



Evaluating factor XIII specificity for glutamine-containing substrates using a matrix-assisted laser desorption/ionization time-of-flight mass spectrometry assay



Prakash G. Doiphode, Marina V. Malovichko, Kelly Njine Mouapi, Muriel C. Maurer*

Department of Chemistry, University of Louisville, Louisville, KY 40292, USA

ARTICLE INFO

Article history:

Received 23 November 2013

Received in revised form 26 March 2014

Accepted 11 April 2014

Available online 19 April 2014

Keywords:

Factor XIII

Transglutaminase

Coagulation

Substrate specificity

Kinetics

Mass spectrometry

ABSTRACT

Activated factor XIII (FXIIIa) catalyzes the formation of γ -glutamyl- ϵ -lysyl cross-links within the fibrin blood clot network. Although several cross-linking targets have been identified, the characteristic features that define FXIIIa substrate specificity are not well understood. To learn more about how FXIIIa selects its targets, a matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS)-based assay was developed that could directly follow the consumption of a glutamine-containing substrate and the formation of a cross-linked product with glycine ethylester. This FXIIIa kinetic assay is no longer reliant on a secondary coupled reaction, on substrate labeling, or on detecting only the final deacylation portion of the transglutaminase reaction. With the MALDI-TOF MS assay, glutamine-containing peptides derived from α_2 -antiplasmin, *Staphylococcus aureus* fibronectin binding protein A, and thrombin-activatable fibrinolysis inhibitor were examined directly. Results suggest that the FXIIIa active site surface responds to changes in substrate residues following the reactive glutamine. The P₋₁ substrate position is sensitive to charge character, and the P₋₂ and P₋₃ substrate positions are sensitive to the broad FXIIIa substrate specificity pockets. The more distant P₋₈ to P₋₁₁ region serves as a secondary substrate anchoring point. New knowledge on FXIIIa specificity may be used to design better substrates or inhibitors of this transglutaminase.

© 2014 Elsevier Inc. All rights reserved.

In blood coagulation, the serine protease thrombin cleaves the N-terminal portions of the fibrinogen α and β chains. The resultant fibrin monomers polymerize noncovalently into linear protofibrils and then laterally into fibers, forming a blood clot that is held together by noncovalent forces [1,2]. To make this blood clot more resistant to degradation, covalent cross-links are introduced into the fibrin network [3,4]. Thrombin aids in this process by helping to activate factor XIII (FXIII)¹ via cleavage of the FXIII R37–G38 activation peptide bond. The resultant transglutaminase, FXIIIa, then catalyzes the formation of γ -glutamyl- ϵ -lysyl cross-links within fibrin and fibrin–ligand complexes [5,6]. The reaction mechanism of FXIIIa involves the thiol group of Cys314 attacking the carboxyam-

ide of a reactive glutamine (Q) side chain. Ammonia is released and a thioester is generated. Deacylation then occurs as the amino group from a Lys (K) side chain targets the thioester. The resultant product of the transglutaminase-catalyzed reaction contains an isopeptide bond between the side chains of the Q and K residues [5].

FXIIIa targets several sites within the fibrin $\alpha\beta\gamma$ environment, leading to formation of a rigid blood clot [7]. Other key FXIIIa substrates include α_2 -antiplasmin (α_2 AP), fibronectin, thrombin-activatable fibrinolysis inhibitor (TAFI), plasminogen activation inhibitor-1, vitronectin, and *Staphylococcus aureus* fibronectin binding protein A [8]. Although a number of FXIIIa substrates have been identified, an obvious consensus sequence for the Q-containing substrate has been difficult to establish [6,8–11].

A variety of methods have been developed to study interactions between FXIIIa and its substrates. In several assays, the lysine-like substrate is detected by colorimetric [12,13], fluorescent [14,15], or radioactive [16–18] methods. With this approach, the glutamine-containing reaction must occur before the lysine-like reaction can be monitored. Another strategy for measuring enzymatic activity is a coupled ultraviolet–visible (UV–Vis) assay in which ammonia released from the reactive glutamines is detected via a separate

* Corresponding author.

E-mail address: muriel.maurer@louisville.edu (M.C. Maurer).

¹ Abbreviations used: FXIII, factor XIII; FXIIIa, activated FXIII; α_2 AP, α_2 -antiplasmin; TAFI, thrombin-activatable fibrinolysis inhibitor; UV–Vis, ultraviolet–visible; MALDI–TOF MS, matrix-assisted laser desorption/ionization time-of-flight mass spectrometry; GEE, glycine ethylester; Fnb A, fibronectin binding protein A; EDTA, ethylenediaminetetraacetic acid; TFA, trifluoroacetic acid; α -CHCA, α -cyano-4-hydroxycinnamic acid; ZED1301, FXIIIa inhibitory peptide (Ac-Asp-MA-Nle-Leu-Pro-Trp-Pro-OH); MA, Michael acceptor unit.

secondary colorimetric assay [19–22]. This resultant ammonia is used to convert α -ketoglutarate to glutamate in a nicotinamide adenine dinucleotide (NADH)-dependent reaction that is monitored at 340 nm. Again, the glutamine-containing reaction is not being monitored directly. An additional method is to work with synthetic peptides in which the reactive glutamine is replaced with colorimetric Glu-pNA or fluorimetric Glu-AMC [23,24]; with such reporter groups, the direct release of *p*-nitroaniline (pNA) or amidomethylcoumarin (AMC) can finally be monitored and is a real strength of this assay design. A drawback is that each substrate needs to have this modified Glu residue introduced into the sequence. Moreover, the bulky reporter groups may hinder interactions with the FXIIIa active site surface.

A valuable advance for monitoring FXIIIa kinetics would be an assay that directly records consumption of substrate and formation of product within a single platform. The use of low sample volumes is also desirable. Our newly developed matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS) kinetic assay achieves these goals (Fig. 1). In the assay, the FXIIIa-catalyzed reaction between a Q-containing substrate and excess glycine ethylester (GEE) is followed. GEE is routinely used as a lysine mimic in the coupled UV-Vis assays [19–22]. The individual transglutaminase reactions are quenched at distinct time points, and MALDI-TOF mass spectra are collected to monitor for loss of Q-containing substrate and gain of the cross-linked product (Q-containing substrate)-GEE (gain of 86 *m/z*). Overall, the MALDI-TOF MS kinetic assay is no longer reliant on a secondary coupled reaction, on substrate labeling, or on detecting only the final deacylation portions of the transglutaminase reaction [12,22,23].

A series of glutamine-containing substrates were chosen to further test the MALDI-TOF MS assay. Using a nomenclature similar to that of the proteases, the reactive Q of FXIIIa substrates occupies the P₁ position. Amino acids can be assigned as...P₄ P₃ P₂ P₁^(reactive Q) and then to the right of the reactive Q as P₋₁ P₋₂ P₋₃ P₋₄... (Table 1). Regions to explore in kinetic studies include the P₋₁ and P₋₂ positions located near the reactive Q residue and also the putative substrate recognition segment P₋₈ to P₋₁₁ [20,21,25]. This P₋₈ to P₋₁₁ segment is proposed to serve

as an extra substrate anchor onto the extended FXIIIa active site region. Three FXIIIa substrates containing reactive Q residues are α_2 AP, *S. aureus* fibronectin binding protein A (FnB A), and TAFI (Table 1). Characteristics of these substrates are described below.

α_2 AP (464 residues) serves as a potent inhibitor of the fibrinolytic agent plasmin. Cleavage of the P12–N13 amide bond generates a 452-residue protein starting with an Asn [26,27]. FXIIIa rapidly cross-links Q2 of N1– α_2 AP to K303 of the fibrin(ogen) α -chain [26,28,29]. While this N1– α_2 AP is tethered to a blood clot, its C-terminal domain remains free to inhibit plasmin [26]. Previous coupled UV-Vis kinetic assays with α_2 AP (1–15) (Table 1) have indicated that the Q2 at the P₁ position is the reactive glutamine of this peptide with the Q4 at the P₋₂ position, serving a supporting role in binding [20]. α_2 AP also contains a ¹⁰LLKL¹³ stretch (P₋₈ to P₋₁₁ region) that is part of a putative substrate recognition exosite [20,25].

S. aureus colonizes human tissue during vascular injury by linking its FnB A to fibrinogen [30,31]. Similar to the α_2 AP (1–15) sequence, FnB A contains glutamines at the P₁ and P₋₂ positions (Table 1) [32–34]. By contrast, *S. aureus* FnB A has a basic R at the P₋₁ position, whereas α_2 AP (1–15) has an acidic E. As found in α_2 AP, *S. aureus* FnB A possesses K residues at the P₋₈ and P₋₉ positions.

TAFI modifies lysine residues within the fibrin network, thereby reducing the affinity of plasminogen for the blood clot. As a result, less plasmin is generated for fibrinolysis [14,35,36]. Similar to α_2 AP and *S. aureus* FnB A, TAFI also contains two glutamines (Table 1). However, with TAFI, a polar serine and the small flexible glycine are situated between the two glutamines. Thus, the second Q residue is switched from the P₋₂ position to the P₋₃ position. As observed with α_2 AP and *S. aureus* FnB A, TAFI (1–15) contains a basic arginine within the P₋₈ to P₋₁₁ region.

Studies to characterize how FXIIIa selects its substrate targets will benefit from the development of assays that directly follow the Q-containing substrate step. Our newly developed MALDI-TOF MS kinetic assay achieves this goal by recording the consumption of the Q-containing substrate and formation of the Q-GEE product within a single assay platform. This MS-based assay is no longer reliant on a secondary coupled reaction, on substrate labeling, or on detecting only the final deacylation portions of the transglutaminase reaction. Using our direct monitoring assay, the kinetic properties of α_2 AP (1–15), *S. aureus* FnB A (100–110, 100–114), and TAFI (1–15) were successfully determined. Aside from native peptide sequences, individual E, S, and/or R substitutions were also examined. Results indicate that the FXIIIa active site surface responds to changes in the substrate residue positions P₋₁ and P₋₂ and also to those of a more distant exosite involving the P₋₈ to P₋₁₁ residues. Additional studies with this MALDI-TOF MS assay will aid in further deciphering the key features of a good FXIIIa substrate or inhibitor.

Materials and methods

Materials for MALDI-TOF assay

Human cellular FXIII expressed in *Saccharomyces cerevisiae* was kindly provided by Paul Bishop (ZymoGenetics, Seattle, WA, USA). The lyophilized FXIII was reconstituted in 18 M Ω deionized water, aliquoted, and stored at –70 °C until future use. The concentration of the FXIII A₂ stock solutions were determined on a Cary 100 UV-Vis spectrophotometer using an extinction coefficient of 1.49 (ml mg⁻¹ cm⁻¹) at 280 nm. Plasma-derived bovine thrombin (Sigma-Aldrich) was used to activate the FXIII A₂. All of the Q-containing substrate peptides (Table 1) were synthesized and purified by New England Peptide (Gardner, MA, USA). The

MALDI-TOF Mass Spectrometry Based Kinetic Assay

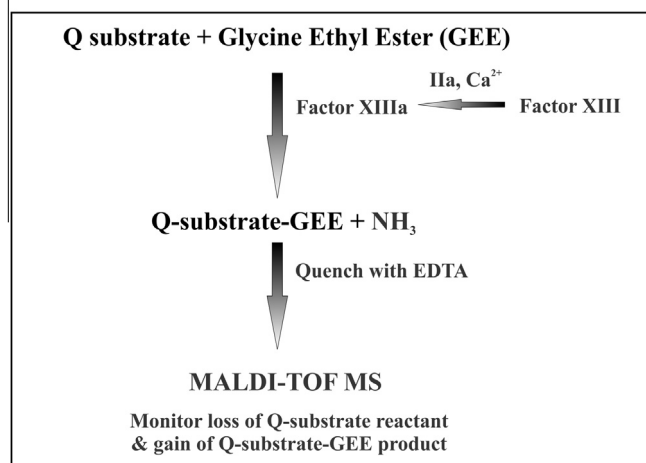


Fig. 1. Flow chart describing the MALDI-TOF MS assay. In this assay, factor XIII is activated by thrombin in the presence of calcium. The resultant FXIIIa is then responsible for catalyzing the reaction between the Q-containing substrate (Q-substrate) and glycine ethylester (GEE) that serves as the lysine mimic. Ethylenediaminetetraacetic acid (EDTA) is added to scavenge calcium away from the FXIIIa and quench the transglutaminase reaction. MALDI-TOF MS is used to monitor loss of the Q-substrate reactant and formation of the Q-substrate-GEE product.

Download English Version:

<https://daneshyari.com/en/article/1172792>

Download Persian Version:

<https://daneshyari.com/article/1172792>

[Daneshyari.com](https://daneshyari.com)