VWF excess and ADAMTS13 deficiency: a unifying pathomechanism linking inflammation to thrombosis in DIC, malaria, and TTP

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Summary
Absent or severely diminished activity of ADAMTS13 (A Disintegrin And Metalloprotease with a Thrombospondin type 1 motif, member 13) resulting in the intravascular persistence and accumulation of highly thrombogenic ultra large von Willebrand factor (UL-VWF) multimers is the pathophysiological mechanism underlying thrombotic thrombocytopenic purpura. Reduced VWF-cleaving protease levels, however, are not uniquely restricted to primary thrombotic microangiopathy (TMA), e.g. thrombotic thrombocytopenic purpura, but also occur in other life-threatening thrombotic conditions: severely decreased ADAMTS13 activity is seen in severe sepsis, disseminated intravascular coagulation (DIC) and complicated malarial infec-tion. The clinical relevance of these secondary thrombotic microangiopathies is increasingly recognised, but its therapeutic implications have not yet been determined. The presence of a secondary TMA in certain diseases may define patient groups which possibly could benefit from ADAMTS13 replacement or a VWF-targeting therapy. This short-review focuses on the role of UL-VWF multimers in secondary TMA and discusses the potential of investigational therapies as candidates for the treatment of TTP. In conclusion, prospective clinical trials on the effectiveness of protease replacement in vivo seem reasonable. Carefully selected patients with secondary TMA may benefit from therapies primarily intended for the use in patients with TTP.

Keywords
ADAMTS13, VWF, TTP, DIC, sepsis, malaria

Background
Von Willebrand factor (VWF) is a multifunctional acute-phase glycoprotein with a central role in primary haemostasis (1, 2). Assembled as multimers in endothelial Weibel Palade bodies, VWF is exocytosed in response to various endogenous stimuli such as inflammatory cytokines, histamine, thrombin, fibrin, or to exogenous desmopressin, tethering passing platelets to damaged endothelial sites in the presence of high shear rates (3–7).

Newly secreted ultra large VWF multimers (UL-VWF) have a high spontaneous platelet-binding potential and consequently must be enzymatically degraded into smaller, less thrombogenic units before entering the circulation. A specific metalloproteinase, ADAMTS13, cleaves VWF after binding to the VWF-A2 domain (8). Fluid shear stress is mandatory to stretch and unfold these multimers into scissile strings in order to facilitate enzymatic processing (9, 10). This cleavage process, which is accelerated by platelets and FVIII (11), prevents elongation and accumulation of hyper-reactive UL-VWF strands on activated endothelial surface (12).

The pathophysiology of thrombotic thrombocytopenic purpura, the prototypical TMA, is characterised by an abnormal persistence of UL-VWF multimers caused by insufficient or absent ADAMTS13 activity, which can be acquired through protease-directed autoantibodies in idiopathic TTP (13) or inherited due to homogeneous ADAMTS13 gene mutations in congenital TTP (14).

The giant strand-shaped multimers with a weight up to > 1 GDa and a length up to > 1 mm (10, 15, 16) spontaneously bind, activate and aggregate platelets and thereby provide the basic molecular substrate for thrombus formation in small blood vessels (16, 17). A widespread deposition of platelet-rich clots throughout the body may occur with occlusion of microvessels and subsequent organ failure. Erythrocytes passing through UL-VWF/platelet/fibrin meshwork in the microvasculature can become fragmented, leading to mechanical haemolysis with schistocyte formation, a diagnostic hallmark of thrombotic microangiopathies (18). Micrsvessel thrombosis in TMA predominantly affects organs with low or absent CD36-expression, especially the kidneys (19). This may suggest some (as yet unproven) restriction of the cleavage activity of ADAMTS13 to CD36-expressing cells.

In addition to platelet-activating properties, VWF promotes leucocyte adhesion to endothelial cells (20) and VWF multimers provide large surfaces for the activation of the complement cascade, thereby directly linking haemostasis to inflammation (21). Not only in haemolytic-uraemic syndrome, but also in cases of TTP, complement-mediated inflammation may play a substantial role in disease pathogenesis (22–24). It has been hypothesised (21)
that in those TTP patients with only slightly reduced enzyme levels an underlying heterozygous ADAMTS13 gene defect may be turned into clinically overt disease by a simultaneous defect in factor H, a circulating glycoprotein regulating the alternative complement pathway. The coincidence of insufficiently degraded multimers with aberrant complement regulation results in uncontrolled synthesis of anaphylatoxins and membrane attack complex, damaging host tissue and promoting thrombosis. This proposed mechanism could explain cases of TTP or TTP-like disorders refractory to extensive plasma exchange-based treatment and may provide a new therapeutic target.

Acquired TTP is a devastating and highly fatal condition if not treated immediately. The primary therapeutic intent is to replenish ADAMTS13 activity by urgent plasma exchange or, if this is not immediately available, by plasma transfusion. Plasma exchange therapy has dramatically decreased overall mortality in acquired TTP from nearly 90% to ≤25% (25, 26). Urgent replacement of ADAMTS13 facilitates cleavage of UL-VWF multimers into smaller, less thrombogenic forms and interrupts the vicious cycle of platelet entrapment and thrombus formation. Replacement therapy using recombinant ADAMTS13 is a reasonable therapeutic approach for TTP since it would specifically replenish the lacking enzyme (27–29). This is a promising approach for TTP due to congenital deficiency of ADAMTS13, but its usefulness in acquired forms of TTP is questionable, since even high-dose replacement may not be able to surmount the endogenous ADAMTS13 inhibitors present in those patients (30). One option to overcome this would be the use of recombinant enzyme variants with increased resistance to inhibition by autoantibodies, which have recently been tested in vitro with promising results (31). Alternatively, one could use anti-VWF agents specifically targeting the VWF-A1 domain to interrupt the process of binding and activating platelets. Two anti-VWF aptamers, ARC1779 (32–40) and ARC15105 (41), have been extensively investigated so far, although their clinical development is currently on hold (30). ALX-0081 (INN: caplacizumab, Ablynx) is a selective and highly potent bivalent humanized nanobody specifically targeting the GPIb-binding site of VWF (42, 43). Recent results from a phase II trial indicate that caplacizumab is effective in patients with acquired TTP (44).

However, the problem of excessive UL-VWF is not exclusively restricted to primary TMAs, but also seems to be a major determinant of prognosis in other life-threatening inflammatory disorders (45–52). Reduced ADAMTS13 activity strongly correlates with the severity of coagulopathy and in-hospital mortality in patients with disseminated intravascular coagulation (DIC) (53) and is further associated with a poor outcome in septic organ failure (54). Thrombotic microangiopathy can complicate sepsis and may even persist after resolution of DIC, prolonging organ dysfunction and influencing prognosis (55).

Although the pathophysiology of ADAMTS13 deficiency in sepsis differs from that in TTP, in a subset of patients with severe sepsis, clinical and laboratory features closely resemble those seen in TTP (50), suggesting the potential for therapeutic utility of enzyme replacement or VWF-A1 targeting therapeutic strategies in patients with severely depressed ADAMTS13 activity unrelated to primary TTP.

This review focuses on the role of UL-VWF multimers in severe sepsis and other secondarily acquired thrombotic microangiopathies and discusses the potentials of ADAMTS13 replacement and VWF-targeting agents as future therapeutics.

**Thrombotic microangiopathies in systemic inflammation**

Often TTP follows an acute episode of inflammation, which possibly triggers autoantibody formation (56, 57).

The unifying pathology linking both systemic inflammation and primary TMAs is acute dysfunctional endothelial cell activation, indicated by high VWF antigen (VWFAg) and VWF propeptide (VWFpp) in both sepsis and TTP (58, 59).

As an acute phase reactant VWF is up-regulated and secreted in inflammatory conditions (30, 56, 60). Its plasma levels usually correlate with the extent of systemic inflammation. Its simultaneously released VWFpp is considered a suitable marker of endothelial cell activation since it is not consumed in (patho-)physiologic haemostatic processes.

An inflammation-related imbalance between overwhelming VWF-release and inhibited and/or exhausted cleaving capacity of ADAMTS13 may finally result in a substantial accumulation of highly adhesive UL-VWF strings compromising the microcirculation. Accordingly, the inverse relation between high VWF antigen levels and low ADAMTS13 activity is associated with the extent of inflammation and the severity of organ failure (61). This is in line with the finding that the enzymatic processing of VWF multimers after endothelial release is predictive for mortality in systemic inflammatory conditions (62).

In a large cohort of a heterogeneous patient population with systemic inflammatory response syndrome (SIRS), VWF levels were significantly higher in non-survivors with an independent (APACHE Score-adjusted) two-fold increased hazard of death for those in the highest VWF-level-tertile (62). In contrast, patients with the highest ADAMTS13 levels showed a four-fold lower mortality rate compared to those with lowest ADAMTS13 levels (hazard ratio HR=0.22; confidence interval CI 0.07–0.74). This is supported by another study which found that VWF levels continuously increase with growing severity of inflammation while ADAMTS13 levels simultaneously decline, culminating in the accumulation of uncleaved VWF molecules (63). Mounting VWF/protease-imbalance therefore was proportional to the risk of developing TMA and consequently organ failure. Accordingly, the ratio between high VWFAg/VWFpp and low ADAMTS13 levels (median 25 ± 8.5%) may be a sensitive marker of disease severity and a useful tool to stratify patients at risk of developing inflammatory TMA (63).

Similar findings were observed in patients with severe sepsis, both in adults (54) and infants (52, 64). Consistently, reduced ADAMTS13 activity determined disease severity (52, 65), organ failure and mortality (64, 65). A causal relationship to sepsis is
indicated by the observation that ADAMTS13 levels discriminated between differences in inflammation, endothelial dysfunction, renal failure, haemodynamic compromise and mortality only in patients with severe sepsis, but not in those with organ failure unrelated to sepsis (54). Severely septic patients with ADAMTS13 levels below 43% showed a significantly lower survival rate, independent of the presence of DIC.

However, two studies, a septic mouse model (66) and a clinical study in septic adults (67) yielded different results; although VWF parameters inversely correlated with ADAMTS13 activity there was no relation between either of these parameters and disease severity or outcome. This apparently negative finding reported by Kremer Hovinga et al. may be explained by fact that their patients’ median ADAMTS13 level was 60% of normal, which is rather high compared to the levels found in other studies yielding positive results, and thus is likely to have been well above a critical threshold value. In contrast, in the study reported by Ono et al. (50) showing a positive correlation between ADAMTS13 level and clinical outcome ADAMTS13 levels were markedly lower, possibly due to a genetic polymorphism in the ADAMTS13 gene which is frequently found in Japan but which has not been reported in Caucasians. Another potential cause for the discrepancy may be the presence or absence of DIC. While Ono et al. (50), who found a clear inverse association between outcome and median ADAMTS13 activity (43%), only included septic patients with DIC (calculated by the LMHW-DIC score) Kremer Hovinga et al. (67) provide no information on DIC prevalence in their septic study population (20% with septic shock, median ADAMTS13 activity 60%). Supporting a possible role of DIC, Habe and colleagues (68) investigated 69 patients with DIC (36% with septic DIC; median ADAMTS13 activity 35%) and found an inverse correlation between the ISTH-DIC score and ADAMTS13 activity with a trend of lower enzyme activity in non-survivors (25 vs 38%). However, the patient population was very heterogeneous including only 36% with septic DIC, but 40% with malignancy and 14% with various conditions. Finally, similarly to Ono et al., Martin et al. (54) described a significant association between outcome and ADAMTS13 activity (median 43%); however, this finding was independent of disease severity (APACHE II score) and the presence of DIC (assessed by ISTH DIC score). Albeit the application of different DIC scoring systems (JMHW vs ISTH DIC score) and the inclusion of patients with different underlying pathologies causing DIC makes it difficult to compare these studies.

Several molecular mediators of endothelial activation with consequent VWF liberation have been investigated so far. The insufficient cleavage capacity of ADAMTS13 likely is also a multifactorial event and may be caused by exhaustion, direct inhibition, reduced synthesis, proteolytic cleavage and enhanced clearance (66, 69–71) (►Table 1).

- Exhaustion. A consumptive ADAMTS13 deficiency occurs when inflammation-triggered substrate supply exceeds the enzyme’s proteolytic capacity. The reciprocal correlation between VWF and ADAMTS13 was observed in several in vitro and in vivo studies ranging from an experimental endotoxinemia model (72) to prospective clinical trials. Lipopolysaccharide exposure results in a significant reduction of ADAMTS13–activity, hepatic ADAMTS13-mRNA expression, and platelet counts, whereas both VWF antigen and neutrophil counts increase. The opposing changes in VWF and ADAMTS13 blood levels also correspond to respective changes in hepatic mRNA expression patterns (73). Finally, in a septic mouse model, ADAMTS13 activity was mainly determined by VWF levels and preserved in VWF-gene-deleted septic animals (66). However, even without accompanying inflammation, stimulating secretion of VWF by administration of Desmopressin decreases ADAMTS13 levels (69, 74).

- Inhibition. In vitro experiments on platelet-rich plasma showed that inflammatory cytokines released from activated leukocytes and endothelial cells at an early stage of inflammation stimulate the synthesis of UL-VWF multimers–strings (both interleukin [IL]-8 and tumour necrosis factor [TNF] α) and concurrently inhibit their cleavage (IL-6 alone) (5). This corresponds to the finding that both reduced ADAMTS13 activity (54) and high IL-6 predict poor prognosis in TTP (5) and septic shock (75, 76). Finally, DIC is associated with the consumption of circulating clotting factors, including FVIII, which normally accelerates VWF proteolysis by ADAMTS13 (11).

- Proteolytic cleavage. A moderate correlation between ADAMTS13 levels and granulocyte elastase-mediated fibrin degradation products in septic adults (50) suggests a contributory role of proteolytic cleavage by neutrophils to diminished ADAMTS13 activity in inflammation. This would be in line with the observation that neutrophil elastase levels are significantly elevated not only in sepsis but also in acute TTP, where both neutrophil extracellular traps (77) and the plasma elastase concentration (71) were found to contribute to disease severity. Accordingly, elastase inhibitors abolished the ADAMTS13 decrease after LPS challenge in mice (73).

- Thrombin, plasmin and reactive oxygen species, all commonly elevated in sepsis and SIRS, as well as specific microbial proteases may also impair the catalytic activity of ADAMTS13 (70, 78, 79). The latter has been shown for Bacillus anthracis (80) and conceivably might apply to other bacteria more frequently causing bloodstream infections.

- Competition. Thrombospondin-1 (TSP-1) is a matrix-glycoprotein which is elevated in inflammatory conditions including human sepsis and which is considered an early marker of sepsis-related DIC (81). It regulates the inflammatory response by activating leukocytes and enhancing chemotaxis and was linked to mortality in a murine sepsis model (82). Thrombospondin-1 is known to be essentially involved in haemostatic processes but its role regarding the development of secondary TMA has not been evaluated yet. By competing with ADAMTS13 for the interaction with the VWF-A3 domain (83), thrombospondin-1 protects newly secreted haemostatically reactive UL-VWF multimers against degradation and slows the rate of their proteolysis (84, 85). Whether increased thrombospondin-1 levels are of clinical relevance regarding the emergence of TMAs in sepsis and SIRS is unclear.
However, patients with sepsis represent a heterogeneous population who suffer from various comorbidities. Host characteristics and risk factors finally determining the development of secondary TMA in inflammation are not well-defined. Coagulopathy in sepsis is a complex process characterized not only by tissue factor related thrombin generation but also by an impaired fibrinolysis, referred to as “fibrinolytic shut down” (86, 87). This may also play a role regarding the occurrence of UL-VWF strings, since plasminogen activation on endothelial cells acts as a natural backup for ADAMTS13 to degrade obstructive platelet-VWF complexes (88). Septic patients lacking both pathways of multimer degradation (ADAMTS13 and fibrinolytic enzymes) may be at high risk of additionally developing TMA. The relative contribution of fibrinolytic enzymes to VWF cleavage, however, is questionable, given that patients suffering from congenital TTP are a priori not expected to have a defect in fibrinolysis.

Also, the contributing role of complicating DIC regarding the development of TMA in sepsis is uncertain. One recent study specifically addressed this issue and found that the occurrence of ADAMTS13 deficiency (median 30%; interquartile range [IQR] 9–45) and its predictive value for mortality in severe sepsis are independent of the presence of DIC (76). Indeed, TMA associated organ failure can outlast DIC and perpetuate microvascular compromise despite the resolution of plasmatic coagulopathy (55).

### Thrombotic microangiopathy in DIC

The scientific subcommittee on DIC of the International Society on Thrombosis and Haemostasis (ISTH) defines DIC as an acquired syndrome characterised by a systemic intravascular activation of coagulation with loss of localisation, which causes damage to the microvasculature, and if sufficiently severe, can produce organ dysfunction (89). DIC is the worst manifestation of a globally activated and deteriorated coagulation system caused by heterogeneous disorders of which sepsis may be one of the most common. It is not a primary disorder but can be superimposed upon and can complicate various medical conditions with high morbidity and mortality.

In contrast to TMAs, where there is little activation of the coagulation cascade, tissue factor-driven coagulopathy in DIC results in excessive thrombin generation with activation and subsequent exhaustion of all coagulation components and platelets, which finally culminates in disseminated depositions of fibrin-rich (rather

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<th>Mode of action</th>
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than platelet-rich thrombi, organ damage and bleeding. Table 2 shows laboratory features of TTP, DIC and coagulopathy in malaria.

A recently published study on patients with various thrombocytopenic disorders found low ADAMTS13 levels to occur in a substantial subset of these patients. Low protease levels were significantly associated with high VWF Ag levels and low platelet counts (51). The most pronounced decrease in VWF proteolytic activity was found in individuals with severe thrombocytopenia. The abnormalities in VWF proteolysis were indistinguishable from that of TTP patients compared to patients with other thrombocytopenic disorders (51).

Another trial investigating patients with sepsis-related DIC yielded similar results: 16% of DIC patients had severely reduced ADAMTS13 activity levels (<5%) with a clinical and chemical profile indistinguishable from that of patients with primary TMA. DIC patients with severely reduced protease activity fulfilled all clinical and laboratory criteria for TTP, indicated by marked non-immune haemolysis (mean LDH: 2481 U/l, mean haemoglobin: 8.3 g/dl), severe thrombocytopenia (mean platelet count 7×10^9/l), impaired renal function (mean creatinine: 1.9 mg/dl) and altered consciousness in 47% (50). The incidence of renal failure, but not that of other organs, was significantly higher in those patients with ADAMTS13 activity <20%, with a strong correlation between UL-VWF multimers and creatinine levels (50). Furthermore, other studies confirmed a higher incidence of renal failure in septic patients with deficient ADAMTS13 (90) and/or high VWFpp levels (67, 90), irrespective of the presence or absence of DIC (54). The higher incidence of kidney damage in septic patients with low enzyme activity might be explained by the suspected, but not yet proven, restriction of ADAMTS13 on CD36-expressing cells, as mentioned above.

Considering that survival in septic shock is substantially lower in those patients with early renal failure (91), therapeutic interventions targeting UL-VWF multimers may be expected to reduce mortality.

Patients with DIC have significantly higher VWFAg and VWFpp levels and lower ADAMTS13 activity than those with related underlying diseases but without DIC. Patients with VWFpp levels >500 U/dl have a significantly lower survival rate (39 vs 61%; p<0.01) (68). The negative impact of high VWFpp on survival is also characteristic for primary TMA. Furthermore, in DIC a low activity of ADAMTS13 strongly correlates with the severity of coagulopathy and in-hospital mortality in adults and is a predictor of poor survival (53).

### Thrombotic microangiopathy in severe falciparum malaria

Microvessel obstruction by infected red cells is a crucial event in malarial coagulopathy, the extent of which is proportional to the disease severity (92). Mechanisms of microvascular compromise in malaria are complex and have been the focus of recent investigations (92–96). As in sepsis and SIRS, disease-related early endothelial cell activation with stimulated Weibel Palade exocytosis is considered a hallmark of malarial disease (94).

Parasite-induced endothelial tissue factor expression, activation of the complement cascade, impairment of coagulation inhibitors and dysfunctional fibrinolysis are contributing and intertwining mechanisms of coagulopathy in severe malaria, finally leading to DIC in 33–50% (92).

However, a common feature of patients infected with plasmodium falciparum is a drop in platelet count (60–80%) (97), which occurs already in the earliest phase of blood stage infection (even before clinical symptoms appear) and becomes progressively more severe in later stages (98). Early thrombocytopenia is not due to an incipient consumptive coagulopathy (99) but rather to the first obvious sign of incipient

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Table 2: Laboratory features of TTP, DIC and coagulopathy in malaria.

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* Median platelet count was 16×10^9/l (range:5–120) (118) and 25.5 ×10^9/l (range:2–124) (119), respectively.

~ Microangiopathic haemolytic anaemia. # Prevalence of thrombocytopenia in malaria ranges from 60%–80% (97). Degree of coagulation derangement varies from mild to severe and correlates with parasitaemia and disease severity (120). In a study on patients (18–40 years) with acute, uncomplicated plasmodium falciparum malaria the mean serum lactate dehydrogenase (LDH) activity was 789 ± 35 IU in males (n=76) and 634 ± 35 IU in females (n=76), considered to result from both red cell lysis and hepatic injury (121). § Anaemia is common in malarial infection, typically presenting with normal erythrocyte indices and a low reticulocyte count (122). Its pathogenesis is multifactorial including intravascular destruction of parasitized red blood cells (RBCs), extravascular clearance of erythrocytes and bone marrow suppression with dyserythropoiesis (123). Black water fever is a very rare form of anaemia and typically associated with a preceding use of quinine and the presence of glucose-6-phosphatase dehydrogenase (G6PD) deficiency. No routine laboratory finding is, however, highly specific for malaria. Platelets <150×10^9/l along with fever, enlarged spleen and serum bilirubin level >1.3 mg/dl was most predictive of malaria in a prospective study on 1900 European febrile travellers (124). In a retrospective analysis the cumulative presence of fever, thrombocytopenia and splenomegaly had a sensitivity of 71% and specificity of 88% (125). DIC occurs in up to 30% of previously naive patients with complicated falciparum malaria (92). Elevated plasminogen activator inhibitor-1 (PAI-1) and reduced tissue plasminogen activator (t-PA) levels indicate a thrombotic phenotype of DIC. ** Fragmented red blood cells (schistocytes and helmet cells). A schistocyte count >1% is highly suggestive of TTP-HUS (18).
VWF-mediated microvessel occlusion. Acutely exocytosed UL-VWF multimers are a critical component linking plasmodium-infected erythrocytes to the endothelial surface and platelets, thereby providing the basis for microthrombus formation with early platelet consumption and ensuing microangiopathic organ damage (Figure 1). This would be in line with the observation that patients with blood group 0, who constitutively express significantly lower VWF levels, are relatively resistant to develop severe malaria (100).

However, since infected red blood cells (RBCs) cannot directly bind to the vascular surface without appropriate receptors, the exact mechanism underlying cytoadherence remained unclear until recently. Platelets have already been recognised as bridging structures between activated brain endothelial cells lacking CD36 and parasited erythrocytes. However, Bridges and colleagues have demonstrated the critical role of VWF to accomplish red cell adherence from results of a laminar flow model. Accordingly, endothelial adhesion and sequestration of infected RBC is mediated by UL-VWF strings, which are littered with platelets abounding in CD36 molecules (93, 101, 102). Platelet microparticles transferring antigens to the RBC surface also increase cytoadherence, at least to cerebral endothelial cells, and may have a triggering role (103).

This proposed microangiopathic mechanism of VWF-mediated microvascular disease in malaria resembling TTP is emphasised by studies showing that both VWFAg and VWFpp levels are markedly elevated early (94, 96), especially in severe and cerebral malaria, indicating massive endothelial perturbation. Both continuously increase with ongoing malarial infection – while platelets proportionally decrease (98) – and strongly correlate with disease severity (94, 104). Due to the short half-life of VWFpp, its levels rapidly fall with disease recovery (94), indicating the acuity of endothelial activation in malaria. These observations could also be reproduced in an experimental human malarial infection model in healthy volunteers, where VWF levels started to rise immediately after plasmodium inoculation with activated VWF reaching levels usually only found in TTP (98). The significant correlation between VWF and low platelet count was even stronger with the activated, GPIba-binding-domain expressing form of VWF, suggesting the latter to be the important mediator of thrombocytopenia during early malaria (98).

Studies investigating ADAMTS13 activity in both uncomplicated and complicated malaria further highlight the parallels between TTP and malarial microvascular disease (96). As observed in severe sepsis and DIC, the presence of procoagulant UL-VWF multimers in malaria is accompanied by a persistently reduced ADAMTS13 level and strongly correlates with increased VWF collagen binding activity, which indicates the multimers’ hyper-reactive phenotype (95). Using recombinant ADAMTS13, malaria-associated UL-VWF strings that capture RBCs could rapidly be cleaved from the endothelium in vitro, a finding which emphasizes a disease-contributing role of enzyme deficiency in malaria. Underlying mechanisms are likely similar to those suspected in other conditions with secondary TMA. In addition to consumptive deficiency due to high substrate level, inhibition by inflammatory cytokines, acquired autoantibodies and non-immune mechanisms (microbial proteases, thrombin, plasmin, haemoglobin) including genetic polymorphisms (105) may play a role (Table 1).

To date, four clinical studies specifically investigated ADAMTS13 activity in the context of malaria and demonstrated that severe and cerebral, but not uncomplicated malaria, are
associated with ADAMTS13 deficiency. However no study yet exists demonstrating a definitive impact of protease-deficiency on outcome in malaria, and it is still unclear if low ADAMTS13 levels really represent a causal pathology mediating malarial disease or rather constitute an epiphenomenal consequence; also, the cut-off value of ADAMTS13 activity which is predictive of a complicated course of disease remains to be defined. More research is required to identify a causal relationship between reduced ADAMTS13 activity and mortality.

**Potential future therapeutic approaches in secondary TMAs**

**Inhibition of VWF release**

Anti-inflammatory agents such as steroids (106) or TNFa-inhibitors (107) block the release of acute phase reactants including VWF. However, their usefulness in sepsis related TMA has not been established, given that the evidence-level of recommendation regarding the use of steroids in sepsis has been downgraded by the Surviving Sepsis Campaign already in 2008 in view of inconsistent trial results.

**Inhibition of VWF activity**

The VWF-GPIb interaction is thought to be „un-druggable“ by conventional small molecule medicinal chemistry, since the area of contact between the A1 domain and the GPIb receptor is considered far too extensive for a small molecular weight antagonist to block (108). However, several small molecule inhibitors have been investigated so far.

**Anti-VWF aptamers**

Aptamers are oligonucleotides with drug-like properties that share some of the attributes of monoclonal antibodies, and were designed to block VWF-platelet binding in patients with TTP (29). Two aptamers have been investigated in more detail to date. The aptamer ARC1779 (32–40) is a first generation, nuclease resistant aptamer, binding to the activated VWF A1-domain with high potency and specificity. It has, however, a rapid elimination half-life and low subcutaneous bioavailability, which properties limit its therapeutic potential. ARC15105 (41) is a chemically advanced follower with a higher affinity to VWF and better pharmacokinetic profile than ARC1779, but it was never advanced into clinical development.

**Anti-VWF nanobody**

ALX-0081 (INN: caplacizumab) is a bivalent, and therefore highly potent humanised nanobody specifically targeting the GPIb binding site of VWF. It directly inhibits platelet binding to UL-VWF multimers.

In phase I studies ALX-0081 proved to be a potent and safe inhibitor of VWF mediated platelet aggregation over a wide range of doses, even when administered in combination with aspirin, clopidogrel and heparin. The international Phase II TITAN trial, a randomised controlled trial including 75 patients with acquired TTP was recently completed and yielded promising results. Caplacizumab significantly decreased time to platelet count normalisation by 39 %, with finally 81 % of patients achieving complete remission. The treatment was generally well tolerated and associated with fewer exacerbations as compared to placebo, with a manageable increase in bleeding tendency (44).

**VWF/GPIb inhibitor**

Agkisacucetin (Anfibatide), a non-enzymatic lectin-like protein purified from snake venom, is a first-in-class antagonist of the platelet GPIb receptor, which mediates binding of platelets to VWF at high shear flow. Considered as a novel antithrombotic agent, Anfibatide has been tested in vitro and in vivo, where it showed good antithrombotic effects without increasing the risk of bleedings in a phase I study in healthy volunteers (109). Anfibatide is currently evaluated in a randomised multi-centre phase II trial on patients with acute coronary syndrome [ClinicalTrials.gov NCT01585259] (110).

**ADAMTS13 replacement**

**Plasma exchange**

Case series and observational studies suggest that plasma exchange may be an effective adjunct therapy and may contribute to a better outcome in patients with severe sepsis and septic shock (111–114). In one case of TMA complicating septic DIC treatment with plasma exchange resulted in an improvement of TMA-related laboratory abnormalities along with a marked increase in consciousness. Nguyen et al. also reported beneficial effects of plasma exchange in critically ill children with secondary multiple organ failure-associated ADAMTS13 deficiency (115).

However, although plasma exchange is the gold standard of care in TTP, its value in the therapy of septic patients and its effectiveness in sepsis-related TMA remain to be established.

**Recombinant ADAMTS13 (rADAMTS13) and auto-antibody-resistant ADAMTS13 variants**

The administration of recombinant human ADAMTS13 completely prevented the emergence of thrombotic microangiopathy in ADAMTS13-deficient mice challenged with UL-VWF multimers (27). Accordingly, Plaimauer et al. found a linear relation between the inhibitor titer and the effective concentration of rADAMTS13 required for the reconstitution of ADAMTS13 activity in the presence of neutralising autoantibodies (28). However, the usefulness of rADAMTS13 in acquired TTP may be limited by an overwhelming amount of autoantibodies counteracting its effects. This drawback might be overcome by using gain-of-function ADAMTS13 variants, generated by site-directed mutagenesis, with
enhanced cleaving activity and increased resistance to binding and subsequent inhibition by autoantibodies (31).

Given that ADAMTS13 deficiency is considered multifactorial in secondary TMA, where it is at least partly caused by exhaustion and reduced synthesis, enzyme replacing strategies may have some therapeutic value there.

Complement inhibitor

Though the contributing role of the complement cascade regarding the pathogenesis and disease severity in TTP remains unknown, it is well conceivable that complement-targeting treatments could be of some benefit in selected patients with secondary TMA. Eculizumab, a terminal complement inhibitor, significantly improved renal function in patients with atypical haemolytic–uremic syndrome (116), the prototype of complement-mediated TMA, and has also demonstrated therapeutic efficacy in a case of TTP recalcitrant to daily plasma exchange therapy (117). One advantage of eculizumab in the treatment of septic TMA may be its high specificity to the complement component 5 (C5) resulting in preserved functions of the up-stream complement system. Considering the importance of humoral host defence in sepsis, eculizumab may accomplish both a sufficient suppression of excessive complement activation and the maintenance of essential immunologic function in sepsis related TMA. However, further studies on the prevalence and implications of complement defects in secondary TMA are needed.

Conclusion

ADAMTS13 proteolytically controls and prevents the occurrence of microvascular obstruction by VWF-platelet complexes under physiologic conditions. Absent or severely reduced enzyme activity causes the accumulation of ULVWF multimers, giant molecules with high spontaneous platelet-activating capacity. The longer the VWF multimers the higher is their procoagulant potential.

Recent evidence suggests that a pathomechanism formerly considered unique to TTP may take place also in other thrombotic and inflammatory diseases where it contributes to mortality. Whether these findings can finally be translated into therapeutic practice needs further investigation. To date, it is unclear whether VWF-directed therapy in patients with severely reduced ADAMTS13 levels or a high VWFpp/ADAMTS13 ratio may improve or even prevent sepsis/DIC related organ failure. VWF-targeted and enzyme replacement therapies would possibly help to eliminate microvascular occlusion and restore vital organ function. Since the use of recombinant ADAMTS13 significantly reduces the number of UL-VWF strings in vitro, prospective clinical trials of the effectiveness of protease replacement in vivo seem reasonable. It may be that carefully selected patients with secondary TMA could benefit from therapeutic approaches primarily intended for the use in TTP.

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Conflicts of interest

Dr. Bernd Jilma has served as the head of the data safety monitoring board of the TITAN trial [ALX-0081 (INN: caplacizumab)] sponsored by Abylnx and has been an investigator in TTP trials [ARC1779] sponsored by Archemix and a trial of ADAMTS13 in congenital TTP. None of the other authors has any conflicts of interest to disclose.

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