

HUS and the case for complement

Edward M. Conway

Centre for Blood Research, Life Sciences Institute, Department of Medicine, Faculty of Medicine, University of British Columbia, Vancouver, BC, Canada

Hemolytic-uremic syndrome (HUS) is a thrombotic microangiopathy that is characterized by microangiopathic hemolytic anemia, thrombocytopenia, and renal failure. Excess complement activation

underlies atypical HUS and is evident in Shiga toxin–induced HUS (STEC-HUS). This *Spotlight* focuses on new knowledge of the role of *Escherichia coli*–derived toxins and polyphosphate in modulating

complement and coagulation, and how they affect disease progression and response to treatment. Such new insights may impact on current and future choices of therapies for STEC-HUS. (*Blood*. 2015;126(18):2085-2090)

Introduction

Hemolytic-uremic syndrome (HUS) is a thrombotic microangiopathy (TMA) with microvascular and arteriolar wall thickening, swollen endothelial cells, and fibrin- and platelet-rich thrombi which compromise blood supply to end organs, particularly the kidney.^{1,2} Fragmentation of erythrocytes occurs from shear stress across partially obstructed vessels. Thrombocytopenia is caused by platelet consumption in clots and the reticuloendothelial system.

HUS is classified as either typical or atypical. Typical HUS is most often acquired from food contaminated with enterohemorrhagic *Escherichia coli* that produce Shiga-like toxins (Stx)³ and is thus referred to as Shiga toxin–induced HUS (STEC-HUS). Other microorganisms have also been implicated (<5%-10%).⁴ After exposure to STEC-HUS, an incubation period of ~3 days ensues, followed by 2 to 5 days of watery diarrhea, nausea, and fever (~30%). Gastroenteritis, with bloody diarrhea proceeds in ~50% to 70%, and is usually self-limited, but rarely is complicated with massive hemorrhage and/or bowel perforation.⁵ A minority of STEC-exposed patients subsequently develop HUS (5%-25%), and this often has serious long-term sequelae.⁶ Approximately 25% of STEC-HUS patients develop chronic renal insufficiency. Neurologic involvement (eg., strokes, seizures) is evident in 10% to 20% of cases and these account for much of the 1% to 5% mortality. Other organs may also be involved.

The remaining noninfective cases of HUS are referred to as atypical (aHUS). aHUS is chronic and recurring with >50% developing end-stage renal failure, and has an early mortality of 10% to 25%. There is overwhelming evidence that aHUS is caused by excess complement activation.⁷ Thus, most patients respond well to the complement inhibitor, eculizumab. In contrast, specific therapies for STEC-HUS are lacking.⁴

Recent advances in our understanding of complement and coagulation and the role of *E coli*–derived toxins (Stx and serine protease autotransporters of *Enterobacteriaceae* [SPATEs]) and polyphosphate in modulating these pathways, as discussed in this *Spotlight*, may help explain why STEC-HUS is less responsive to eculizumab, and hopefully aid in the rational design of STEC-HUS therapies.

The complement system

Complement comprises over 30 soluble and membrane-bound proteins, coordinated to eliminate pathogens and damaged cells.⁸ Complement activation proceeds via 3 pathways: classical (CP), lectin (LP), and alternative (AP). The CP is triggered by antigen-antibody complexes recognized by C1q, and the LP by sugars recognized primarily by mannose-binding lectin. This causes activation of proteases that cleave C4 and C2 to form the C4b2a LP/CP C3 convertase which proteolyzes C3 into C3b and C3a. The AP constitutively generates fluid-phase C3b-like C3H₂O. This binds to factor B (FB) that is cleaved by factor D (FD), yielding C3(H₂O)Bb. C3(H₂O)Bb cleaves C3, yielding C3b that is necessary to form surface-bound C3 convertase, C3bBb. With more C3b, substrate specificity of the convertase shifts to C5, generating C5a and C5b. C5b triggers assembly of the C5b-9 membrane attack complex (MAC) that lyses target pathogens/cells.

Complement is tightly regulated. The major fluid-phase AP-negative regulator, factor H (FH), competes with FB binding to C3b, acts as a cofactor for factor I (FI) cleavage/inactivation of C3b, and accelerates convertase decay. FH also interacts with thrombomodulin and von Willebrand factor (VWF), augmenting FI inactivation of C3b.⁹⁻¹¹ Membrane glycoproteins, CD55 and CD46, also promote convertase decay.¹² Anaphylatoxins C3a and C5a are modulated by their receptors¹³ and/or degraded by plasmin,¹⁴ matrix metalloproteinases,¹⁵ and activated thrombin-activatable fibrinolysis inhibitor (TAFIa).¹⁶ MAC formation is suppressed by CD59, mortalin/GRP75,¹⁷ clusterin, and vitronectin. Recently, we showed that polyphosphate also interferes with MAC assembly.¹⁸

aHUS: molecular defects causing excess complement activation

Mutations of genes that encode complement components, or antibodies that alter their function, account for ~60% to 70% of patients with inherited or sporadic aHUS. Affected genes and associated

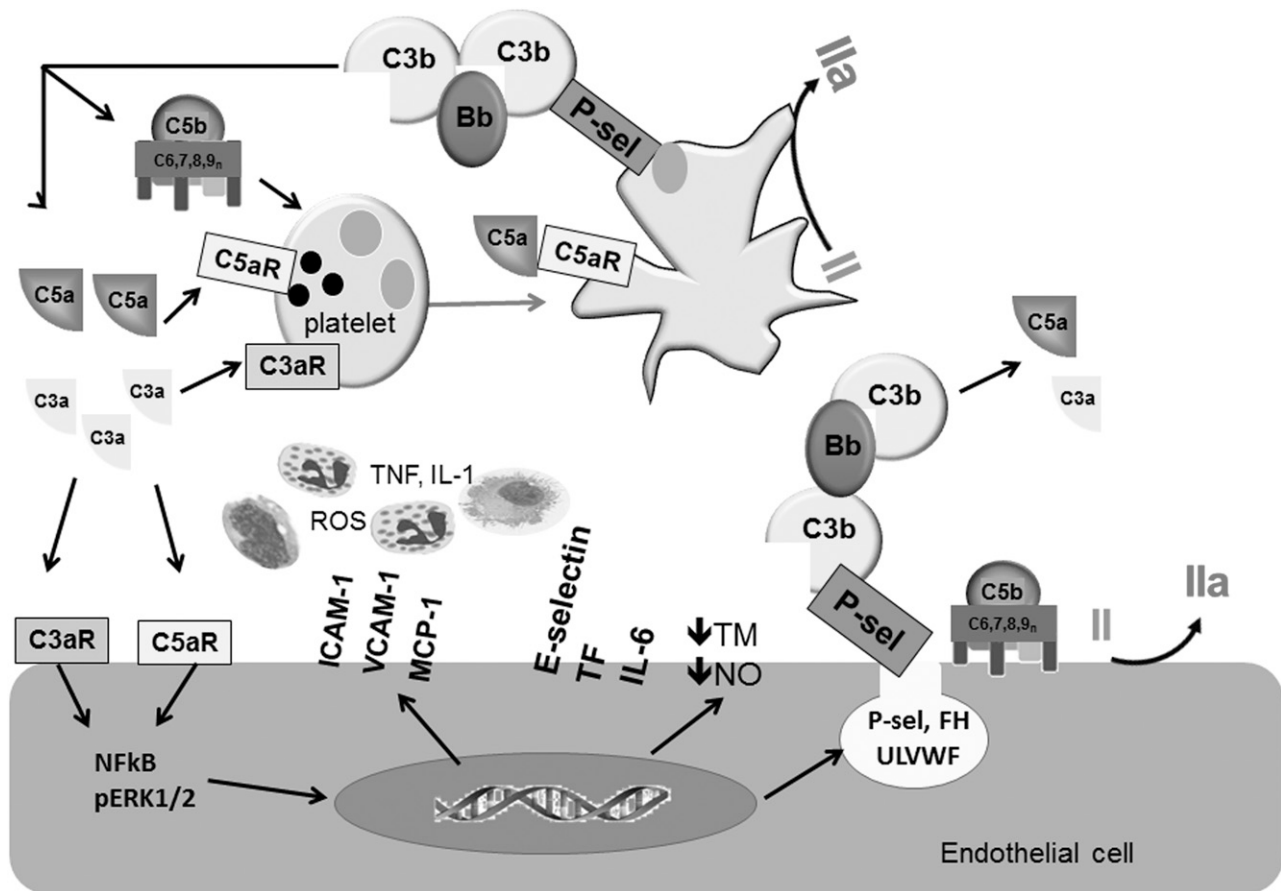


Figure 1. Complement activation and TMA. The scheme highlights some of the mechanisms by which unregulated complement activation provokes the development of microvascular thrombi via unremitting activation/damage to the endothelium and platelet. With excess complement activation, the abundant anaphylatoxins C3a and C5a bind to their widely expressed cognate receptors. Endothelial cells and platelets can thus be activated, whereupon intracellular proinflammatory and procoagulant signaling cascades are recruited (eg, nuclear factor κ B [NF κ B], phospho-ERK 1/2 [pERK1/2]). In endothelial cells, these upregulate expression of adhesion molecules (intercellular adhesion molecule-1 [ICAM-1], vascular cell adhesion molecule [VCAM-1], E-selectin), the release of proinflammatory cytokines (eg, monocyte chemoattractant protein 1 [MCP-1], interleukin-6 [IL-6]), exposure of TF, suppressed expression of the anticoagulant/anticoagulant thrombomodulin (TM), and reduced release of nitric oxide (NO). P-selectin, FH, and ULVWF multimers are secreted from endothelial cell Weibel-Palade bodies, and they are released from α -granules and/or the cytoplasm (for FH) of platelets. P-selectin and ULVWF facilitate platelet adhesion/aggregation, but also are receptors for C3b binding, allowing assembly of the AP C3 and C5 convertases (C3bBb and C3bBbC3b, respectively). These further amplify complement activation, with release of more C3a and C5a. Downstream, complement activation yields terminal pathway complexes (C5b-7, sC5b-9) which induce TF exposure, endothelial membrane “flipping” to support prothrombinase assembly, and the transformation of prothrombin (II) to the procoagulant, platelet activating and proinflammatory protease, thrombin (IIa). Thrombin and other procoagulant enzymes (eg, factor Xa) also feed back to cleave C5, fueling further activation of complement. C3a, C5a, and cell-released chemokines also recruit inflammatory cells, which in turn exacerbate endothelial damage via the release of reactive oxygen species and cytokines (eg, tumor necrosis factor α [TNF α], high-mobility group box 1 [HMGB1]). Activated platelets also release their granule contents which are primarily procoagulant and proinflammatory. Platelet and endothelial microparticles, released in response to exposure to C3a and C5a, carry complement factors and express TF to further promote coagulation and complement. ULVWF multimers support platelet adhesion/aggregation and the formation of thrombi, as well as further activation of complement. In aHUS, eculizumab interferes with cleavage of C5 to C5b and C5a, and effectively reverses the unremitting, self-propelling activation of the cascades that otherwise result in the TMA.

frequencies include FH (20%-30%), FI (5%-10%), CD46 (10%-15%), thrombomodulin (3%-4%), C3 (2%-10%), and FB (1%-4%).¹⁹⁻²¹ Hybrid forms of FH and FH-related proteins (3%-5%) have reduced activity,²² and autoantibodies against FH (5%-10%) reduce its binding to endothelium or C3b.²³ A mutation in diacylglycerol kinase- ϵ is a rare cause of aHUS, but its link to complement is not well established.^{24,25}

Spiraling activation of complement and coagulation

Complement overactivation, as occurs in aHUS, undermines vasculo-protective properties via several mechanisms²⁶ (Figure 1). C3a/C5a stimulate secretion of cytokines, promote leukocyte adhesion²⁷ and

tissue factor (TF) expression, suppress thrombomodulin, and induce P-selectin and VWF release. C3a/C5a also activate platelets,²⁸ causing granule secretion, exposure of P-selectin, and release of procoagulant microparticles. P-selectin recruits leukocytes and platelets and is a C3b receptor²⁹ for AP convertase assembly, amplifying complement activation. C5b-7 activates TF on monocytes,³⁰ and sC5b-9 activates platelets and endothelial cells, induces VWF and cytokine release, and promotes prothrombinase assembly and release of TF-expressing microparticles.³¹ Thrombin loops back to liberate C5a and generate a more damaging MAC.^{32,33}

Defects in processing VWF are linked to another TMA, thrombotic thrombocytopenia purpura.³⁴ But VWF also contributes to complement activation, and likely participates in HUS-associated thrombosis. When secreted, VWF is anchored to endothelium as ultra-large (ULVWF) multimeric strings.³⁵ These are normally cleaved by a disintegrin and metalloproteinase with a thrombospondin

type 1 motif, member 13 (ADAMTS13) into smaller, less procoagulant forms. FH binds to VWF and also reduces multimers,^{36,37} limiting complement and platelet activation/aggregation. Loss-of-function FH mutations that cause aHUS would facilitate formation of platelet-rich thrombi, and allow unregulated complement activation with generation of procoagulant products.

Overall, it is not surprising that a drug that restrains complement is effective for aHUS. Eculizumab interferes with C5 cleavage to C5a and C5b-9³⁸; its efficacy in aHUS is >80%.³⁹

STEC-HUS and complement activation

Does complement activation occur in STEC-HUS? If so, does it explain the TMA? Most patients infected with STEC exhibit heightened complement activation,⁴⁰ with increased generation of C3a, Bb, and sC5b-9, and C3 and C9 deposits on platelet-leukocyte aggregates and microvesicles.⁴¹⁻⁴³ Complement activation therefore likely contributes to endothelial damage and thrombosis in STEC-HUS. Clinical validation is, however, lacking, as there have not been controlled studies of eculizumab for STEC-HUS. In 2 small reports of affected children with neurologic impairment, benefit for some appeared to be derived from eculizumab.^{44,45} A similar result was reported for a small number of French adults.⁴⁶ However, review of its use in the large European outbreak did not reveal any benefit.⁴⁷ Controlled, prospective studies are needed. Nonetheless, the stunning success of eculizumab in aHUS has not been seen with STEC-HUS, suggesting that modulating complement, by itself, is inadequate.

Shiga toxin: direct effects on multiple pathways

The most frequent cause of STEC-HUS is *E coli* strain 0157:H7, but many others have been identified.^{48,49} Stx is the key virulence factor.⁵⁰⁻⁵² Upon ingestion, STEC colonize the gut, adhere to epithelial cells, destroy the brush border villi, and cause diarrhea.⁵² Stx is secreted through the epithelium where it can contact blood, but only trace amounts of circulating free toxin are found. Rather, Stx preferentially binds to platelets, neutrophils, monocytes, and possibly erythrocytes.^{53,54} Transfer of toxin to cells in target tissues is achieved via binding to globotriaosylceramide (Gb3) and globotetraosylceramide (Gb4), expressed by endothelial cells of the intestine, brain and kidney, podocytes, mesangial cells, and renal tubular epithelial cells.⁵⁵ An intriguing alternative virulence mechanism has been described in which Stx is internalized by circulating blood cells, and then released in microvesicles, which in turn are transferred into glomerular and peritubular capillary endothelial cells via Gb3-independent pathways for uptake by renal cells.⁵⁶ Whatever the mechanism, intracellular Stx interferes with ribosomal apparatus and blocks protein synthesis.⁵⁷ Stx also induces release of P-selectin and VWF from platelets and endothelial cells, which further activates platelets and neutrophils, induces neutrophil extracellular trap formation,⁵⁸ and amplifies the AP^{29,59} (Table 1). It interferes with FH, rendering the endothelium more vulnerable to complement.⁶⁰ Stx also stimulates release of C3- and C9-bearing microparticles from platelets and monocytes^{41,42} and in animal models, causes AP-dependent microvascular thrombosis, apoptosis of renal tubular cells, and podocyte loss and dysfunction.^{61,62}

Thus, at some stage(s) of the syndrome, Stx directly or indirectly activates complement and, in turn, coagulation. However, Stx has many complement-independent effects (Table 1). Upon internalization, Stx triggers proinflammatory signals,⁵⁵ promotes chemokine release, upregulates leukocyte adhesion molecules and TF, suppresses endothelial TF pathway inhibitor and thrombomodulin, induces platelet-neutrophil interactions,⁴¹ promotes release of TF-bearing microparticles,⁶³ and induces neutrophil production of reactive oxygen species.⁶⁴ Stx also interferes with ULVWF cleavage by ADAMTS13⁶⁵ and FH,⁶⁶ enhancing platelet adhesion/aggregation. Overall, Stx uses multiple means to promote thrombosis, not all of which are sensitive to anticomplement interventions.

SPATES: aiding Stx in crime

In addition to Stx, >25 SPATEs are produced by pathogenic strains of *E coli*⁶⁷ and their activities may further explain the variable response of STEC-HUS to eculizumab. SPATE proteases cleave/inactivate chemokines and adhesion molecules, disrupt leukocyte chemotaxis, transmigration, and activation, and dampen inflammatory and prothrombotic responses.⁶⁸ Several SPATEs have recently been shown to modulate complement (Table 1). For example, the serine protease Pic is secreted by strains of HUS-inducing STEC, including 0104:H4. Pic suppresses complement activation by proteolyzing C2, C3/C3b, and C4/C4b.^{68,69} Pic also synergizes with FI/FH to inactivate C3b, and des-arginates C3a, rendering it less active. EspP is a serine protease strongly associated with *E coli* 0157:H7.⁷⁰ EspP dampens complement activation by proteolyzing C3/C3b and C5.⁷¹ Interestingly, EspP also inactivates coagulation factor V and α 2-antiplasmin, contributing to mucosal bleeding and facilitating bacterial invasion. StcE is a metalloprotease that is also secreted by *E coli* 0157:H7.^{72,73} Correlated with high virulence,⁷⁴ StcE cleaves C1-esterase inhibitor (C1-INH), but enhances its capacity to neutralize C1s and MASPs. StcE also binds to host cell surfaces (eg, platelets, endothelial cells) and tethers C1-INH so that it can protect the pathogen and host cell by subverting complement activation.

By circumventing immune destruction, several SPATE proteases confer a survival advantage to the STEC, favoring persistence, invasion, and migration. Therapeutic interventions that suppress complement might therefore not be of benefit.

Polyphosphate

Beyond SPATEs, *E coli* and other bacteria rely on multiple mechanisms to evade complement.^{75,76} Polyphosphate (polyP) is an anionic, linear polymer of phosphate that is found in all cells.⁷⁷⁻⁷⁹ Identified first in cytoplasmic granules, it is localized in several cell compartments, and, in *E coli*, is prominent in the membrane.⁸⁰ In prokaryotes, polyP exhibits prosurvival properties as an energy source, a metal ion chelator, and a molecular chaperone, and is essential for microorganism pathogenicity.^{81,82} In mammals, polyP promotes coagulation at several steps in the cascade.^{83,84} We recently determined that polyP also dampens complement activation (Table 1), a finding in line with reports that a mutant form of *Neisseria meningitidis* with excess polyP, is protected against complement-mediated killing.⁸⁵

PolyP destabilizes C5b,6, and reduces binding of the resultant C5b-7 and C5b-8 to the target membrane.¹⁸ Thus, polyP in the

Table 1. Summary of diverse biological effects of Stx, SPATEs, and polyphosphate that modulate activation of complement and coagulation during STEC-HUS

Summary of diverse biological effects of Stx, SPATEs, and polyphosphate	
Stx	
Direct effects that promote complement activation	<ul style="list-style-type: none"> • Induces release and cell surface expression of P-selectin (receptor for C3b) • Induces release of VWF (site for AP convertase assembly) • Interferes with functions of FH • Induces release of C3- and C9-bearing microparticles from platelets and monocytes
Direct effects that induce endothelial damage and activation of coagulation	<ul style="list-style-type: none"> • Activates endothelial cells, platelets, monocytes, and neutrophils • Induces release of reactive oxygen species from neutrophils • Induces expression of TF • Suppresses expression of TM and TF pathway inhibitor • Induces endothelial expression of leukocyte adhesion molecules (e.g., ICAM-1, VCAM-1) • Induces platelet-neutrophil interactions • Interferes with cleavage of ULVWF by ADAMTS13 and FH • Promotes release of TF-bearing microparticles • Induces proinflammatory cytokine (e.g., TNF, IL-8) release by endothelial cells and monocytes
SPATEs	
Pic	<ul style="list-style-type: none"> • Proteolyzes and inactivates C2, C3/C3b, C4/C4b • Des-arginates C3a
EspP	<ul style="list-style-type: none"> • Proteolyzes and inactivates C3/C3b, C5 • Inactivates coagulation factor V and α2-antiplasmin
StcE	<ul style="list-style-type: none"> • Cleaves C1-INH and increases its neutralization of C1s and MASP • Binds to host/pathogen surfaces and tethers C1-INH
PolyP	<ul style="list-style-type: none"> • Promotes coagulation activation at several steps in cascade • Destabilizes C5b,6 and interferes with MAC assembly on host and pathogen • Binds to C1-INH and increases its neutralization of C1s and MASP • Binds to FH (function unknown)

ICAM, intercellular adhesion molecule; IL, interleukin; TM, thrombomodulin; TNF, tumor necrosis factor; VCAM, vascular cell adhesion molecule.

membrane of STEC likely provides a barrier against MAC assembly.⁸⁰ PolyP binds to C1-INH¹⁸ and enhances its activity, suppressing the CP and LP.⁸⁶ PolyP also binds to FH,¹⁸ although the functional consequences are unknown. Interestingly, like polyP, C1-INH and FH are released from activated platelets, where they may coat and protect host cells⁸⁷ from complement activation, convertase assembly, and MAC binding/integration. Similar to C1-INH, FH may also be recruited to pathogen surfaces for immune evasion.⁷⁶ We speculate that released polyP binds to FH, C1-INH, and/or other cationic proteins on the surface of host cells and pathogenic *E coli*, providing an additional barrier against complement-mediated damage. Once again, this may help explain why a complement inhibitor alone may not be sufficient to resolve the manifestations of STEC-HUS.

Conclusions

For aHUS, where disturbances in complement regulation define a “starting point” for amplification of cascades that lead to TMA, intervening with eculizumab is effective and currently the favored first-line treatment. The situation is more complex for STEC-HUS. The virulence factor Stx(s), which is requisite for development of STEC-HUS, has complement-activating properties, but also triggers endothelial injury, podocyte and renal tubular damage, platelet activation and thrombosis, via pathways that likely vary during the course of the syndrome. It follows that preventing C5 cleavage with eculizumab may not be uniformly effective in abrogating the associated TMA. Furthermore, pathogenic *E coli* that cause HUS also use multiple means to prolong survival and enhance virulence, partly by encapsulating its toxin in microvesicles, synthesizing polyP, and secreting SPATEs along with Stx(s). These either evade or dampen complement-dependent immune-mediated killing, and again may help explain

the suboptimal response to eculizumab. They also point to potential shortcomings of using HUS models that rely solely on Stx induction,^{50,51} and underline the importance of seeking alternative therapeutic approaches. In that respect, efforts are under way to delineate the pathways by which the toxins traffick to target organs, and to characterize factors synthesized and/or secreted by enterohemorrhagic *E coli*. These are uncovering potential strain-specific targets to reduce pathogen persistence, replication and adhesion, biofilm formation, and Stx invasion.⁸⁸ Closer to the clinic, vaccines and neutralizing anti-Stx antibodies are in development,⁸⁹⁻⁹¹ and these will hopefully reduce the incidence and severity of STEC-HUS.

Acknowledgments

E.M.C. is supported by operating grants from the Canadian Institutes for Health Research (CIHR), the Natural Sciences and Engineering Research Council of Canada (NSERC), and the Canada Foundations for Innovation (CFI).

E.M.C. holds a CSL Behring Research Chair and a Tier 1 Canada Research Chair in Endothelial Cell Biology, and is an Adjunct Scientist with the Canadian Blood Services.

Authorship

Contribution: E.M.C. researched and wrote the manuscript.

Conflict-of-interest disclosure: The author declares no competing financial interests.

Correspondence: Edward M. Conway, Centre for Blood Research (CBR), University of British Columbia, 4306-2350 Health Sciences Mall, Vancouver, BC V6T 1Z3, Canada; e-mail: ed.conway@ubc.ca.

References

- Gasser C, Gautier E, Steck A, Siebenmann RE, Oechslin R. Hemolytic-uremic syndrome: bilateral necrosis of the renal cortex in acute acquired hemolytic anemia [in German]. *Schweiz Med Wochenschr.* 1955;85(38-39):905-909.
- Hofer J, Rosales A, Fischer C, Giner T. Extra-renal manifestations of complement-mediated thrombotic microangiopathies. *Front Pediatr.* 2014;2:97.
- Frank C, Werber D, Cramer JP, et al; HUS Investigation Team. Epidemic profile of Shiga-toxin-producing *Escherichia coli* O104:H4 outbreak in Germany. *N Engl J Med.* 2011;365(19):1771-1780.
- Salvadori M, Bertoni E. Update on hemolytic uremic syndrome: diagnostic and therapeutic recommendations. *World J Nephrol.* 2013;2(3):56-76.
- Grisaru S. Management of hemolytic-uremic syndrome in children. *Int J Nephrol Renovasc Dis.* 2014;7:231-239.
- Rosales A, Hofer J, Zimmerhackl LB, et al; German-Austrian HUS Study Group. Need for long-term follow-up in enterohemorrhagic *Escherichia coli*-associated hemolytic uremic syndrome due to late-emerging sequelae. *Clin Infect Dis.* 2012;54(10):1413-1421.
- Sperati CJ, Moliterno AR. Thrombotic microangiopathy: focus on atypical hemolytic uremic syndrome. *Hematol Oncol Clin North Am.* 2015;29(3):541-559.
- Ricklin D, Lambris JD. Complement-targeted therapeutics. *Nat Biotechnol.* 2007;25(11):1265-1275.
- Delvaeye M, Noris M, De Vriese A, et al. Thrombomodulin mutations in atypical hemolytic-uremic syndrome. *N Engl J Med.* 2009;361(4):345-357.
- Rayes J, Roumenina LT, Dimitrov JD, et al. The interaction between factor H and VWF increases factor H cofactor activity and regulates VWF prothrombotic status. *Blood.* 2014;123(1):121-125.
- Feng S, Liang X, Kroll MH, Chung DW, Afshar-Kharghan V. von Willebrand factor is a cofactor in complement regulation. *Blood.* 2015;125(6):1034-1037.
- Kim DD, Song WC. Membrane complement regulatory proteins. *Clin Immunol.* 2006;118(2-3):127-136.
- Bosmann M, Ward PA. Role of C3, C5 and anaphylatoxin receptors in acute lung injury and in sepsis. *Adv Exp Med Biol.* 2012;946:147-159.
- Barthel D, Schindler S, Zipfel PF. Plasminogen is a complement inhibitor. *J Biol Chem.* 2012;287(22):18831-18842.
- Bellac CL, Dufour A, Krisinger MJ, et al. Macrophage matrix metalloproteinase-12 dampens inflammation and neutrophil influx in arthritis. *Cell Reports.* 2014;9(2):618-632.
- Campbell WD, Lazoura E, Okada N, Okada H. Inactivation of C3a and C5a octapeptides by carboxypeptidase R and carboxypeptidase N. *Microbiol Immunol.* 2002;46(2):131-134.
- Saar Ray M, Moskovich O, Iosefson O, Fishelson Z. Mortalin/GRP75 binds to complement C9 and plays a role in resistance to complement-dependent cytotoxicity. *J Biol Chem.* 2014;289(21):15014-15022.
- Wat JM, Foley JH, Krisinger MJ, et al. Polyphosphate suppresses complement via the terminal pathway. *Blood.* 2014;123(5):768-776.
- Mele C, Remuzzi G, Noris M. Hemolytic uremic syndrome. *Semin Immunopathol.* 2014;36(4):399-420.
- Noris M, Caprioli J, Bresin E, et al. Relative role of genetic complement abnormalities in sporadic and familial aHUS and their impact on clinical phenotype. *Clin J Am Soc Nephrol.* 2010;5(10):1844-1859.
- Bresin E, Rurali E, Caprioli J, et al; European Working Party on Complement Genetics in Renal Diseases. Combined complement gene mutations in atypical hemolytic uremic syndrome influence clinical phenotype. *J Am Soc Nephrol.* 2013;24(3):475-486.
- Francis NJ, McNicholas B, Awan A, et al. A novel hybrid CFH/CFHR3 gene generated by a microhomology-mediated deletion in familial atypical hemolytic uremic syndrome. *Blood.* 2012;119(2):591-601.
- Abarrategui-Garrido C, Martínez-Barricarte R, López-Trascasa M, de Córdoba SR, Sánchez-Corral P. Characterization of complement factor H-related (CFHR) proteins in plasma reveals novel genetic variations of CFHR1 associated with atypical hemolytic uremic syndrome. *Blood.* 2009;114(19):4261-4271.
- Lemaire M, Frémeaux-Bacchi V, Schaefer F, et al. Recessive mutations in DGKE cause atypical hemolytic-uremic syndrome. *Nat Genet.* 2013;45(5):531-536.
- Bruneau S, Néel M, Roumenina LT, et al. Loss of DGKE induces endothelial cell activation and death independently of complement activation. *Blood.* 2015;125(6):1038-1046.
- Conway EM. Reincarnation of ancient links between coagulation and complement. *J Thromb Haemost.* 2015;13(suppl 1):S121-S132.
- Monsinjon T, Gasque P, Chan P, Ischenko A, Brady JJ, Fontaine MC. Regulation by complement C3a and C5a anaphylatoxins of cytokine production in human umbilical vein endothelial cells. *FASEB J.* 2003;17(9):1003-1014.
- Verschoor A, Langer HF. Crosstalk between platelets and the complement system in immune protection and disease. *Thromb Haemost.* 2013;110(5):910-919.
- Del Conde I, Cruz MA, Zhang H, López JA, Afshar-Kharghan V. Platelet activation leads to activation and propagation of the complement system. *J Exp Med.* 2005;201(6):871-879.
- Langer F, Spath B, Fischer C, et al. Rapid activation of monocyte tissue factor by antithymocyte globulin is dependent on complement and protein disulfide isomerase. *Blood.* 2013;121(12):2324-2335.
- Wiedmer T, Esmon CT, Sims PJ. Complement proteins C5b-9 stimulate procoagulant activity through platelet prothrombinase. *Blood.* 1986;68(4):875-880.
- Huber-Lang M, Sarma JV, Zetoun FS, et al. Generation of C5a in the absence of C3: a new complement activation pathway. *Nat Med.* 2006;12(6):682-687.
- Krisinger MJ, Goebeler V, Lu Z, et al. Thrombin generates previously unidentified C5 products that support the terminal complement activation pathway. *Blood.* 2012;120(8):1717-1725.
- Zheng XL. ADAMTS13 and von Willebrand factor in thrombotic thrombocytopenic purpura. *Annu Rev Med.* 2015;66:211-225.
- Turner N, Nolasco L, Nolasco J, Sartain S, Moake J. Thrombotic microangiopathies and the linkage between von Willebrand factor and the alternative complement pathway. *Semin Thromb Hemost.* 2014;40(5):544-550.
- Feng S, Liang X, Cruz MA, et al. The interaction between factor H and Von Willebrand factor. *PLoS One.* 2013;8(8):e73715.
- Nolasco L, Nolasco J, Feng S, Afshar-Kharghan V, Moake J. Human complement factor H is a reductase for large soluble von Willebrand factor multimers—brief report. *Arterioscler Thromb Vasc Biol.* 2013;33(11):2524-2528.
- Gruppo RA, Rother RP. Eculizumab for congenital atypical hemolytic-uremic syndrome. *N Engl J Med.* 2009;360(5):544-546.
- Licht C, Greenbaum LA, Muus P, et al. Efficacy and safety of eculizumab in atypical hemolytic uremic syndrome from 2-year extensions of phase 2 studies. *Kidney Int.* 2015;87(5):1061-1073.
- Noris M, Mescia F, Remuzzi G. STEC-HUS, atypical HUS and TTP are all diseases of complement activation. *Nat Rev Nephrol.* 2012;8(11):622-633.
- Stähl AL, Sartz L, Karpman D. Complement activation on platelet-leukocyte complexes and microparticles in enterohemorrhagic *Escherichia coli*-induced hemolytic uremic syndrome. *Blood.* 2011;117(20):5503-5513.
- Thurman JM, Marians R, Emlen W, et al. Alternative pathway of complement in children with diarrhea-associated hemolytic uremic syndrome. *Clin J Am Soc Nephrol.* 2009;4(12):1920-1924.
- Ferraris JR, Ferraris V, Acquier AB, et al. Activation of the alternative pathway of complement during the acute phase of typical haemolytic uraemic syndrome. *Clin Exp Immunol.* 2015;181(1):118-125.
- Lapeyraque AL, Malina M, Frémeaux-Bacchi V, et al. Eculizumab in severe Shiga-toxin-associated HUS. *N Engl J Med.* 2011;364(26):2561-2563.
- Pape L, Hartmann H, Bange FC, Suerbaum S, Bueltmann E, Ahlenstiel-Grunow T. Eculizumab in typical hemolytic uremic syndrome (HUS) with neurological involvement. *Medicine (Baltimore).* 2015;94(24):e1000.
- Delmas Y, Vendrely B, Clouzeau B, et al. Outbreak of *Escherichia coli* O104:H4 haemolytic uraemic syndrome in France: outcome with eculizumab. *Nephrol Dial Transplant.* 2014;29(3):565-572.
- Menne J, Nitschke M, Stingle R, et al; EHEC-HUS consortium. Validation of treatment strategies for enterohaemorrhagic *Escherichia coli* O104:H4 induced haemolytic uraemic syndrome: case-control study. *BMJ.* 2012;345:e4565.
- Miko A, Rivas M, Bentancor A, Delannoy S, Fach P, Beutin L. Emerging types of Shiga toxin-producing *E. coli* (STEC) O178 present in cattle, deer, and humans from Argentina and Germany. *Front Cell Infect Microbiol.* 2014;4:78.
- Loos S, Ahlenstiel T, Kranz B, et al. An outbreak of Shiga toxin-producing *Escherichia coli* O104:H4 hemolytic uremic syndrome in Germany: presentation and short-term outcome in children. *Clin Infect Dis.* 2012;55(6):753-759.
- Siegler RL, Pysker TJ, Lou R, Tesh VL, Taylor FB Jr. Response to Shiga toxin-1, with and without lipopolysaccharide, in a primate model of hemolytic uremic syndrome. *Am J Nephrol.* 2001;21(5):420-425.
- Sauter KA, Melton-Celsa AR, Larkin K, Troxell ML, O'Brien AD, Magun BE. Mouse model of hemolytic-uremic syndrome caused by endotoxin-free Shiga toxin 2 (Stx2) and protection from lethal outcome by anti-Stx2 antibody. *Infect Immun.* 2008;76(10):4469-4478.
- Melton-Celsa AR. Shiga Toxin (Stx) classification, structure, and function. *Microbiol Spectr.* 2014;2(2).

53. Ståhl AL, Sartz L, Nelsson A, Békássy ZD, Karpman D. Shiga toxin and lipopolysaccharide induce platelet-leukocyte aggregates and tissue factor release, a thrombotic mechanism in hemolytic uremic syndrome. *PLoS One*. 2009; 4(9):e6990.
54. Brigotti M, Carnicelli D, Arfilli V, et al. Identification of TLR4 as the receptor that recognizes Shiga toxins in human neutrophils. *J Immunol*. 2013; 191(9):4748-4758.
55. Obrig TG. Escherichia coli Shiga toxin mechanisms of action in renal disease. *Toxins (Basel)*. 2010;2(12):2769-2794.
56. Ståhl AL, Arvidsson I, Johansson KE, et al. A novel mechanism of bacterial toxin transfer within host blood cell-derived microvesicles. *PLoS Pathog*. 2015;11(2):e1004619.
57. Endo Y, Tsurugi K, Yutsudo T, Takeda Y, Ogasawara T, Igarashi K. Site of action of a Vero toxin (VT2) from Escherichia coli O157:H7 and of Shiga toxin on eukaryotic ribosomes. RNA N-glycosidase activity of the toxins. *Eur J Biochem*. 1988;171(1-2):45-50.
58. Fuchs TA, Kremer Hovinga JA, Schatzberg D, Wagner DD, Lämmle B. Circulating DNA and myeloperoxidase indicate disease activity in patients with thrombotic microangiopathies. *Blood*. 2012;120(6):1157-1164.
59. Geelen J, van den Biggelaar M, Linssen P, van der Velden T, Mertens K, Monnens L. The effect of shiga toxin on weibel-palade bodies in primary human endothelial cells. *Nephron Extra*. 2014; 4(2):101-107.
60. Poolpol K, Orth-Höller D, Speth C, et al. Interaction of Shiga toxin 2 with complement regulators of the factor H protein family. *Mol Immunol*. 2014;58(1):77-84.
61. Morigi M, Galbusera M, Gastoldi S, et al. Alternative pathway activation of complement by Shiga toxin promotes exuberant C3a formation that triggers microvascular thrombosis. *J Immunol*. 2011;187(1):172-180.
62. Locatelli M, Buelli S, Pezzotta A, et al. Shiga toxin promotes podocyte injury in experimental hemolytic uremic syndrome via activation of the alternative pathway of complement. *J Am Soc Nephrol*. 2014;25(8):1786-1798.
63. Brigotti M, Carnicelli D, Ravanelli E, et al. Molecular damage and induction of proinflammatory cytokines in human endothelial cells exposed to Shiga toxin 1, Shiga toxin 2, and alpha-sarcin. *Infect Immun*. 2007;75(5): 2201-2207.
64. Gomez SA, Abrey-Recalde MJ, Panek CA, et al. The oxidative stress induced in vivo by Shiga toxin-2 contributes to the pathogenicity of haemolytic uraemic syndrome. *Clin Exp Immunol*. 2013;173(3):463-472.
65. Lo NC, Turner NA, Cruz MA, Moake J. Interaction of Shiga toxin with the A-domains and multimers of von Willebrand Factor. *J Biol Chem*. 2013; 288(46):33118-33123.
66. Orth D, Khan AB, Naim A, et al. Shiga toxin activates complement and binds factor H: evidence for an active role of complement in hemolytic uremic syndrome. *J Immunol*. 2009; 182(10):6394-6400.
67. Dautin N. Serine protease autotransporters of enterobacteriaceae (SPATEs): biogenesis and function. *Toxins (Basel)*. 2010;2(6):1179-1206.
68. Ayala-Lujan JL, Vijayakumar V, Gong M, Smith R, Santiago AE, Ruiz-Perez F. Broad spectrum activity of a lectin-like bacterial serine protease family on human leukocytes. *PLoS One*. 2014; 9(9):e107920.
69. Abreu AG, Fraga TR, Granados Martínez AP, et al. The serine protease Pic from enteroaggregative Escherichia coli mediates immune evasion by the direct cleavage of complement proteins. *J Infect Dis*. 2015;212(1): 106-115.
70. Orth D, Ehrlenbach S, Brockmeyer J, et al. EspP, a serine protease of enterohemorrhagic Escherichia coli, impairs complement activation by cleaving complement factors C3/C3b and C5. *Infect Immun*. 2010;78(10):4294-4301.
71. Weiss A, Joerhs H, Brockmeyer J. Structural and functional characterization of cleavage and inactivation of human serine protease inhibitors by the bacterial SPATE protease EspP α from enterohemorrhagic E. coli. *PLoS One*. 2014; 9(10):e111363.
72. Latham WW, Bergsbaken T, Welch RA. Potentiation of C1 esterase inhibitor by StcE, a metalloprotease secreted by Escherichia coli O157:H7. *J Exp Med*. 2004;199(8):1077-1087.
73. Yu AC, Worrall LJ, Strynadka NC. Structural insight into the bacterial mucinase StcE essential to adhesion and immune evasion during enterohemorrhagic E. coli infection. *Structure*. 2012;20(4):707-717.
74. Kobayashi N, Lee K, Yamazaki A, et al. Virulence gene profiles and population genetic analysis for exploration of pathogenic serogroups of Shiga toxin-producing Escherichia coli. *J Clin Microbiol*. 2013;51(12):4022-4028.
75. Berends ET, Kuipers A, Ravesloot MM, Urbanus RT, Rooijackers SH. Bacteria under stress by complement and coagulation. *FEMS Microbiol Rev*. 2014;38(6):1146-1171.
76. Blom AM, Hallström T, Riesbeck K. Complement evasion strategies of pathogens-acquisition of inhibitors and beyond. *Mol Immunol*. 2009;46(14): 2808-2817.
77. Caen J, Wu Q. Hageman factor, platelets and polyphosphates: early history and recent connection. *J Thromb Haemost*. 2010;8(8): 1670-1674.
78. Rao NN, Kornberg A. Inorganic polyphosphate regulates responses of Escherichia coli to nutritional stringencies, environmental stresses and survival in the stationary phase. *Prog Mol Subcell Biol*. 1999;23:183-195.
79. Achbergerová L, Nahálka J. Polyphosphate—an ancient energy source and active metabolic regulator. *Microb Cell Fact*. 2011;10:63.
80. Castuma CE, Huang R, Kornberg A, Reusch RN. Inorganic polyphosphates in the acquisition of competence in Escherichia coli. *J Biol Chem*. 1995;270(22):12980-12983.
81. Moreno SN, Docampo R. Polyphosphate and its diverse functions in host cells and pathogens. *PLoS Pathog*. 2013;9(5):e1003230.
82. Gray MJ, Wholey WY, Wagner NO, et al. Polyphosphate is a primordial chaperone. *Mol Cell*. 2014;53(5):689-699.
83. Müller F, Mutch NJ, Schenk WA, et al. Platelet polyphosphates are proinflammatory and procoagulant mediators in vivo. *Cell*. 2009;139(6): 1143-1156.
84. Travers RJ, Smith SA, Morrissey JH. Polyphosphate, platelets, and coagulation. *Int J Lab Hematol*. 2015;37(Suppl 1):31-35.
85. Zhang Q, Li Y, Tang CM. The role of the exopolyphosphatase PPX in avoidance by Neisseria meningitidis of complement-mediated killing. *J Biol Chem*. 2010;285(44):34259-34268.
86. Lameignere E, Wijeyewickrema L, O'Byrne A, et al. Polyphosphate acts as a cofactor for C1-inhibitor-mediated regulation of the classical pathway of complement. *J Thromb Haemostasis Suppl*. 2015;2(suppl 13):616.
87. Licht C, Pluthero FG, Li L, et al. Platelet-associated complement factor H in healthy persons and patients with atypical HUS. *Blood*. 2009;114(20):4538-4545.
88. Melton-Celsa AR, O'Brien AD. New therapeutic developments against Shiga toxin-producing Escherichia coli. *Microbiol Spectr*. 2014;2(5).
89. Andrade GR, New RR, Sant'Anna OA, et al. A universal polysaccharide conjugated vaccine against O111 E. coli. *Hum Vaccin Immunother*. 2014;10(10):2864-2874.
90. Choi KS, Kim SH, Kim ED, et al. Protection from hemolytic uremic syndrome by eyedrop vaccination with modified enterohemorrhagic E. coli outer membrane vesicles. *PLoS One*. 2014; 9(7):e100229.
91. Melton-Celsa AR, Carvalho HM, Thuning-Roberson C, O'Brien AD. Protective efficacy and pharmacokinetics of human/mouse chimeric anti-Stx1 and anti-Stx2 antibodies in mice. *Clin Vaccine Immunol*. 2015;22(4):448-455.