Proteolytic Inactivation of ADAMTS13 by Thrombin and Plasmin

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Abbreviations: ADAMTS – A Disintegrin And Metalloproteinase with ThromboSpondin type-1 repeats, TF – tissue factor, TSP1 - thrombospondin type-1 repeat, TTP – thrombotic thrombocytopenic purpura, VWF – von Willebrand factor, VWF:CBA - VWF collagen binding assay

1
Abstract

The multimeric size and the function of circulating von Willebrand factor are modulated via its proteolytic cleavage by the plasma metalloproteinase, ADAMTS13. It is unclear how ADAMTS13 activity is regulated within the vascular system. In the absence of a regulatory mechanism, ADAMTS13 activity might compromise platelet adhesion at sites of vascular injury. We hypothesised that at sites of vascular injury, ADAMTS13 activity could be regulated locally by coagulation proteinases. Initiation of coagulation in human plasma resulted the disappearance of added full-length recombinant ADAMTS13. This loss was inhibited by hirudin. Using purified proteins, we showed that ADAMTS13 is proteolysed at several cleavage sites by thrombin in a time- and concentration-dependent manner. Furthermore, this proteolysis ablated ADAMTS13 activity against purified von Willebrand factor. Preincubation of thrombin with soluble thrombomodulin, but not heparin, inhibited the proteolysis of ADAMTS13, suggesting the involvement of thrombin exosite I (and not exosite II) in ADAMTS13 recognition. Plasmin also cleaved ADAMTS13 into similar fragments, resulting in the loss of ADAMTS13 activity. This study demonstrates the susceptibility of ADAMTS13 to proteolytic inactivation and suggests possible roles for thrombin and plasmin at sites of vascular injury.
Introduction

Von Willebrand factor (VWF) is a large (2050 amino acid/~250 kDa) multidomain glycoprotein whose functions are critical for normal haemostasis\(^1\). VWF has two principal roles that influence the haemostatic process - a) it acts as a carrier protein for coagulation factor VIII\(^2\), b) it mediates rapid adhesion of platelets to sites of vascular perturbation. The latter is one of the first haemostatic events following endothelial disruption and occurs through the specific binding of VWF to exposed sub-endothelial matrix proteins (principally collagen)\(^3\). Once immobilised, VWF affinity for the GP Ib-IX-V receptor complex on the surface of circulating platelets is significantly enhanced\(^4\). This results in the tethering of platelets at sites of vascular damage and the formation of a primary platelet plug. Tethered platelets are subsequently activated and expose phosphatidylserine-rich surfaces that are critical for efficient thrombin generation to occur\(^5\).

VWF is constitutively secreted into the blood by endothelial cells as multimers of varying size that differ predominantly in the number of component VWF units. A significant proportion is also stored within Weibel-Palade bodies, predominantly as “ultra-large” multimers (UL-VWF)\(^6\) that may exceed 2x10\(^4\) kDa\(^1\). This pool is released upon demand in response to endothelial cell activation\(^7\).

The properties of circulating VWF are, in part, dependent upon its molecular size. Larger VWF multimers not only bind circulating platelets more readily than smaller forms, but also undergo marked conformational changes in response to the rheological forces exerted by the circulating blood\(^8\). Under normal flowing conditions, VWF multimers circulate in a globular form. However, when VWF is exposed to increased shear forces, these molecules unravel into a “string-like” conformation. This increases the number of exposed platelet/matrix binding sites and thus enhances the platelet tethering potential of the VWF molecule.
Thrombotic thrombocytopenic purpura (TTP) is rare, life-threatening condition, linked to UL-VWF function. These patients develop platelet-rich thrombi in their microvasculature, which may lead to renal failure, neurological dysfunction and ultimately, death. Such clinical sequelae are, at least in part, related to the elevated plasma UL-VWF levels detected in affected individuals. These increased UL-VWF levels may promote platelet-rich thrombus formation by more readily tethering/aggregating circulating platelets. However, the precise stimulus that precipitates an episode of TTP remains unclear. The cause of increased UL-VWF levels in most TTP patients has been attributed to deficiency in ADAMTS13. This enzyme cleaves VWF in the A2 domain at the Y1605-M1606 bond and so converts UL-VWF into smaller forms that demonstrate a reduced adhesive/prothrombotic potential. Consequently, TTP sufferers with congenital defects in ADAMTS13, or those with autoantibody inhibitors that compromise ADAMTS13 function, are unable to process UL-VWF normally.

ADAMTS13 is secreted into the blood as an active enzyme with a plasma concentration of ~1 µg/ml. It is a stable enzyme with a plasma half-life of ~2-3 days. Plasma ADAMTS13 is a 1349 residue protein with a molecular mass of 180-200 kDa. The metalloproteinase domain contains a highly conserved active site that includes three histidine residues that coordinate a Zn$^{2+}$ ion. The disintegrin-like domain and the first thrombospondin type-1 repeat (TSP1) domain precede the cysteine-rich domain and spacer domain, followed by seven consecutive TSP1 and two C-terminal CUB domains.

It is uncertain how ADAMTS13 activity is both regulated and localised. ADAMTS13 cleaves VWF inefficiently in the absence of shear forces, or a denaturant (e.g. urea), that “unravels” VWF and permits access of ADAMTS13 to the scissile bond. Although VWF is normally protected from proteolysis by its tertiary structure, it seems unlikely that ADAMTS13
activity in blood is not specifically regulated. In this study, we describe the proteolytic cleavage/inactivation of ADAMTS13 by thrombin and FXa. Moreover, we also demonstrate the proteolytic inactivation of ADAMTS13 by plasmin, another protease generated specifically at sites of vessel damage, following thrombin-dependent fibrin deposition.
Materials and Methods

Expression and Purification of Recombinant ADAMTS13

Full-length ADAMTS13 cDNA (a gift from Dr F Sheiflinger, Baxter, Austria\textsuperscript{18}) was cloned in-frame into the mammalian expression vector, pcDNA3.1myc/His, which fused a myc epitope and polyhistidine tag to the C-terminus of ADAMTS13. Stably transfected HEK293 cells were generated by standard techniques, using G418 for selection. Cleared serum-free conditioned medium was desalted and thereafter, ADAMTS13 was purified with a Ni\textsuperscript{2+}-HiTrap Column (AmershamPharmacia, UK), according to manufacturer’s instructions. Purified fractions were dialysed into 20mM Tris-HCl (pH 7.8) and analysed by Western blotting using either anti-ADAMTS13 protease domain (a gift from Dr F Sheiflinger, Baxter\textsuperscript{18}) or anti-myc epitope monoclonal antibodies (Invitrogen, UK). Recombinant ADAMTS13 purity was assessed by non-reducing SDS-PAGE and Coomassie staining. ADAMTS13 concentration was determined with a BCA total protein assay kit (Perbio, UK).

Proteolysis of ADAMTS13 in Human Plasma

To 800µl of citrated normal human plasma, or plasma defibrinated with Reptilase Reagent (Pentapharm, Switzerland), recombinant ADAMTS13 (80nM) was added to increase the ADAMTS13 concentration. To three 100µl samples of ADAMTS13-spiked plasma, 8pM lipidated human tissue factor (TF) (Baxter, Austria) and phospholipids (PS:PC:PE – 20:60:20) (Avanti, USA) were added. Thereafter, 0.4mM fluorogenic thrombin substrate, Z-Gly-Gly-Arg-AMC-HCl (Bachem, UK) was added and samples recalcified to initiate thrombin generation. Thrombin generation was monitored in real-time using a Fluoroscan plate reader (Thermolabs, UK) and quantified by comparison to a standard curve generated at the same time. From parallel, samples lacking the thrombin substrate, sub-samples (25µl) were removed and analysed by
Western blotting using the anti-ADAMTS13 protease domain antibody. In some experiments, 5mg/ml GPRP-amide (Sigma) was added to whole plasma samples to prevent polymerisation and cross-linking of fibrin. 100U/ml hirudin was added to one sample to specifically inhibit thrombin.

**Enzymatic Proteolysis of ADAMTS13 by Thrombin, FXa and Plasmin**

250nM ADAMTS13 was incubated with 0.9-90nM human α-thrombin (Sigma, UK), FXa (Calbiochem, UK) or plasmin (Calbiochem, UK) in 20mM Tris-HCl (pH7.8), 5mM CaCl₂, 150mM NaCl at 37°C. At time points (0-6 hours), ADAMTS13 fragmentation was analysed by Coomassie staining and Western blotting (with anti-protease domain or anti-myc mAb). Experiments using heparin or TM were performed as above except that thrombin (50nM) was preincubated for 5 minutes with equimolar or a 10-fold molar excess of unfractionated heparin (CP Pharmaceuticals, UK) or soluble rabbit TM (Haematologic Technologies), respectively. To assess the relative specificity of thrombin for ADAMTS13 over other ADAMTS-family members, parallel experiments in which ADAMTS4 was treated with 9nM thrombin for 0-6 hours were analysed by Western blotting using ADAMTS4 protease domain-specific polyclonal antibodies (also a gift of Dr H. Nagase, Imperial College London).

**Purification of VWF from Human Plasma**

VWF was purified from normal human plasma as described by Bowen. Briefly, 600ml fresh frozen plasma was cryoprecipitated. The pellet was dissolved in 10ml 20mM Tris-HCl (pH 8). Samples were cleared before purification by gel filtration using a Sepharose CL-2B HiPrep 26/60 column (AmershamPharmacia, UK). VWF concentration in each fraction determined using a specific VWF ELISA, as previously described. The purity of VWF was assessed by
reducing SDS-PAGE and silver staining.

**ADAMTS13 Activity Assays**

To assess the effect of proteolysis upon ADAMTS13 activity, 1mM ADAMTS13 was incubated with and without 9nM thrombin or plasmin for 16 hours at 37°C, as outlined above. ADAMTS13 activity assays were performed essentially as described\(^\text{14}\) and using a VWF collagen-binding assay (VWF:CBA) (Technoclone, UK). Briefly, 5µg of each ADAMTS13 sample (with and without proteinase treatment) was preincubated with 10mM BaCl\(_2\) for 10 minutes at 37°C. Thereafter, samples were incubated at 37°C with 8nM purified human VWF in reaction conditions containing 1.5M urea, 10mM BaCl\(_2\), 5mM NaCl, 0.5mM CaCl\(_2\), 15mM Tris-HCl (pH 8.0). At 0, 1, 2, 4 and 6 hours, sub-samples were removed for analysis of VWF proteolysis. Samples were mixed with 2x reducing SDS loading buffer and heated for 20 minutes at 95°C. Samples were separated on 5% polyacrylamide gels and electroblotted onto a PVDF membrane. Proteolysis of VWF was examined by Western blotting using a polyclonal anti-VWF antibody (Dako, UK). Parallel samples were also analysed in triplicate with a VWF:CBA, according to manufacturer’s instructions. ADAMTS13 activity was thus assessed indirectly by examining its effect upon VWF collagen-binding function. VWF:CBA results were normalised by parallel sampling in the absence of recombinant ADAMTS13.
Results

Purification of Recombinant ADAMTS13

Recombinant myc/6xHis-tagged ADAMTS13 was expressed into conditioned medium from HEK293 cells that had been stably transfected with the vector, pcDNA3.1/ADAMTS13/myc/His. ADAMTS13 (Mr ~200kDa) was purified and concentrated by affinity chromatography, see result of SDS-PAGE (Fig. 1, lane 1). Western blot analysis using an anti-ADAMTS13 protease domain antibody highlighted the presence of the same prominent ~190-210kDa band (Fig. 1, lane 2), in accordance with a previous report\(^{18}\). However, analysis of this same sample using an anti-myc epitope antibody revealed the presence of an additional but lower intensity ~40kDa band (Fig. 1, lane 3). This minor species was believed to be a C-terminal ADAMTS13 degradation fragment (containing the myc/His tag) that was co-purified with the full-length enzyme.

ADAMTS13 is Proteolysed Following Activation of Coagulation in Human Plasma

Western blotting to detect endogenous ADAMTS13 in human plasma was ineffective due to its low concentration. Addition of recombinant ADAMTS13 to defibrinated plasma allowed detection of a single ~200kDa ADAMTS13 band. Adding relipidated TF, phospholipids and Ca\(^{2+}\) initiated thrombin generation that was quantified in real-time using a fluorogenic substrate and an automated plate reader. This demonstrated that thrombin was generated in ADAMTS13-spiked plasma samples with a characteristic lag-time followed by a rapid propagation phase (Fig. 2A). In parallel samples lacking the thrombin substrate, aliquots were removed at different time points (0-30 minutes), and the effect of coagulation upon ADAMTS13 was examined by Western blotting using the anti-ADAMTS13 protease domain monoclonal antibody. In samples in which thrombin was generated (Fig. 2B), a decrease in the intensity of the full-length
ADAMTS13 band was observed. This demonstrated that activation of the coagulation cascade caused the rapid loss of full-length ADAMTS13. In the absence of a procoagulant stimulus (i.e. in samples lacking TF, phospholipids and Ca\textsuperscript{2+}), no such depletion in full-length ADAMTS13 over the 30 minute time-course (not shown). To further assess this loss of ADAMTS13, whole human plasma (containing fibrinogen) was supplemented with ADAMTS13 and also 5mg/ml GPRP-amide, to prevent fibrin polymerisation and subsequent cross-linking. Analysis of these samples by Western blotting using the anti-protease domain mAb (Fig. 2C) revealed that without supplemented ADAMTS13, no ADAMTS13 was detected in plasma (Fig. 2C - lane 1). In the absence of procoagulant stimulus, recombinant ADAMTS13 was detected as a single ~200kDa band (Fig. 2C - lane 2). Following a 30 minute incubation at 37°C with TF, Ca\textsuperscript{2+} and phospholipids, the intensity of the full-length ADAMTS13 band was markedly reduced (Fig. 2C - lane 3), but this was completely inhibited by hirudin (Fig. 2C - lane 4). This implied that the loss of ADAMTS-13 from plasma was mediated in part via a mechanism involving thrombin.

**Thrombin Cleaves ADAMTS13 at Several Sites**

Degradation of ADAMTS13 by thrombin and/or FXa was considered a possible mechanism for the removal of full-length ADAMTS13 from plasma. To circumvent the technical difficulties associated with blotting/detecting specific proteins in plasma samples, this was examined further using purified proteins. ADAMTS13 was incubated with 9nM human thrombin for 16 hours in a reaction containing 150mM NaCl and 5mM Ca\textsuperscript{2+}. Reducing SDS-PAGE and Coomassie staining revealed that without the addition of thrombin, full-length ADAMTS13 was clearly visible as a band of ~200kDa (Fig. 3A). In the same sample containing thrombin, the full-length ADAMTS13 band disappeared completely (Fig. 3A). Furthermore, lower molecular weight bands that were not readily visualised in either the thrombin or ADAMTS13 preparations
alone, became evident (Fig. 3A – see newly emerged arrows). These smaller bands represent degradation products arising from thrombin cleavage. This finding not only demonstrated that ADAMTS13 is susceptible to proteolysis by thrombin, but also that cleavage occurs at multiple sites.

Under non-reducing conditions (Fig. 3B), after incubation with thrombin, a similar disappearance of the band corresponding to full-length ADAMTS13 was observed. Two major ADAMTS13 proteolytic fragments were resolved, one of ~150kDa and another of ~40kDa (Fig. 3B), the latter band migrating with a similar molecular weight to that of thrombin. The presence of only two major cleavage products under non-reducing conditions suggested that most of the ADAMTS13 cleavage fragments, with the exception of the 40kDa C-terminal species, remain covalently associated by disulfide bond linkages.

Western blot analysis of these same samples using the anti-ADAMTS13 protease domain antibody revealed that under reducing conditions, the protease domain was present within a ~28kDa band (Fig. 3C). Under non-reducing conditions and after treatment with thrombin, the protease domain was present in the ~150kDa band (Fig. 3C).

Analysis of these samples using the anti-myc antibody revealed that a ~40kDa C-terminal fragment of ADAMTS13 was generated after thrombin treatment under both reducing and non-reducing conditions (Fig. 3D). This implies that at the most C-terminal cleavage site, thrombin proteolyses ADAMTS13 between unpaired cysteine residues and that the C-terminal fragment (containing all or most of the two CUB domains) is no longer covalently associated with the rest of the ADAMTS13 molecule.

*Time- and Concentration-Dependent Cleavage of ADAMTS13 by Thrombin*

Incubation of ADAMTS13 with varying concentrations of human thrombin (0.9-90nM or
0.1-10 U/ml) for 0-6 hours followed by Western blotting with the anti ADAMTS13 protease domain antibody, revealed that ADAMTS13 was cleaved by thrombin in a time- and concentration-dependent fashion (Fig. 3E). In the presence of 0.9nM thrombin, no cleavage of the ~200kDa full-length ADAMTS13 band was observed after one hour incubation. At 2 hours, an additional faint ~150kDa band became evident and by 6 hours, this band and a band at ~28kDa were also observed.

Using 9nM thrombin, the rate of ADAMTS13 proteolysis was appreciably enhanced. After only 30 minutes, little full-length ADAMTS13 remained. Moreover, the three major fragmentation bands (~150kDa, ~40kDa & ~28kDa, respectively) were prominent. Over time, the high Mr band disappeared, whereas the ~28kDa band became stronger. With 90nM thrombin, there was complete loss of full-length ADAMTS13 within 30 minutes. Interestingly, the gain in the 28kDa band intensity did not necessarily reflect the loss in intensity of the higher Mr bands. One possible cause of this might have be the presence of an additional non-specific/less favoured cleavage site within the protease domain that with time, generated a fragment that was only marginally smaller than the 28kDa band.

Experiments were also performed using FXa. These demonstrated that FXa caused ADAMTS13 fragmentation similarly to that caused by thrombin. However, the rate of ADAMTS13 proteolysis by FXa was less than that of thrombin (Fig. 3F). Even at 50nM FXa, proteolysis of ADAMTS13 was markedly slower than when 9nM thrombin was used (Fig. 3E).

**Characterisation of ADAMTS13 Proteolysis by Thrombin**

To further analyse ADAMTS13 fragmentation, partially proteolysed ADAMTS13 species were visualised by Western blotting. As before, we detected either N-terminus-containing fragments with the anti-ADAMTS13 protease domain antibody (Fig. 4A) or C-terminus-
containing fragments with the anti-myc antibody (Fig. 4B). In these experiments, short incubations using limited quantities of thrombin were employed to facilitate visualisation of proteolytic intermediates of ADAMTS13 (Fig. 4A & 4B – lanes 2). In addition, increased quantities of ADAMTS13 were treated with thrombin and blots developed for extended periods to better identify transient proteolytic intermediates that, over time, would be further cleaved into smaller species (Fig. 4A, lane 3 & Fig. 4B, lane 4).

Limited treatment of ADAMTS13 with thrombin revealed at least three protease domain-containing fragments (Fig. 4A – lane 2) of ~150kDa, ~40kDa and ~28kDa (designated Frag 1, 5 & 6, respectively). Frag 1 and 6 were the major species. However, proteolysis using increased quantities of ADAMTS13 (Fig. 4A – lane 3) indicated that at least a further three intermediate species were generated by thrombin treatment (Frag 2, 3 & 4) – the faintness of these bands, visible on the original blot, is suggestive of additional, yet less favoured, thrombin cleavage sites. In Fig. 4C, the approximate locations of Frag 1-6 are predicted based on their mobilities and identification by the detecting antibodies.

Similar analysis of C-terminal, myc epitope-containing fragments (Fig. 4B) revealed a fragmentation pattern that corroborated and complemented the data obtained using the anti-ADAMTS13 protease domain antibody. The major species generated by thrombin were Frag A (~160kDa) and Frag D (~40kDa), again highlighting the potential preference for proteolysis at the most C-terminal and N-terminal sites. Interestingly, the ~40kDa C-terminal fragment that was co-purified with ADAMTS13 during its original preparation could not be resolved from Frag D. This might imply that both species have arisen via cleavage at the same proteolytically sensitive site.

Due to their predicted high molecular weight, we could not resolve the 170kDa, 160kDa
Crawley et al.: Regulation of ADAMTS13 by Thrombin

and 150kDa fragments that would correspond to the C-terminal aspects of Frag 4, 5 and 6 (Fig. 4C). These species probably co-migrated as Frag A. Once again, proteolysis of increased quantities of ADAMTS13 indicated two additional intermediate species were generated (Frag B & C), very possibly representing the C-terminal portion of Frag 2 and 3. Again, the location of these fragments is predicted in Fig. 4C.

*Proteolysis by Thrombin Ablates ADAMTS13 Activity*

C-terminal truncation of ADAMTS13 past the spacer domain has previously been reported to abolish ADAMTS13 activity\(^{23,24}\). It would seem that after proteolysis by thrombin, the protease domain to TSP1-8 remains covalently intact through disulphide bond linkages. Although a ~40kDa C-terminal fragment of ADAMTS13, containing the CUB domains, is proteolytically removed by thrombin, the loss of these domains does not appear to significantly affect the VWF-cleaving activity of ADAMTS13 in static assays\(^{23,24}\). Thus, to ascertain whether thrombin cleavage might simultaneously abrogate ADAMTS13 activity, we compared the ability of full-length and thrombin-proteolysed ADAMTS13 to cleave purified human VWF. Human VWF was purified from plasma by cryoprecipitation followed by gel filtration (Fig. 5A). The first protein fractions eluted (B8-C6) contained VWF. The purity of VWF was assessed by SDS-PAGE and silver staining (Fig. 5B). Fractions B8-B10 were free of major contaminating proteins and were thus used as an ADAMTS13 substrate. ADAMTS13 pre-incubated in the absence of thrombin was fully active and generated characteristic 176kDa and 140kDa VWF bands (Fig. 5C). Moreover, the intensity of the 250kDa VWF band diminished over the 6 hour time-course. ADAMTS13 that had been pre-treated with thrombin exhibited little or no VWF-cleaving activity. As the VWF Western blot detection method is qualitative, we also examined ADAMTS13 cleavage of VWF in a quantitative manner by monitoring loss of VWF function.
with a VWF:CBA (Fig. 5D). These data corroborated the VWF fragmentation results. At 6 hours, the VWF in samples containing ADAMTS13 preincubated without thrombin exhibited ~5% original collagen binding activity due to loss of VWF function arising from ADAMTS13-dependent proteolysis. In contrast, VWF incubated with thrombin-treated ADAMTS13 still exhibited ~90% of the original collagen binding potential. Taken together, these data confirm that once proteolysed by thrombin, ADAMTS13 is no longer able to efficiently cleave VWF and so inhibit its physiological function.

**Effect of Thrombin Co-Factors Upon ADAMTS13 Proteolysis**

To further assess the nature of the interaction of thrombin with ADAMTS13 and also to examine the way in which thrombin-dependent ADAMTS13 inactivation might itself be regulated/localised to the site of vascular injury, we investigated the effect of thrombin co-factors (TM and heparin) upon the rate of ADAMTS13 proteolysis. For these experiments, we preincubated 50nM thrombin for 5 minutes with either equimolar (50nM) or 10-fold molar excess (500nM) of either soluble rabbit TM or heparin. Thereafter, we examined the extent of ADAMTS13 fragmentation over a 1 hour incubation period.

**i) Thrombomodulin**

Preincubation of thrombin with soluble rabbit TM markedly inhibited the ability of thrombin to proteolyse ADAMTS13 (Fig. 6A). In the absence of TM, proteolysis went to near completion after 1 hour of incubation with thrombin. At this time, only trace amounts of full-length ADAMTS13 could be detected. However, with preincubation of thrombin with equimolar quantities of TM, the rate of full-length ADAMTS13 band depletion was reduced. The rate at which the smallest fragmentation band (Frag 6) was generated was also similarly impaired. As expected, the effect of a 10-fold molar excess of TM was more pronounced. Indeed, after 30
minutes incubation with thrombin, very little ADAMTS13 fragmentation was observed, by comparison to the no TM control. After 1 hour, we estimated that >50% of the remaining ADAMTS13 was still intact.

ii) Heparin

Preincubation of thrombin with equimolar or a 10-fold molar excess of heparin, had no inhibitory effect upon ADAMTS13 fragmentation, by comparison to samples lacking heparin (Fig. 6B). Indeed, at a 10-fold excess, the rate of ADAMTS13 cleavage may have been slightly enhanced, when compared to the no heparin and equimolar heparin samples. The reason for this very moderate enhancement is uncertain but may be due to bridging of ADAMTS13 and thrombin by heparin.

Specificity of Thrombin for ADAMTS13

To help establish whether thrombin proteolysed ADAMTS13 in a specific fashion, we compared the susceptibility of another ADAMTS-family member (ADAMTS4) to thrombin cleavage. In these experiments, no proteolysis of ADAMTS4 was detected with 9nM thrombin within 6 hours (Fig. 6C), suggesting that ADAMTS4 is resistant to thrombin-mediated proteolysis. This finding implies that ADAMTS13, which has a similar domain structure to ADAMTS4, is specifically recognised/cleaved by thrombin.

ADAMTS13 is Rapidly Proteolysed and Inactivated by Plasmin

Thrombin and FXa are generated rapidly at the site of vascular injury. However, other proteases are also locally activated following initiation of coagulation that might also influence ADAMTS13 function. We examined the susceptibility of ADAMTS13 to cleavage and inactivation by plasmin. Following limited proteolysis of ADAMTS13 (9nM plasmin for 30 minutes – Fig. 7A), plasmin generated a similar, but not identical, fragmentation of ADAMTS13
to that observed for proteolysis by thrombin (see Fig. 4A & 4B) – consistent with its limited specificity for cleavage after certain arginine residues\textsuperscript{25}. However, plasmin also cleaves after lysine, which potentially explains the presence of alternative ADAMTS13 fragments. Of particular interest was the additional cleavage site in the protease domain that generated Frag p7. This fragment was not generated by thrombin. Plasmin cleaved ADAMTS13 more rapidly than thrombin (compare Fig. 7B & Fig. 3, respectively). Indeed, at 0.9nM plasmin full-length ADAMTS13 was rapidly depleted/proteolysed. The effect of plasmin proteolysis upon ADAMTS13 activity was assessed using purified human VWF, as before. Pretreatment of ADAMTS13 with plasmin had the same effect upon VWF-cleaving activity as did thrombin. No proteolysis of VWF by plasmin-treated ADAMTS13 was detected by Western blotting (Fig. 7C). Furthermore, only a small reduction in VWF:CBA was observed in plasmin treated samples, whereas untreated ADAMTS13 generated characteristic VWF cleavage products and rapidly abolished the VWF collagen binding activity (Fig. 7D).
Discussion

In this study, we have addressed how ADAMTS13 might be regulated. To date, no specific protease inhibitor has been identified that might localise and restrict the known role of ADAMTS13 activity in prevention of unwanted VWF adherence and platelet aggregation to the intact endothelium. The absence of a specific inhibitor was considered surprising in view of the crucial role played by inhibitors in regulating proteases that participate in normal haemostasis. A single report has implicated interleukin-6 as a possible regulator of ADAMTS13, however, the mechanism for this and its significance in normal ADAMTS13 regulation remain uncertain. Accordingly, we have investigated proteolysis as a regulatory mechanism.

We have shown the rapid disappearance of full-length recombinant ADAMTS13 from defibrinated human plasma in which thrombin generation had been initiated (Fig. 2). One explanation for this was the proteolytic depletion of ADAMTS13 by a component of the coagulation cascade. Further evidence for this was provided using whole plasma, in which the depletion of full-length ADAMTS13 was inhibited by hirudin. Thereafter, we confirmed that in vitro low concentrations of thrombin cleaves purified ADAMTS13 (Fig. 3 & 4) (and not ADAMTS4, Fig. 6) at several sites. Interestingly, FXa cleaved ADAMTS13 with a very similar fragmentation pattern to that of thrombin. However, FXa proteolyzed ADAMTS13 appreciably slower. As thrombin is more active against ADAMTS13 and given that the local concentrations of thrombin that are generated in vivo invariably exceed those of FXa, the proteolysis of ADAMTS13 by thrombin is more likely to be of physiological importance.

It was apparent that the proteolysis of ADAMTS13 by thrombin occurs preferentially at two sites (Fig. 3). The first appears to be located close to the end of the protease domain and the second, near to the start of the first CUB domain. ADAMTS13 fragments generated by cleavage
at these sites were rapidly generated. With the exception of the most C-terminal site, thrombin
cleaved at sites located between paired cysteine residues. Thus after proteolysis, the resulting
ADAMTS13 cleavage fragments remain covalently associated, with the exception of the ~40kDa
C-terminal CUB domain-containing region (Fig. 3).

The physiological relevance of proteolytic removal of the CUB domains, located at the
C-terminus of ADAMTS13, is uncertain. These domains are not required for VWF-cleaving
activity measured under static conditions\textsuperscript{23,24}. Moreover, certain strains of mice have been
identified that express an ADAMTS13 variant that lacks the CUB domains (and also TSP1-7 and
-8)\textsuperscript{27}, further suggesting that the functions of these domains are not critical \textit{in vivo}. Although
auto-antibodies against the CUB domains have been found in patients with acquired TTP\textsuperscript{28}, it is
unclear whether these compromise ADAMTS13 function. The recent study reporting that CUB
domain-derived peptides inhibit VWF-cleaving activity under flowing, but not static, conditions
might however imply a potentially important functional role to these domains\textsuperscript{29}. This latter
finding suggests that removal of the CUB domains by thrombin might further contribute to
ADAMTS13 inactivation \textit{in vivo}. Further studies are now required to determine the precise
cleavage site and physiological effect of the proteolytic removal of the CUB domains.

ADAMTS13 is secreted into plasma, where it circulates as an active enzyme. At sites of
endothelial disruption, VWF binds collagen and unfolds in response to shear forces. This in turn
facilitates platelet tethering – critical for efficient platelet plug formation. If ADAMTS13 were to
have no specific inhibitor, its VWF-cleaving activity would be predicted to oppose the essential
tethering of platelets once the scissile bond in VWF is exposed by rheological forces. The data
presented here suggest an effective regulatory mechanism in which ADAMTS13 could be
proteolytically inactivated by thrombin at sites of vascular perturbation, even at the low thrombin
concentrations observed during the initial lag phase of coagulation. During initial platelet plug
formation, platelets adhere to VWF and then exposed collagen. However, although this first
platelet layer might resist ADAMTS13-mediated VWF cleavage/platelet release due to its tight
association with collagen, subsequent layers of platelets will be more remote from collagen. In vivo imaging of thrombus development in mice has demonstrated that thrombin activity/fibrin
deposition are generated very rapidly (within 20 seconds) after vascular injury. Interestingly,
this study also reveals the "blebbing-off" of tethered platelets from the back of the developing
thrombus prior to thrombin generation/fibrin deposition. This observation might be explained
by/correspond to the action of uninhibited ADAMTS13 upon VWF-bound platelets and highlight
a need for local ADAMTS13 inactivation at the site of injury.

Thrombin has multiple physiological substrates and roles. Following the initiation of
coeagulation, thrombin confers its prothrombotic activity by cleaving fibrinogen, feedback
activation of up-stream clotting factors and the activation of tethered platelets. Together, these
functions promote rapid thrombus formation. We now propose a model describing a novel
procoagulant function for thrombin. Uniquely, this involves the inactivation of an enzyme
(ADAMTS13) to promote thrombus growth, as opposed to its well characterised roles in
activation of PAR molecules and clotting factors V, VIII, X, XI and XIII. Recently, several
studies have attempted to implicate moderate ADAMTS13-deficiency in the pathogenesis of
certain thrombotic disorders. Our data suggests that reductions in ADAMTS13 activity might
arise as a result of increased thrombin-dependent inactivation during the course of a thrombotic
event. Therefore, additional studies are now required to ascertain whether partial deficiency of
ADAMTS13 is a primary cause, or merely a consequence, of thrombosis.

Physiologically, the activity and specificity of thrombin is modulated, in part, by co-
factors. Competition between co-factors for exosite binding have been proposed as a mechanism for directing the activities of thrombin\textsuperscript{37}. In binding TM, the specificity of thrombin towards fibrinogen, protein C and PAR molecules is dramatically altered\textsuperscript{38}. This serves to restrict its procoagulant activities to the site of vascular injury. This effect is partly due to the occupancy of exosite I on thrombin with TM\textsuperscript{39}. We found that if thrombin was preincubated with TM, its ability to proteolyse ADAMTS13 was appreciably compromised (Fig. 6). This might therefore implicate exosite I as a possible site of interaction with ADAMTS13. As preincubation of thrombin with heparin had no effect upon the rate of ADAMTS13 cleavage, it seems unlikely that exosite II\textsuperscript{40} on thrombin plays a significant role in its interaction with ADAMTS13.

Interestingly, there seemed to be a slight enhancement of ADAMTS13 proteolysis by thrombin in the presence of a 10 fold molar excess of heparin. This might be attributable to the bridging of thrombin and ADAMTS13 by heparin but further investigation is required to confirm this contention.

Whilst examining the susceptibility of ADAMTS13 to proteolysis by coagulation proteinases, we found that plasmin cleaved ADAMTS13 at a faster rate than thrombin, and gave a similar, but not identical fragmentation pattern. Physiologically, plasmin is generated following thrombin-dependent fibrin deposition and mediates fibrinolysis\textsuperscript{41}. Plasminogen activators bind to fibrin as it is deposited. Consequently, given the sensitivity of ADAMTS13 to proteolysis at low plasmin concentrations (Fig. 7), such cleavage might also be of physiological importance with respect to the modulation of ADAMTS13 activity at sites of haemostatic plug formation. Proteolysis of ADAMTS13 by plasmin might also suggest a role for VWF in tissue repair. Post-injury vessel regeneration is associated with plasmin activity. If this process should also involve a role for VWF, plasmin-dependent ADAMTS13 inactivation during tissue repair might serve to
maintain VWF function. It is interesting to note that whereas plasmin proteolytically activates several different prometalloproteinases\textsuperscript{42}, it uniquely has an inactivating role against ADAMTS13.

Together, our data provide a plausible mechanism by which the thrombin-dependent inactivation of ADAMTS13 might be specifically localised to the site of haemostatic plug formation. Whereas thrombin (and plasmin) might rapidly inactivate ADAMTS13 at the site of vascular injury (and so favour platelet aggregation), on the surface of the adjacent uninjured endothelium, any thrombin would preferentially bind to TM. Consequently, thrombin-dependent ADAMTS13 proteolysis would be inhibited. ADAMTS13 activity in this location would serve to limit platelet aggregation from spreading beyond the site of injury.
References


34. Bohm M, Antoine G, Plaimauer B, Krause M, Scheiflinger F: Determination of ADAMTS-13 activity in 70 patients with venous thrombosis with ischemic stroke revealed the first case of partial and congenital ADAMTS-13 deficiency as possible cause of ischemic


Acknowledgements

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Figure Legends

Fig. 1. Purification of recombinant human ADAMTS13. ADAMTS13 was expressed by HEK293 cells stably transfected with the expression vector, pcDNA3.1/ADAMTS13/myc/His. ADAMTS13 was purified from conditioned medium Ni$^{2+}$-HiTrap columns. Purified material was electrophoresed on a 7.5% polyacrylamide gel under non-reducing conditions and stained with Coomassie (lane 1). Lane 2 & 3 - Western blot analysis under reducing conditions with the anti-ADAMTS13 protease domain antibody and anti-myc epitope antibody, respectively. Lad = marker ladder.

Fig. 2. ADAMTS13 is proteolysed following thrombin generation in human plasma. Recombinant ADAMTS13 (final concentration 80nM) was added to defibrinated normal pooled citrated human plasma. A) Thrombin generation was initiated by the addition of 8pM TF, Ca$^{2+}$ and phospholipids (PL) (PS:PC:PE – 20:60:20). The amount of thrombin generated in defibrinated ADAMTS13-spiked plasma was monitored in triplicate in real-time by a Fluoroscan plate reader using a fluorogenic thrombin substrate. Readings were quantified with a standard curve generated in parallel. B) Parallel plasma samples were also analysed for ADAMTS13 fragmentation. At 0, 10, 20 and 30 minutes, ADAMTS13 degradation was assessed by Western blotting under reducing conditions using an anti-ADAMTS13 protease domain antibody. C) Citrated whole human plasma (containing 5mg/ml GPRP-amide) was incubated with either TF/PL/Ca$^{2+}$ (lane 1), 125nM ADAMTS13 (lane 2), TF/PL/Ca$^{2+}$ and 125nM ADAMTS13 (lane 3) or TF/PL/Ca$^{2+}$, 125nM ADAMTS13 and 100U/ml hirudin (lane 4) for 30 minutes at 37°C. Samples were analysed by Western blotting using an anti-ADAMTS13 protease domain antibody.
Fig. 3. Purified ADAMTS13 is cleaved by thrombin and FXa in a concentration- and time-dependent fashion. 400nM purified ADAMTS13 was incubated with and without 9nM thrombin (FIIa) in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\) at 37°C for 16 hours. Samples were electrophoresed on a 4-15% polyacrylamide gel under both reducing (A) and non-reducing (B) conditions followed by Coomassie staining. Thrombin was detected as a ~36kDa band (white arrowhead), full-length ADAMTS13 and its proteolytic fragments are identified by arrows. C) Western blot analysis of ADAMTS13 with and without complete thrombin proteolysis (under reducing and non-reducing conditions) using the anti-ADAMTS13 protease domain antibody. D) Western blot analysis of ADAMTS13 with and without complete thrombin proteolysis (under reducing and non-reducing conditions) using the anti-myc epitope antibody. E) Varying concentrations of thrombin (0-90nM) were incubated with 250nM ADAMTS13 at 37°C for 0 - 6 hours in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\). Samples were analysed under reducing conditions by Western blotting using the anti-ADAMTS13 protease domain antibody. F) 50nM FXa was incubated with 250nM ADAMTS13 at 37°C for 0 - 6 hours in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\) and analysed as in (E).
Fig. 4. Identification of ADAMTS13 fragments generated by thrombin. Recombinant ADAMTS13 (250nM) was incubated with 9nM human thrombin at 37°C for 30 minutes in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\), causing partial proteolysis of ADAMTS13. Samples were analysed by Western blotting. A) ADAMTS13 detected using the anti-ADAMTS13 protease domain antibody. ADAMTS13 (lane 1) was incubated with thrombin (lane 2). Increased quantities of ADAMTS13 (400nM) was incubated with thrombin and extended development of blots (lane 3). B) Probing of samples from (A) using the anti-myc antibody. ADAMTS13 (lane 1), ADAMTS13 treated with thrombin (lanes 2, 3 & 4). Lanes 3 and 4 represent same sample, lane 3 contains 25% of the volume loaded in lane 4. C) The predicted identities of the cleavage fragments in panel A (Frag 1-6) and panel B (Frag A-D) are represented. The locations of the predicted cleavage sites are approximate. Fragment boundaries/sizes were estimated from the molecular weights calculated from (A) and (B). The fragments generated by cleavage at the favoured cleavage sites are shown in bold.
Crawley et al.: Regulation of ADAMTS13 by Thrombin

Fig. 5. Proteolysis of ADAMTS13 by thrombin abolishes ADAMTS13 enzymatic activity towards purified human VWF. Human VWF was purified from human plasma by gel filtration. After gel filtration (A), the purity of VWF in each fraction was assessed by reducing SDS-PAGE on a 7.5% polyacrylamide gel, followed by silver staining (B). Fractions B8-B12 contained VWF, visualised as a ~250kDa band (arrow) and were free of major contaminating proteins. Only the highest purity VWF from fractions B8-B10 were used as a substrate for ADAMTS13. ADAMTS13 that had been pre-treated with and without 9nM thrombin in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\) for 16 hours at 37˚C was assayed for activity. Sub-samples were removed at 0, 1, 2 and 6 hours and assessed by Western blotting using an anti-VWF polyclonal antibody (C). ADAMTS13 generated characteristic VWF proteolytic fragments of 176kDa and 140kDa. Samples studied in (C) were also analysed using a functional assay based on collagen binding (VWF:CBA) as outlined in Experimental Procedures (D). Results shown ±SEM, (n = 3).
Fig. 6. Co-factors of thrombin and ADAMTS13 proteolysis. Recombinant ADAMTS13 (250nM) was treated with 50nM thrombin that had been preincubated at 37°C with either equimolar or a 10-fold molar excess of either soluble rabbit TM (A) or heparin (B) in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca^{2+}. ADAMTS13 fragmentation was monitored under reducing conditions using the anti-protease domain antibody. A) Thrombin was preincubated with either 50nM or 500nM soluble rabbit TM for 5 minutes. Thereafter, the ability of each thrombin•TM mix to cleave ADAMTS13 was monitored at 0, 30 minutes and 1 hour time points and was compared to that of thrombin alone. B) Thrombin was preincubated with either 50nM or 500nM heparin and analysed as in (A). C) The specificity of thrombin for ADAMTS13 relative to other ADAMTS-family members was gauged by incubating 250nM ADAMTS4 in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca^{2+} with 9nM thrombin for 0 – 6 hours. Samples were analysed by Western blotting using a polyclonal anti-ADAMTS4 protease domain antibody.
Fig. 7 Proteolytic inactivation of ADAMTS13 by plasmin. Recombinant ADAMTS13 (250nM) was incubated with 9nM human plasmin in 20mM Tris-HCl (pH 7.8) containing 150mM NaCl and 5mM Ca\(^{2+}\) at 37°C for 30 minutes, causing partial proteolysis of ADAMTS13 (A). Samples were analysed by Western blotting using either anti-ADAMTS13 protease domain (lane 1) or anti-myc epitope antibodies (lane 2). The rate of plasmin-dependent ADAMTS13 proteolysis was assessed by incubating varying concentrations of plasmin (0-90nM) with 250nM ADAMTS13 at 37°C for 30 minutes to 6 hours (B). Samples were analysed under reducing conditions by Western blotting using the anti-ADAMTS13 protease domain antibody. To assess the effect of plasmin proteolysis upon ADAMTS13 function, ADAMTS13 was pre-treated with or without 9nM plasmin for 16 hours at 37°C and thereafter assayed for activity by two methods, as outlined in Experimental Procedures. Sub-samples were removed from each assay at 0, 1, 2, 4 and 6 hours and assessed by Western blotting using an anti-VWF polyclonal antibody (C). ADAMTS13 generated characteristic VWF proteolytic fragments of 176kDa and 140kDa that were visualised after 1 hour and increased in intensity over the 6 hour time-course. There was also a concomitant decrease in the intensity of the 250kDa VWF band. Analysis of VWF function as in Fig 6D using a VWF:CBA (D). Results shown ±SEM, (n = 3).
Crawley et al.: Regulation of ADAMTS13 by Thrombin

Fig 1

![Image of gel electrophoresis](Image)

Lane: 1 2 3 Lad

- 250kDa
- 45kDa

Purified ADAMTS13

Fig. 2

A) Thrombin (nM)

B) +TF/+PL/+Ca²⁺

0 10 20 30 40 50 60

Time (mins)

C) Lane:

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<td>-</td>
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ADAMTS13
Fig. 4

A) 

Lad 1 2 3

ADAMTS13

Frag 1 ~150kDa
Frag 2 ~120kDa
Frag 3 ~95kDa
Frag 4 ~50kDa
Frag 5 ~40kDa
Frag 6 ~28kDa

B) 

Lad 1 2 3 4

ADAMTS13

Frag A ~160kDa
Frag B ~100kDa
Frag C ~80kDa
Frag D ~40kDa

C) 

Disintegrin-like

Metalloprotease Domain
Cy-rich & Spacer
TSPI Domains
CUB

ADAMTS13

Frag 1
Frag 2
Frag 3
Frag 4
Frag 5
Frag 6

ADAMTS13

Frag D
Frag C
Frag B
Frag A
Crawley et al.: Regulation of ADAMTS13 by Thrombin

Fig. 5

A) 

B) 

C) 

ADAMTS13

No Thrombin

37°C 16 hours

0 hrs 1 hr 2 hrs 6 hrs 0 hrs 1 hr 2 hrs 6 hrs

VWF

170kDa

140kDa

ADAMTS13

90nM Thrombin

37°C 16 hours

0 hrs 1 hr 2 hrs 6 hrs Lad

170kDa

140kDa

D) 

% VWF / Fibrin

0 20 40 60 80 100

0 1 2 3 4 5 6

Time (hours)
Fig. 6

A) TM:Thrombin

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B) Heparin:Thrombin

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C) ADAMTS4/Thrombin

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Proteolytic Inactivation of ADAMTS13 by Thrombin and Plasmin

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