysis by thrombin was studied by incubating 4.0 mol/L of purified FXIII variants with 0 to 10 U/mL human a-thrombin (Sigma) in 0.05 mol/L Tris-HCl, 1 mmol/L EDTA, pH 7.5 for 1 hour at 37°C. The reactions were stopped by the addition of equal (1:1) volume of reducing buffer 100 mmol/L Tris-HCl, 0.1 mol/L DTT, 4% SDS, 0.2% bromophenol blue, 20% glycerol, pH 6.8 with immediate boiling for 5 minutes, and 25 µL was loaded on the gel. The time course of FXIII proteolysis was studied by incubating 4.0 mol/L of purified FXIII variants with 0.4 U/mL human a-thrombin at 37°C. The reaction was stopped at 0 to 240 minutes as described for the dose-response reactions, and 25 µL was loaded on the gel.

Reverse-phase high-performance liquid chromatography (HPLC)

Reverse-phase HPLC was performed to analyze the FXIII activation peptide release using a 0.46 × 25 cm silica C18 (5 µm, 300 Å) column (Pepmap C18; Perseptive Biosystems Inc) on a Biocad Sprint automated chromatography system (also from Perseptive Biosystems, Framingham, MA), according to a method previously described by Janus and coworkers.25 The column was equilibrated with 1 column volume of 8% buffer A and 15% buffer B in 7 column volumes. The column was further washed with 2 column volumes of 100% buffer B followed by a linear gradient from 85% buffer A/15% buffer B to 100% buffer B in 7 column volumes. Flow-rate was set at 1 mL/min throughout the experiment. Buffer A consisted of 10% acetonitrile/90% 0.083 mol/L sodium phosphate pH 3.1 and buffer B of 40% acetonitrile/60% 0.083 mol/L sodium phosphate pH 3.1. All reagents were HPLC grade and solutions were filtered through 0.22 µm to eliminate particulates. Elution of peptides was detected by absorbance at 205 nm.

Kinetics of activation peptide release from FXIII

Molar quantities of the activation peptides analyzed by reverse-phase HPLC were calculated from the chromatograms by integration of the respective peak areas. The area under the curve was converted into molar quantity by comparison to that of a calibration FXIII activation peptide loaded at known concentration. Purified FXIII was dialyzed against 9.47 mol/L sodium phosphate, 137 mol/L NaCl, 2.5 mol/L KCl, 0.1% PEG, pH 7.4. Dose-response kinetics of FXIII activation peptide release by thrombin were performed by incubating 4.0 mol/L of dialyzed FXIII with 0, 0.01, 0.05, 0.1, 0.2, 0.3, 0.4, 0.5, 0.75, 1, and 10 U/mL (1 U/mL = 9.16 mmol/L) human a-thrombin (Sigma) for 1 hour at 37°C. The reactions were stopped by the addition of 1:10 (vol/vol) of 3 mol/L HClO4, centrifuged for 10 minutes at maximum speed in an Eppendorf centrifuge and 180 µL injected onto the C18 column. Time course of the activation peptide release reaction was studied by incubation of 4.0, 2.0, and 1.0 mol/L of FXIII with 0.5 U/mL human a-thrombin at 37°C. The reactions were stopped at 0, 2, 5, 10, 20, 30, 60, 120, 180, and 240 minutes by subsampling 10 volumes into 1 volume of 3 mol/L HClO4. The samples were centrifuged and 180 µL injected onto the C18 column. A similar method was used for studying the time course of FXIII activation peptide release in the presence of fibrin or fibrin and Gly-Pro-Arg-Pro-Ala-amide. In the presence of fibrin alone, 2.7 mmol/L of FXIII was incubated with 3.1 mmol/L human fibrinogen (Sigma) and 0.2 U/mL human a-thrombin. At this concentration of fibrinogen a clot formed, which was defined in size and did not interfere with the subsampling procedure. The reaction was stopped after 0.25, 0.5, 1, 2, 3, 4, 8, and 16 minutes of incubation and 270 µL injected onto the C18 column. In the presence of fibrin and the antipolymerizing peptide Gly-Pro-Arg-Pro-Ala, 2.7 mmol/L of FXIII was incubated with 3.1 mmol/L human fibrinogen (Sigma), 2 mg/mL Gly-Pro-Arg-Pro-Ala-amide (Sigma), and 0.5 U/mL
human α-thrombin. In these experiments, the reaction was stopped at 2, 4, 6, 8, 10, 12, 14, 16, 18, 20, and 22 minutes and 270 μL injected.

**Catalytic efficiency**

Catalytic efficiencies of the FXIII activation peptide release reaction by thrombin were calculated by fitting the data from the time-course and dose-response experiments analyzed with reverse-phase HPLC to the following equation:

\[
\frac{k_{cat}}{K_m} = \frac{-\ln(1 - \frac{[\text{AP}]}{[\text{AP}]})}{e \times t}
\]

where \([\text{AP}]\) is the concentration of FXIII activation peptide at a given time point, \([\text{AP}]\) the concentration of FXIII activation peptide at full activation, \(e\) the concentration of thrombin, and \(t\) time. Fitting of the data to this equation was performed using the Enzfit for Windows software version 2.0.8 (Biosoft, Cambridge, UK). The data obtained for activation in the presence of fibrin was also fitted to the equation, although it was recognized that this was an approximation due to the complex nature of enhanced FXIII activation by fibrin.

**Mass spectrometry**

Fractions from reverse-phase HPLC containing the FXIII activation peptides were collected for each purified FXIII Val34Leu variant. Molecular weights were analyzed with a single quadrupole, bench top mass spectrometer (Platform II, Micromass, Cheshire, UK). Samples were infused into the ionization source at a flow rate of 4 mL/min using a Harvard syringe pump. The mass spectrometer was fitted with a standard electrospray ionization source. Positive electrospray ionization was affected with a probe tip voltage of 3.5 kV, and a counter electrode voltage of 0.5 kV. Nitrogen was used as both the nebulizing and the drying gas; typical flow rates were a nebulizing gas flow rate of 20 L/h and a drying gas flow rate of 200 L/h. The sampling cone voltage was set to 50 volts. Data were acquired over the m/z range 500 to 2500. Spectra were transposed onto a true molecular mass scale using Maximum Entropy techniques. An external calibration was applied on a separate introduction of horse heart myoglobin (molecular weight, 16,951.49 d).

**Analysis of rates of α- and γ-chain cross-link formation**

Rates of α- and γ-chain cross-link formation were studied by incubating 6.1 μmol/L human fibrinogen (Sigma) with 0.1 μmol/L purified FXIII, 0.5 μmol/L human thrombin (Sigma) and 10 mmol/L calcium in 0.05 mol/L Tris-HCl, 0.15 mol/L NaCl, pH 7.5. The reaction was stopped by the addition of equal (1:1) volume of reducing loading buffer with immediate boiling for 5 minutes, and 25 μL was loaded on an 8% polyacrylamide gel. SDS-PAGE was further performed as described for the FXIII proteolysis studies.

**Turbidity measurements at 350 nm**

Plasma samples from subjects homozygous for either FXIII 34Leu (n = 3) or FXIII 34Val (n = 3) were diluted 2/3 with 0.05 mol/L Tris-HCl, 0.15 mol/L NaCl, pH 7.5 and incubated with 1 μmol/L human thrombin (Sigma) and 16 mmol/L calcium in a 0.5-mL cuvette. Immediately on addition of thrombin/calcium, absorbancy was read every 3 seconds at 350 nm for 5 minutes with a Perkin-Elmer Lambda 4B spectrophotometer. Parameters such as lag phase before start of fibrin polymerization, slope of the polymerization curve (A/min) and maximum absorbancy at full polymerization were recorded. Three replicate measurements were performed for each sample.

**Clot permeation measurements**

Plasma samples (100 μL) from subjects homozygous for either FXIII 34Leu (n = 3) or FXIII 34Val (n = 3) were incubated to allow clot formation with 1 μmol/L human thrombin (Sigma) and 20 mmol/L calcium in open tubes for 2 hours at room temperature in a wet chamber. The tubes containing the clots were connected via plastic tubing to a reservoir containing 0.05 mol/L Tris-HCl, 0.15 mol/L NaCl, pH 7.5 with a pressure drop of 4 cm. After washing the clots, flow rates of buffer through the fibrin gels were measured by timing 6 drops for each tube and weighing each drop for exact volume. Four replicate clots of each sample were analyzed in this way. The permeation coefficient or Darcy constant, which represents the surface of the gel allowing flow through a network and thus provides information on the pore structure, was calculated using the following formula:

\[
K_s = \frac{Q \cdot L}{A \cdot \Delta P}
\]

where Q is the volume of liquid (mL) with the viscosity η (10^{-2} poise) flowing through a clot with length L (1.3 cm) and a cross-sectional area A (0.049 cm²) in time t (seconds) under pressure ΔP (dyne/cm²). The unit of the resulting Darcy constant is cm².

**Scanning electron microscopy of fibrin clots**

After permeation experiments, the clots were fixed by permeating them with a 2% (v/v) glutaraldehyde solution overnight. Clots were recovered from the tubes and further processed by dehydration using a stepwise ethanol gradient, critical point drying, and sputter coating with gold-palladium as previously described. Plasma clots from 3 homozygous FXIII 34Leu subjects and 3 homozygous FXIII 34Val subjects were observed and photographed digitally in at least 6 different areas using a scanning electron microscope (XL 20, Philips Electron Optics, Eindhoven, The Netherlands).

**Results**

**Effect of preactivation on FXIIIa cross-linking activities**

Plasma samples from subjects homozygous for FXIII 34Leu showed greater activity in a pentylamine-incorporation assay, in which there is thrombin activation during the course of the assay, than plasma samples from subjects homozygous for FXIII 34Val or heterozygous subjects. Cross-linking rates in this assay were typically higher for FXIII 34Leu than for FXIII 34Val, with an intermediate response to heterozygous samples and pooled normal plasma (Figure 1A). After preincubation with 5 U/mL human α-thrombin for 30 minutes at 37°C, however, this difference disappeared, and cross-linking activities of plasma samples from subjects homozygous for FXIII 34Leu or FXIII 34Val and heterozygous subjects were similar (Figure 1B).

**Increased proteolysis of FXIII 34Leu by thrombin**

Proteolysis of FXIII subunits by thrombin was studied using SDS-PAGE. Electrophoresis of FXIII on SDS-polyacrylamide gels under reducing conditions showed 2 bands for the A and B subunit migrating very closely together. Activation by thrombin produced a distinct band for the activated A subunit that migrated faster in the gel. We first examined activation of purified FXIII 34Leu and FXIII 34Val by increasing doses of human α-thrombin. Figure 2 shows that FXIII 34Val (panel B) proteolysis started with 0.3 to 0.4 U/mL of thrombin after 1 hour incubation at 37°C, whereas FXIII 34Leu (panel A) proteolysis started with 0.1 to 0.2 U/mL of thrombin and was complete with 1 U/mL. We next investigated FXIII cleavage by adding 0.4 U/mL human α-thrombin and stopping the reaction after increasing times. Proteolysis of FXIII 34Val (panel D) started after 60 to 90 minutes, whereas FXIII 34Leu cleavage (panel C) by the same dose of thrombin started after 10 to 20 minutes and was complete after 120 minutes.

**Analysis of FXIII activation peptides**

The preactivation and SDS-PAGE studies suggested that release of the activation peptide from FXIII 34Leu is more rapid than that from FXIII 34Val, but the incomplete resolution of the bands in
SDS-PAGE did not allow quantification. Reverse-phase HPLC was used to quantify the amount of peptide released from each form of FXIII by thrombin. When activating purified FXIII 34Val with 10 U/mL human α-thrombin for 1 hour at 37°C, one distinct peak for the activation peptide was detected. Activating FXIII purified from a mixed genotype, however, produced 2 peaks, and analysis of activated purified FXIII 34Leu indicated that the 2nd peak represented the 34Leu activation peptide (Figure 3). This separation of the 2 species of activation peptides on the C18 column was confirmed by mass spectrometry. HPLC fractions containing the respective peptide peaks were collected and analyzed for the molecular weight of their peptides. The first peak contained a peptide of molecular weight of 3951.5 d and the second peak a peptide of 3965.0 d. These results confirmed that the first peak contains the FXIII 34Val activation peptide and the 2nd peak the FXIII 34Leu activation peptide. The difference in molecular weight between the peptides can be accounted for by the expected single methyl group (14.0 d) difference between the Leu and Val side chains. They also demonstrated that the thrombin cleavage site between Arg37 and Gly38 is unaltered by the Val34Leu polymorphism.

**Kinetics of FXIII activation**

The separation of the FXIII 34Val from the FXIII 34Leu activation peptide on reverse-phase HPLC allowed us to carry out kinetic
analysis of the activation reaction for each form of FXIII using purified FXIII from a mixed genotype. Purified FXIII was incubated at 4.0 µmol/L with increasing concentrations of human \( \alpha \)-thrombin. HPLC was performed on the incubation mixtures. Peak areas of the FXIII 34V al and FXIII 34Leu activation peptides were calculated from the chromatograms and transformed in molar quantities. More FXIII 34Leu activation peptide was released than FXIII 34V al activation peptide after incubation with thrombin concentrations between 0.01 and 0.5 U/mL for 1 hour at 37°C (Figure 4A), confirming the findings from the gel electrophoresis studies that FXIII 34Leu is activated by lower doses of thrombin than FXIII 34V al. Time-course experiments were also performed. Purified FXIII at 4.0, 2.0, and 1.0 µmol/L was activated with 0.5 U/mL human \( \alpha \)-thrombin at 37°C and the reaction was stopped at increasing times. More FXIII 34Leu activation peptide was released than FXIII 34Val activation peptide early in the reaction (Figure 4B). This faster initial rate was observed regardless of the relative concentrations of FXIII and thrombin; kinetic data are summarized in Table 1.

**Effect of fibrin on the activation peptide release reaction**

Fibrinogen was added at a concentration of 3.1 µmol/L to 2.7 µmol/L purified FXIII and 0.2 U/mL human \( \alpha \)-thrombin. The inset of Figure 5 shows a typical chromatogram from these experiments, with defined peaks for fibrinopeptide A, fibrinopeptide B, FXIII 34Val activation peptide, and FXIII 34Leu activation peptide. In the presence of fibrinogen, the release of both FXIII activation peptides by thrombin was greatly accelerated. FXIII 34Leu activation peptide release remained significantly faster than that of FXIII 34Val. The FXIII 34Leu activation peptide was released at a similar rate as fibrinopeptide A, followed by release of the FXIII 34V al peptide and finally fibrinopeptide B (Figure 5).

**Catalytic efficiency of FXIII activation**

Catalytic efficiencies of thrombin-dependent activation peptide release from the different polymorphic forms of FXIII were calculated in the absence of fibrin, in the presence of fibrin alone, and in the presence of both fibrin and Gly-Pro-Arg-Pro-amide. The 34Leu form showed twice the catalytic efficiency, regardless of the experimental conditions (Table 1). The presence of fibrin greatly enhanced FXIII activation peptide release by thrombin and this catalytic effect was reduced when polymerization of fibrin was inhibited with Gly-Pro-Arg-Pro-amide. The difference in catalytic efficiency between FXIII 34Val and FXIII 34Leu remained in the presence of polymerizing fibrin.

<table>
<thead>
<tr>
<th>Time courses</th>
<th>Dose response</th>
<th>Mean ± SD (n = 4)</th>
<th>+FI +GPRP (n = 3)</th>
<th>+FI –GPRP (n = 4)</th>
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<tr>
<td>k_{cat}/k_{on} (µmol/L)^{-1} × sec^{-1}</td>
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<td>1.0 µmol/L</td>
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<td>0.41</td>
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</tbody>
</table>
Effect of FXIII 34Leu on \( \alpha \)- and \( \gamma \)-chain cross-linking

Cross-linking of the fibrin \( \alpha \)- and \( \gamma \)-chains by the 2 forms of FXIII was studied using reducing SDS-PAGE. Cross-linking of both \( \gamma \)-chains and \( \alpha \)-chains appeared earlier when incubating a mixture of fibrinogen and thrombin with purified FXIII 34Leu than with FXIII 34Val (Figure 6). The fibrinogen preparation used for these experiments had some contamination with fibronectin and FXIII. Due to traces of FXIII in the preparation, some \( \gamma \)-chain dimers were already present without incubation with the isolated FXIII variants. Nonetheless, more \( \gamma \)-chain dimers and \( \alpha \)-chain polymers were formed by incubation with FXIII 34Leu than FXIII 34Val after 5 minutes. After 20 minutes incubation FXIII 34Leu produced more of high molecular weight \( \alpha \)-chain polymers that did not enter the gel and remained in the loading well (Figure 6).

FXIII Val34Leu and fibrin structure

Plasma samples from subjects homozygous for either form of FXIII Val34Leu were used to study effects of this polymorphism on fibrin structure. The lag phase in formation of turbidity at 350 nm was shorter for homozygous FXIII 34Leu (1.08 ± 0.11 minutes) than for homozygous FXIII 34Val samples (1.55 ± 0.15 minutes). The slope of the turbidity curves did not differ, but maximum absorbancy after 5 minutes was slightly lower for the FXIII 34Leu (1.83 ± 0.04) than FXIII 34Val (1.93 ± 0.03), indicating the presence of thinner fibrin fibers in the FXIII 34Leu samples. Consistent with these results, permeation of fibrin clotted for 2 hours was found to be slower in clots from FXIII 34Leu samples when compared with FXIII 34Val. The Darcy constant, which reflects the surface area of the clot available for flow, was 3.6 ± 0.5 \( \times \) 10\(^{-9} \) cm\(^2\) for FXIII 34Leu and 8.7 ± 4.4 \( \times \) 10\(^{-9} \) cm\(^2\) for FXIII 34Val. These structural differences were confirmed by scanning electron microscopy. Clots prepared from plasma samples homozygous for FXIII 34Val (Figure 7A) showed a fibrin meshwork with thick fibers and large pores, whereas clots prepared from plasma samples homozygous for FXIII 34Leu showed a finer fibrin meshwork with thinner fibers and reduced space between the fibrin strands (Figure 7B).

Discussion

The FXIII gene is known to be highly polymorphic, with several common nucleotide sequence variations within the population, which encode amino acid substitutions. Although these amino acid sequence changes could alter the levels and activity of FXIII, to date there has been little formal examination of their effects. We report here the functional characterization of the Val34Leu polymorphism in the A subunit. This polymorphism is located adjacent to the thrombin cleavage activation site of FXIII. Presence of the 34Leu encoding allele has been associated with a protective effect against thrombotic disease in the arterial and venous systems.\(^{13-17}\)

In this study, it is demonstrated that the nature of the amino acid substitution at position 34 influences the activation rate of FXIII. Using SDS-PAGE, the rate of proteolysis of the FXIII A subunit is shown to be faster and to occur at lower thrombin concentrations.
when Leu rather than Val is at position 34. HPLC analysis demonstrates that proteolysis of FXIII occurs at the activation cleavage site (by direct determination of the precise Mr of the activation peptide) and that the catalytic efficiency of thrombin cleavage of the activation peptide is increased approximately 2-fold by substitution of Val by Leu.

Prior studies of the kinetics of activation peptide release from FXIII have used pooled donations of plasma to prepare the protein, and this was carried out without knowledge of the genotype. The estimates of catalytic efficiency obtained in those studies, kcat/Km 0.13 to 0.16 (µmol/L)^-1 × sec^-1,25 agree with that derived in the present study for FXIII 34Val. We now find that the 34Val and 34Leu activation peptides separate on a C18 column, with the 34Val peptide eluting before 34Leu. Also in the study by Janus and coworkers,25 a second eluting peak was observed when the FXIII activation was analyzed by HPLC. Amino acid analysis on the collected fractions demonstrated 1.8 and 2.7 leucine residues in the first and second peaks, respectively. The authors concluded that the second peak was FXIII related, but were unable to identify the presence of the Val34Leu polymorphism in the variants. The second peak was ignored in the reported kinetic analysis, and in retrospect, the catalytic efficiency reported by Janus and colleagues, therefore, appears to be related solely to FXIII 34Val.

Characterization of the later eluting peak in the present study has enabled its identity to be determined as the activation peptide from FXIII 34Leu. This information has allowed an estimate to be made of the catalytic efficiency of its cleavage by thrombin, kcat/Km 0.49 (µmol/L)^-1 × sec^-1. The estimates obtained were robust in relation to the relative ratio of thrombin to its substrate over a range of concentrations of both thrombin and its substrate. In the presence of fibrin, the catalytic efficiencies of FXIII activation peptide release were further increased to 2.15 and 4.81 (µmol/L)^-1 × sec^-1 for FXIII 34Val and FXIII 34Leu, respectively, maintaining the polymorphism-related differences. These fibrin-induced increases are again compatible with previous reports, despite the lack of a rigorous kinetic approach here compared with the prior studies.25-28 Polymerization of fibrin appears to play a role in the FXIII activation process, because inhibition of polymerization with Gly-Pro-Arg-Pro-amide significantly reduced the catalytic function of fibrin in the FXIII activation peptide release reaction.

The finding of an accelerated activation rate of FXIII 34Leu suggests that the pentylamine-incorporation assay, as it is routinely used for screening of FXIII activity levels, is particularly sensitive to the FXIII activation step. The result also provides an explanation for the relatively poor correlation between FXIII activity levels and FXIII A-subunit antigen levels,19,20,23 because the wide variability in activity in relation to the Val34Leu polymorphism would largely override the effect of relatively minor fluctuations in the A-subunit levels under normal conditions. Kangsadampai and Board expressed the 34Val and 34Leu forms of FXIII in vitro and also found that FXIII 34Leu showed higher activity in a pentylamine-incorporation assay.19 However, activation kinetics were not investigated, and their results could have also been related to the increased activation rate of FXIII 34Leu. We found that, once activated, the specific cross-linking activities of the 2 forms of FXIII do not differ.

When analyzing the sequence of the release of the FXIII activation peptides in relation to fibrinopeptides A and B, it was found that the FXIII 34Leu activation peptide was released at approximately the same rate as fibrinopeptide A, whereas release of the FXIII 34Val activation peptide was slower, with a rate in-between those of the 2 fibrinopeptides. These findings suggest that cross-linking activity by FXIII 34Leu is generated at the time of desA fibrin formation, whereas cross-linking activity by FXIII 34Val is generated more at a time when both fibrinopeptides A and B are cleaved from fibrinogen. It has to be borne in mind, however, that the experimental conditions of this experiment are different from those encountered in vivo. Whereas thrombin and fibrinogen concentrations could be considered to resemble those attainable under in vivo conditions, the FXIII concentration of 3.1 µmol/L, necessary to obtain measurable peak areas of the activation peptide by HPLC, was significantly higher than the physiologic plasma concentration of about 0.07 µmol/L.

The observed differences in cleavage of the 2 species of FXIII activation peptides in temporal relation to fibrinopeptide A and B release prompted us to investigate the effect of the FXIII Val34Leu polymorphism on the structure of the cross-linked fibrin clot. In agreement with the accelerated activation of FXIII 34Leu, analysis by SDS-PAGE showed that cross-linking of fibrin α- and γ-chains by FXIII 34Leu occurred earlier than that by FXIII 34Val. At later time points, both forms of FXIII generated similar amounts of cross-links. Polymerization, as followed by turbidity, was also faster in clotting plasma samples homozygous for FXIII 34Leu than in those homozygous for FXIII 34Val. After the initial more rapid rise of turbidity in the FXIII 34Leu samples, however, total turbidity at 5 minutes was lower than that in FXIII 34Val samples, indicating the presence of thinner fibrin fibers.

Biophysical and structural properties of fibrin in homozygous plasma samples clotted for 2 hours, allowing cross-linking by FXIII to reach completion, were analyzed using permeation characteristics of the fibrin gel in addition to electron microscopy. Fibrin that was fully cross-linked by FXIII 34Leu showed a finer structure with thinner fibers and smaller pores, than fibrin cross-linked by FXIII 34Val. Previous studies have shown that the delay in fibrinopeptide B release compared to the relatively fast release of fibrinopeptide A appears to be necessary for enhanced lateral aggregation and thicker fibers.31,32 Our present findings suggest that the accelerated generation of covalent cross-linking activity by FXIII 34Leu influences the lateral aggregation process and affects the molecular structure of the cross-linked fibrin clot. It appears that early cross-linking of fibrin by FXIII 34Leu, which is activated at the time of fibrinopeptide A release, inhibits lateral aggregation of the fibrin fibers, whereas delayed cross-linking by FXIII 34Val, which is activated at the time of desAB fibrin formation, allows for more lateral aggregation.

This study has demonstrated that the Val34Leu polymorphism in the A subunit affects the function of FXIII by increasing the rate of FXIII activation by thrombin and by altering the molecular structure of the cross-linked fibrin gel to one with reduced mass/length ratio of the fibers. The potential consequences of fibrin cross-linked by FXIII 34Leu, with a finer meshwork, thinner fibers, and altered permeation characteristics, on clinical (vascular) outcome remain to be elucidated.

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References


