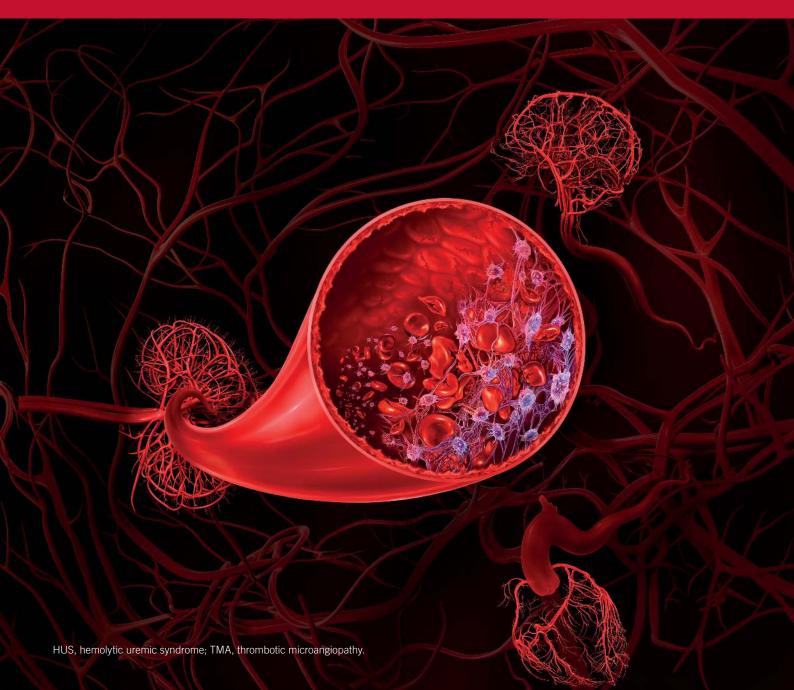
Patients with **atypical-HUS** can be at continuous risk of the life-threatening consequences of unpredictable complement-mediated TMA¹

Chronic, uncontrolled complement activity in atypical-HUS leads to ongoing endothelial injury, organ damage, and potential death^{1,2}

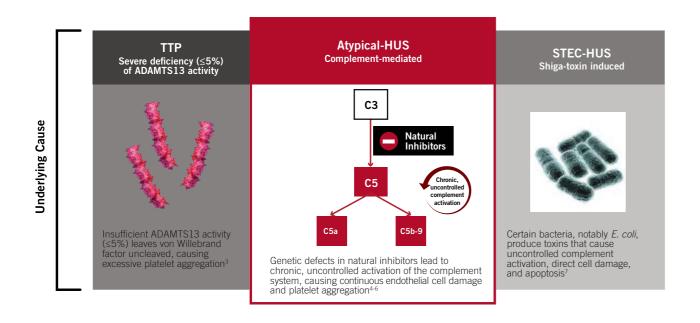


The information in this presentation is intended as educational information for healthcare professionals. It does not replace a healthcare professional's judgment or clinical diagnosis.

What Is Thrombotic Microangiopathy?

TMA manifests as thrombocytopenia, microangiopathic hemolysis, and organ involvement^{3,4}

TMAs present with similar signs and symptoms, but have distinct underlying causes^{3,4}



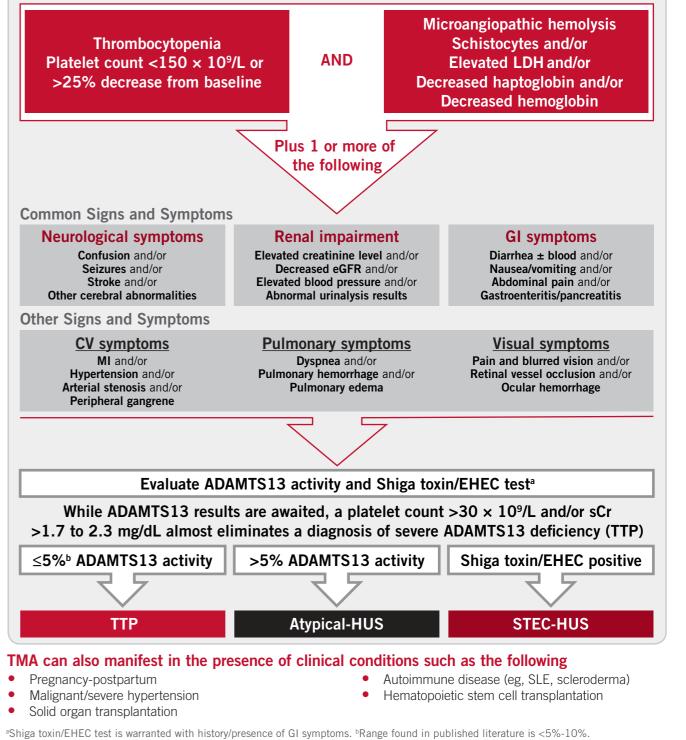
When you suspect TMA

- Rapidly take a sample to test for ADAMTS13 activity PRIOR TO intervention to ensure accurate test results^{3,4}
- Confirm negative for Shiga toxin/EHEC^{3,7}
- Get medical history including previous TMA and other potential causes³
- Obtain family history of TMA or renal impairment^{3,5}

ADAMTS13, a disintegrin and metalloproteinase with a thrombospondin type 1 motif member 13; CV, cardiovascular; eGFR, estimated glomerular filtration rate; EHEC, enterohemorrhagic E. coli; GI, gastrointestinal; HUS, hemolytic uremic syndrome; LDH, lactate dehydrogenase; MI, myocardial infarction; STEC-HUS, Shiga toxinproducing E. coli-hemolytic uremic syndrome; SLE, systemic lupus erythematosus; TMA, thrombotic microangiopathy; TTP, thrombotic thrombocytopenic purpura.

A clinical diagnosis of atypical-HUS in a patient with signs and symptoms of TMA requires exclusion of other underlying causes

Differential diagnosis for TMAs: Atypical-HUS, TTP, and STEC-HUS^{3,4,8,9}



Early identification of atypical-HUS allows for rapid disease management³

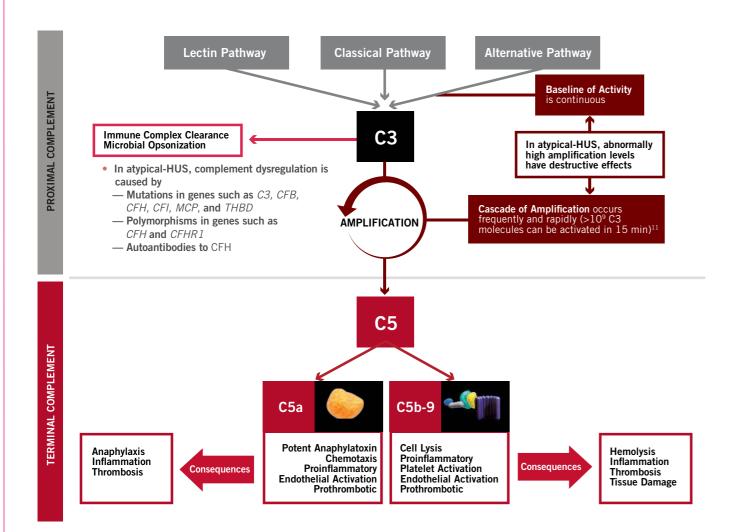




Pathophysiology of Atypical-HUS

Atypical-HUS results from chronic, uncontrolled complement activity that may lead to progressive and life-threatening complications⁴

Genetic mutations, polymorphisms, and autoantibodies lead to chronic, uncontrolled complement activity^{2,7,10}

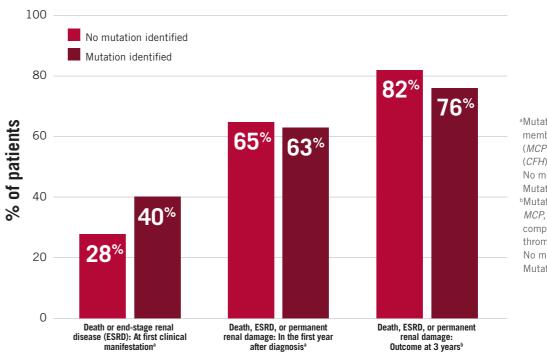


The number of new genetic abnormalities discovered in patients with atypical-HUS continues to increase over time¹²⁻¹⁵

45% of patients with atypical-HUS have an identifable genetic mutation or anticomplement factor H antibody¹⁶

Regardless of whether or not a mutation is identified, patients with atypical-HUS have similarly devastating outcomes^{12,16,17}

High morbidity and mortality regardless of mutation identification



• Identification of genetic complement mutations is not required for atypical-HUS diagnosis or management decisions³

ESRD, estimated glomerular filtration rate; CFH, complement factor H; HUS, hemolytic uremic syndrome.

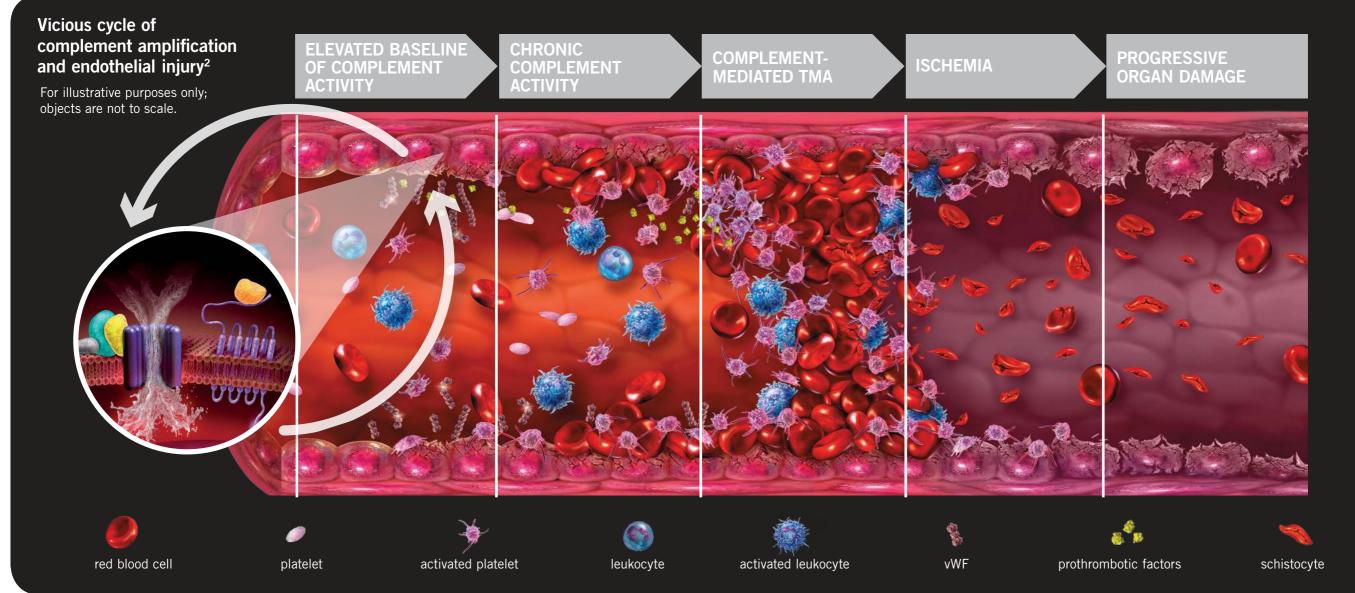
Role of Genetics in Atypical-HUS

5

Role of Genetics in Atypical-HUS

^aMutations consisted of membrane cofactor protein (MCP), complement factor H (CFH), and factor | (CFI), No mutation identified: n = 81 Mutation identified: n = 60. ^bMutations consisted of MCP, CFH, CFI, complement component 3 (*C3*), and thrombomodulin (THBD) No mutation identified: n = 119. Mutation identified: n = 116.

Chronic, uncontrolled complement activity can result in continuous endothelial damage and ongoing risk of TMA^{2,3,18}



Individual with atypical-HUS

- The assembly of multiple C5b-9 complexes on the surface of endothelial cells causes endothelial injury and platelet activation^{2,19-21}
- Binding of C5a to the C5a receptor results in a decrease in the endothelium's anticomplement and antithrombogenic properties^{2,20,22,23}
- Disrupted endothelial cells

6

- Release complement-activating microparticles, resulting in a vicious cycle of endothelial activation, complement amplification, and ongoing endothelial injury^{2,24}
- Release prothrombotic coagulation proteins, activate platelets, and recruit leukocytes, resulting in the formation of thrombi in small blood vessels throughout the body²

- Uncontrolled complement activity causes ongoing vascular endothelial injury, resulting in TMA lesions and progressive organ damage^{1,18,25}
- Biomarkers of complement activation, inflammation, endothelial cell activation and damage, coagulation, and renal damage (eg, Ba, sTNFR1, sVCAM-1, D-dimer, U-Cystatin C) are similarly elevated among patients with atypical-HUS receiving or not receiving plasma exchange or plasma infusion²⁶

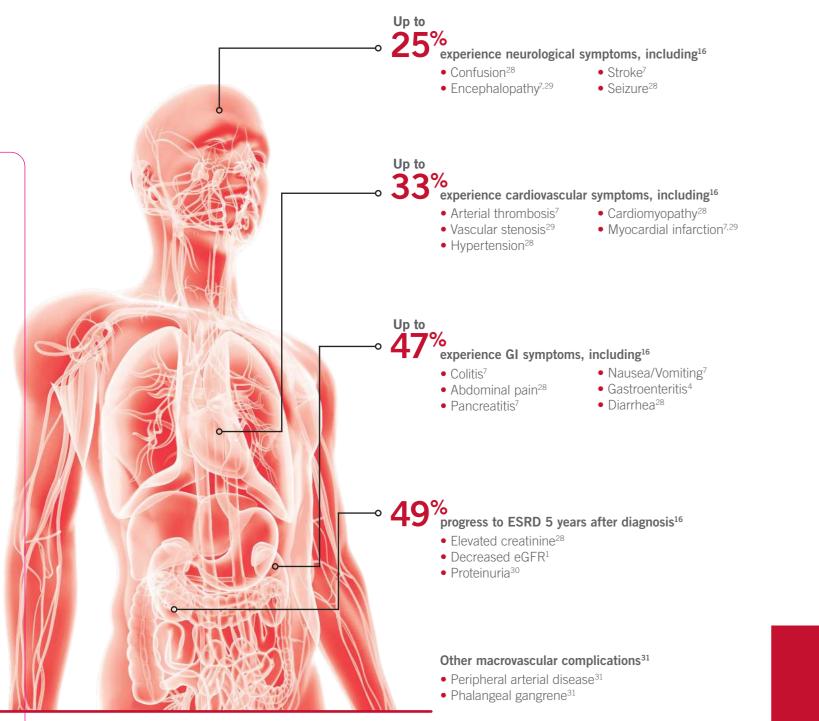
TMA lesions from repeated endothelial injury can progress toward irreversible tissue damage^{1,18,25}

Pathophysiology of Atypical-HUS

Atypical-HUS Is a Systemic, Life-threatening Disease

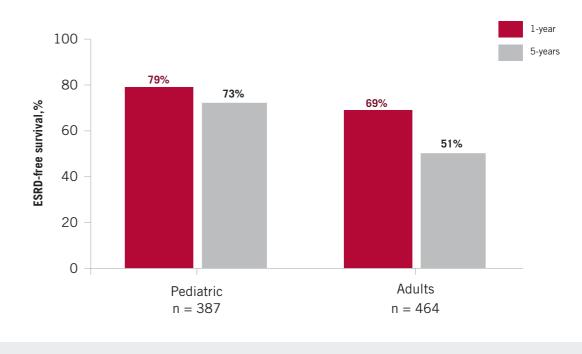
Patients with atypical-HUS are at ongoing risk of systemic, life-threatening, and sudden complications^{1,3,4,27}

Atypical-HUS patients can show involvement in more than one organ system^{7,16,28-30,a}



The risk of TMA is ongoing, unpredictable, and life-threatening in patients with atypical-HUS^{1,12,16,27}

Pediatric patients have a lower risk of developing ESRD compared with adult patients (adjusted hazard ratio 0.55 [95% CI, 0.41–0.73]);sex, race, family history of atypical-HUS, time from initial presentation to diagnosis, and potential complement-activating conditions were not associated with ESRD risk¹⁶



Study description: Global, observational study of atypical-HUS including both retrospective and prospective enrollment. At time of data cutoff (November 30, 2015), 851 patients were enrolled.

Cl, confidence interval; eGFR, estimated glomerular filtration rate; ESRD, end-stage renal disease; Gl, gastrointestinal; HUS, hemolytic uremic syndrome; TMA, thrombotic microangiopathy.

Approximately 50% of adult patients with atypical-HUS are at risk for ESRD and death^{16,a}

^aThe organ-specific symptoms associated with atypical-HUS are reported from published literature and are not limited to only those listed above.

8

^d5-year ESRD-free survival.

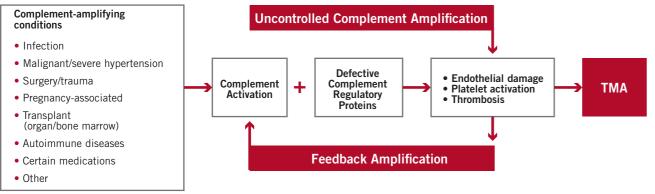
Chronic Risk of ESRD

Chronic Risk of ESRD

Complement-Amplifying Conditions May Unmask Atypical-HUS

Complement-amplifying conditions place patients with atypical-HUS at high risk for TMA manifestations^{2,7}

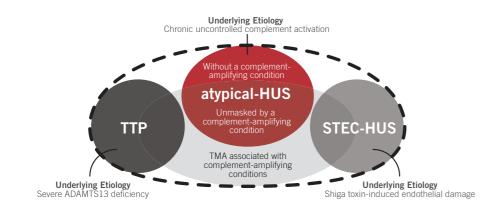
Ongoing complement activity⁸



Adapted from Laurence J, et al. Clin Adv Hematol Oncol. 2016;14:1-15

• 70% of patients (191/273) with atypical-HUS presented their first clinical manifestations while experiencing a complement-amplifying condition¹²

TMAs associated with complement-amplifying conditions³



NOTE: The diagram is for illustrative purposes only; disease areas are not drawn to proportional scale and are not meant to reflect relative incidence

ADAMTS13, a disintegrin and metalloproteinase with a thrombospondin type 1 motif member 13; HUS, hemolytic uremic syndrome; STEC-HUS, Shiga toxin-producing E. coli-hemolytic uremic syndrome; TMA, thrombotic microangiopathy; TTP, thrombotic thrombocytopenic purpura.

If the signs and symptoms of TMA do not rapidly resolve in response to management of the triggering condition, evaluate for atypical-HUS by assessing for STEC-HUS and ADAMTS13 deficiency

Persistent TMA May Suggest Atypical-HUS

Persistent TMA may suggest atypical-HUS

Examples of complement-amplifying conditions include

Malignant hypertension (MHT)/severe hypertension

 Persistent TMA despite hypertension management should raise suspicion for atypical-HUS unmasked by MHT or severe hypertension³²

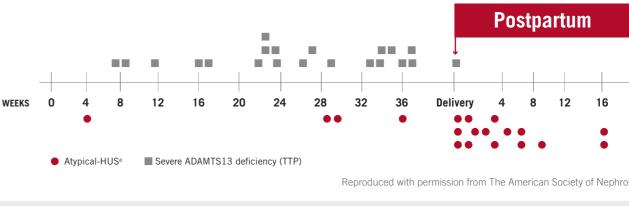
Infection

Suspect atypical-HUS if symptoms of TMA persist after treatment of infection¹²

Pregnancy

• A high clinical suspicion for atypical-HUS should be raised if signs of TMA present postpartum or persist more than 48 hours after delivery/termination^{33,34}

Atypical-HUS is commonly unmasked in the postpartum setting³⁵



Study Description: A retrospective analysis of 19 patients who experienced clinical manifestations of atypical-HUS, and 21 patients who experienced clinical manifestations of TTP, during pregnancy or postpartum. Patients with aHUS were referred between 2000 and 2008 to the laboratory of immunology at Hôpital Européen Georges-Pompidou (Paris, France), a reference center for the evaluation of complement disorders. A diagnosis of atypical-HUS was defined by the coexistence of hemolytic anemia, thrombocytopenia, and acute renal failure. Patients with ADAMTS13 deficiency-associated TTP were identified from previously published cases.

^aDefined as complement dysregulation TMA.

Certain medications used in transplant

Suspect atypical-HUS if TMA symptoms persist after the associated drug (such as calcineurin inhibitors) has been discontinued³

Reproduced with permission from The American Society of Nephrology.

Persistent TMA May Suggest Atypical-HUS

Potential exposure to any complement-amplifying condition may lead to TMA manifestations in patients with atypical-HUS^{4,9,12,16}

Limitations of Plasma Exchange/ **Plasma Infusion in Atypical-HUS**

Examples of factors that may increase risk for TMA manifestations in patients with atypical-HUS include^{7,36,a}



History of renal transplant 1,37

- The risk of atypical-HUS recurrence following transplantation has been reported to range from 20% to more than 80% depending on the presence of a specific genetic mutation³⁸
- Risk for TMA is also deemed high in patients without a genetic mutation who have received a renal transplant^{7,37}
- Risk for allograft loss is high in patients with atypical-HUS^{12,17,39}



Age of patient⁶

Children are considered to be at high risk for recurrent TMA due to the frequency of common events that lead to complement activation in this age group⁶



Identified genetic mutation^{12,27,40}

Clinical studies show that mutations in complement genes are associated with higher risk of TMA^{12,27,40}



Pregnancy/Postpartum^{9,35,41,42}

 Patients with atypical-HUS are at risk for TMA manifestations during the pregnancy/postpartum period^{9,41,42} due to factors such as infection, hemorrhage, and HELLP syndrome^{9,41}



Clinical history or family history of TMA^{36,40,43,44}

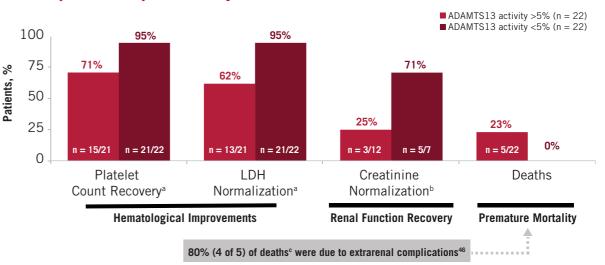
- Multiple TMA manifestations suggests high risk for subsequent TMA in the presence of complement-amplifying conditions^{36,40}
- Patients with family history of disease have a higher rate of disease progression; rate of ESRD has been reported to be between 50-80%12,44

^aThis is not a comprehensive list, but is intended to provide examples of factors that may increase risk for TMA. Anything that amplifies complement is a risk factor for TMA.4,9,12,35

ESRD, end-stage renal disease; HELLP, hemolysis, elevated liver enzymes, low platelet count; HUS, hemolytic uremic syndrome; LDH, lactate dehydrogenase; PE/PI, plasma exchange/plasma infusion; TMA, thrombotic microangiopathy vFW, von Willebrand factor.

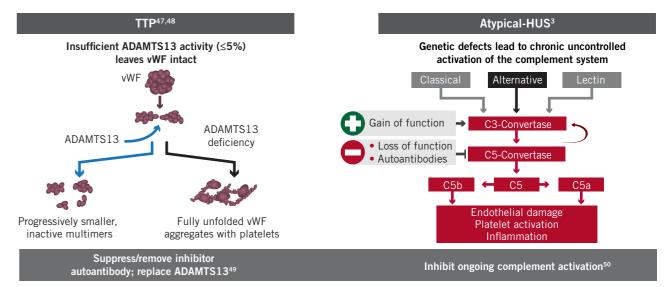
Mortality outcomes differed in patients receiving PE with and without severe ADAMTS13 deficiency⁴⁵

Rates of Renal Recovery and Premature Mortality in Patients With Severe or Nonsevere ADAMTS13 Activity Undergoing PE/PI for the Treatment of TMA⁴⁵ (Follow-up Period up to 21 Days)



Non-ST-segment elevation MI/aspiration pneumonia, non-ST-segment elevation MI/abdominal abscess, multiorgan failure, respiratory failure, sepsis.46

TTP and atypical-HUS are driven by different pathophysiologic processes and have different management goals



Patients with atypical-HUS remain at risk of impaired renal function and death, regardless of hematologic improvement after PE/PI⁸

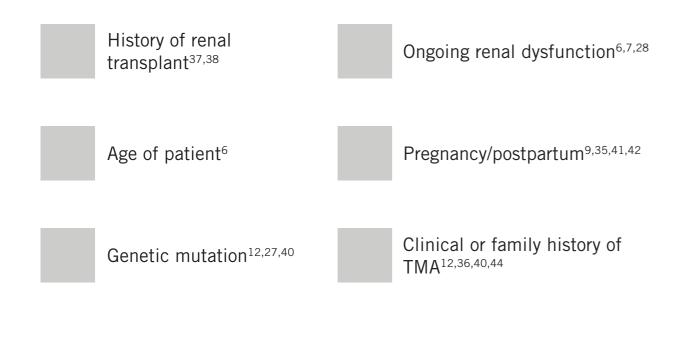
Limitations of PE/PI in Atypical-HUS

^aOf patients with available data. ^bOf patients with abnormal creatinine level at baseline

Factors to Consider for Long-term Management of Atypical-HUS

When developing a management plan for an atypical-HUS patient, risk assessment for TMA manifestations, the patient's unique clinical situation, and an understanding of the unpredictable nature of atypical-HUS should be considered^{36,40,51}

Factors that may increase risk for TMA complications in patients with atypical-HUS^a



^aThis is not a comprehensive list, but is intended to provide examples of factors that may increase risk for TMA. Anything that amplifies complement is a risk factor for TMA.^{9,12,16,40}

References

1. Legendre CM, et al. N Engl J Med. 2013;368:2169-2181. 2. Noris M, et al. Nat Rev Nephrol. 2012;8:622-633 3. Azoulay E, et al. CHEST. 2017;152:424-434. 4. Goodship THJ, et al. Kidney Int. 2017;91:539-551 5. Kato H, et al. Pediatr Int. 2016;58:549-555. 6. Loirat C, et al. Pediatr Nephrol. 2016;31:15-39. 7. Campistol JM. et al. Nefrologia. 2015:35:421-447. 8. Laurence J et al. Clin Adv Hematol Oncol. 2016;14:2-15. 9. Asif A, et al. J Nephrol. 2017;30:347-362. 10. Jokiranta TS. Blood. 2017: 129:2847-2856 11. Maga TK, et al. Hum Mutat. 2010;31:E1445-E1460. 12. Noris M, et al. Clin J Am Soc Nephrol. 2010;5:1844-1859. 13. Bu F, et al. J Am Soc Nephrol. 2014;25:55-64. 14. George JN, et al. N Engl J Med. 2014;371:654-666. 15. Structural Immunology Group. Database of complement gene variants. http:// www.complement-db.org/advance_search_result s.php?dosearch=1&source=lab&dosearc h=1&source=lab&genead%5B%5 D=all&reference=&condition%5B%5D=aHUS&c utoff=0.01. Accessed November 15, 2018. 16. Schaefer F, et al. Kidney Int. 2018;94:408-418. 17. Caprioli J, et al. Blood. 2006;108:1267-1279. 18. Sellier-Leclerc AL, et al. J Am Soc Nephrol. 2007;18:2392-2400. 19. Barbour T. et al. Nephrol Dial Transplant. 2012:27:2673-2685 20. Fang CJ, et al. Br J Haematol. 2008;143:336-348. 21. Loirat C, Frémeaux-Bacchi V. Orphanet J Rare Dis. 2011;6:60. 22. Gastoldi S,et al. Immunobiology. 2012;217:1145-1146. 23. Salant D.J. J Am Soc Nephrol. 2011:22:7-9. 24. Renner B,et al. J Am Soc Nephrol. 2013;24:1849-1862. 25. Nester CM, Thomas CP. Hematology Am Soc Hematol Educ Program. 2012;617-625. 26. Cofiell R.et al. Blood. 2015:125:3253-62. 27. Fremeaux-Bacchi V, et al. Clin J Am Soc Nephrol. 2013;8:554-562 28. Jamme M, et al. PLoS One. 2017;12:e0177894. 29. Hofer J. et al. Front Pediatr. 2014:2:1-16. 30. Krishnappa V, et al. Ther Apher Dial. 2018;22:178-188. 31. Noris M, et al. Nat Rev Nephrol. 2014;10:174-180. 32. Zhang B, et al. Hypertens Res. 2008;31:479-483. 33. Dobyne A, et al. Med J Obstet Gynecol. 2015;3:1064. 34. Sibai BM, et al. Am J Obstet Gynecol. 1993;169:1000-1006. 35. Fakhouri F, et al. J Am Soc Nephrol. 2010;21:859-867. 36. Macia M, et al. Clin Kid J. 2017;10:310-319. 37. Zuber J, et al. Transplant Rev. 2013;27:117-125. 38 Abbas F et al. World J Transplant, 2018;8:122-141 39. Kim S et al. J Korean Med Sci. 2018;33:e4. 40. Fakhouri F, et al. Clin J Am Soc Nephrol. 2017;12:50-59. 41. Bruel A, et al. Clin J Am Soc Nephrol. 2017;12: 1237-1247 42. Huerta A, et al. Kidney Int. 2018;93:450-459. 43. Williams LA, Marques MB. Am J Clin Pathol. 2016;145:158-165. 44. Noris M, Remuzzi G. N Engl J Med. 2009;361:1676-1687. 45. Pishko AM, Arepally GM. Blood. 2014;124:4192. 46. Pishko AM, Arepally GM. Presented at: 56th American Society of Hematology Annual Meeting and Exposition; December 6-9, 2014; San Francisco, CA. Poster 4192. 47. Tsai HM. Int J Hematol. 2010:91:1-19. 48. Sadler JE. Blood. 2008;112:11-18. 49. Tsai HM. Am J Med. 2013;126:200-209

- 50. Laurence J. Clin Adv Hematol Oncol. 2012;10:1-12.
- 51. Olson SR, et al. Am J Nephrol. 2018;48:96-107.





Atypical-HUS is a chronic, unpredictable, genetic disease that can be life threatening^{1,3,4}

Patients with TMA require rapid differential diagnosis to make appropriate management decisions³

Persistent TMA in patients with complement-amplifying conditions may suggest atypical-HUS⁸

The role of PE/PI is limited in management of atypical-HUS⁸

HUS, hemolytic uremic syndrome; PE/PI: plasma exchange/plasma infusion; TMA, thrombotic microangiopathy.

© 2019, Alexion Pharmaceuticals, Inc. All rights reserved. US/UNB-a/0051 Alexion® is a registered trademark of Alexion Pharmaceuticals, Inc.

