# **Hemophilia A, Hemophilia B, Congenital von Willebrand Disease, and Acquired von Willebrand Syndrome**

Shiu-Ki Rocky Hui

## **Hemophilia A**

Hemophilia A is a X-linked disorder due to congenital deficiency of factor VIII (FVIII). The frequency of Hemophilia is estimated to be approximately 1/5000 male births across ethnic group. The degree of bleeding severity with rare exception is directly related to patient's baseline FVIII level and can range from mild to life-threatening  $[3]$ . Although Hemophilia A affects mostly male, in rare occasion the disease can manifest in female via chromosome lionization [4– [6](#page-5-0). In addition, approximately  $30\%$  of Kemophilia A can present as de-novo mutation. Therefore, diagnosis of Hemophilia A should be considered even in the absence of a strong family history  $[7-9]$ .

# **Factor VIII Protein**

 Unlike most other plasma coagulation proteins, which are produced by hepatocytes, FVIII is mainly produced by liver endothelial cells  $[10, 11]$ . This is likely the reason for elevated FVIII even under liver failure.

 Factor VIII protein is produced as a six-domain protein, A1-A2-B-A3-C1-C2  $[12-14]$ . In the Golgi compartment, FVIII is cleaved within the B domain to form the mature FVIII heterodimer that consists of the 200 kDa A1-A2-B heavy chain and the 80 kDa A3-C1-C2 light chain  $[15]$ . Once FVIII is released into circulation, it binds to von Willebrand factor (VWF) which serves to increase FVIII survival and regulate FVIII activity. Upon activation by thrombin, FVIIIa serves as a co-factor for Factor IXa in the

S.-K.R. Hui, M.D.  $(\boxtimes)$ 

Departments of Pathology & Immunology and Pediatrics, Division of Transfusion Medicine & Coagulation, Texas Children's Hospital, 6621 Fannin Street, Suite F0560.14, Houston, TX 77030, USA e-mail: [sxhui@txch.org](mailto:sxhui@txch.org)

activation of factor X to factor Xa and ultimately generation of thrombin  $[15-18]$ . Hemophilia A has been linked to a variety of well-known molecular mutations including inversions, large deletion, frameshift, nonsense, and missense defects  $[19, 20]$ . The severity of the disease can often be predicted by the site and type of mutation in the FVIII gene  $[21 - 24]$ .

# **Diagnosis of Hemophilia A**

 The diagnosis of Hemophilia A begins in recognizing the X-linked inheritance pattern of an unexplained and isolated prolongation of PTT that is corrected in mixing study  $[25]$ . Subsequent laboratory workup should then identify a FVIII deficiency. However, it is important to note that an initial finding of an isolated prolonged PTT with mixing study correction does not immediately imply Hemophilia A [26]. Other PTT pathway factor deficiencies such as Factor IX (FIX) and Factor XI (FXI) deficiencies or even non-clinically significant Factor XII deficiencies will present with the same initial laboratory finding  $[27]$ . Therefore, it is important to order FIX and FXI along with FVIII to rule out other potential congenital bleeding disorders. Specifically important is to consider the von Willebrand disease (VWD), when considering the diagnosis of Hemophilia A, as carrier defect (Type  $2N$ ) or deficiency (Type 1 and 3) of VWF will result in decrease of FVIII. The diagnosis of these specific VWD will be discussed later in this chapter.

 Severity of Hemophilia A is directly related to the degree of deficiency of FVIII activity as measured by one-stage assay, Mild Hemophilia is defined by FVIII activity between 5 snd 40 %, moderate disease is activity 1–5 %, while severe Hemophilia is below 1%. Although FVIII activity via onestage assay in general can predict bleeding phenotype, in very rare situation, there can be discrepant clinical bleeding symptoms with FVIII activity via one-stage assay. In such scenario, a two-stage FVIII assay may be used to better define patient's disease [28, [29](#page-6-0)].

 **7**

#### **Hemophilia B**

 Like Hemophilia A, Hemophilia B is also a X-linked bleeding disorder affecting around 1 in 25,000 male births. In Hemophilia B, factor IX (FIX) is deficient which results in lifelong bleeding symptoms. The severity of bleeding is directly related to the degree of FIX deficiency. Approximately, one third of all cases arise from spontaneous mutation  $[30]$ . Therefore, like Hemophilia A, a lack of family history does not exclude the diagnosis.

# **Factor IX Protein**

FIX protein is a 57 kDa multi-domain protein produced in the liver by hepatocyte. FIX is a vitamin K-dependent protein, requiring gamma-carboxyglutamation to be fully functional. FIX can be activated into FIXa via the intrinsic pathway by FXIa or the extrinsic pathway by tissue factor and FVIIa [31]. FIXa in the presence of calcium, FVIIIa, and phospholipids in turn activates FX into FXa and ultimately thrombin. Hemophilia B has been linked to a number of mutations, especially frameshift, missense, or nonsense mutations within the CpG dinucleotide mutation hotspot. Large and short deletions and insertions complete the genetic mutation profile for Hemophilia B [30].

#### **Diagnosis**

A deficiency in FIX may indicate vitamin K deficiency instead of Hemophilia B, especially in neonates [32]. Since FVII is also a vitamin K-dependent factor, an isolated prolonged PTT without prolongation of PT makes Hemophilia B likely as half-life of FVII is shorter than FIX. Like FVIII in Hemophilia A, severity of Hemophilia B is directly related to the degree of deficiency of FIX activity. Mild Hemophilia B is defined by FIX activity between 5 and 40 %, moderate disease is activity 1–5 %, while severe Hemophilia B is below 1 %.

#### **Congenital von Willebrand Disease**

 VWD remains the most common congenital bleeding disease worldwide across all ethnic groups. Unlike Hemophilia, VWD is an autosomal disorder; therefore it affects both sexes equally and this inheritance pattern helps to distinguish it from Hemophilia A. Bleeding characteristics and severity are greatly affected by its subtypes and can range from joint and muscle bleeding (Type 3) to menorrhagia to mild oral and mucosal bleeding [33, [34](#page-6-0)]. Due to the varia-

tion of clinical presentation, a complete and accurate laboratory workup is important for the subtyping of VWD [35]. The laboratory workup for VWD will be discussed in detail in Chapter XX.

## **von Willebrand Factor**

 Unlike most other coagulation proteins, VWF is not produced by the liver which also helps to explain elevated FVIII and increased thrombotic risk in the setting of liver dysfunction. VWF is produced by both megakarocytes and endothelial cells (Fig. 7.1) as a 2813 amino acid long prepropeptide (Fig. [7.2](#page-2-0) ) in the endoplasmic reticulum, which is then dimerized into 800kD dimers. These dimers are then polymerized into mature VWF multimers up to 20,000kD in length and VWF propeptide dimmers (VWF:pp) are cleaved from the mature multimers as they travel through the golgi. Finally, both the mature VWF and VWF:pp are packaged and stored in Weibel–Palade body of endothelial cells or alpha granules of platelets. Upon activation, mature VWF and VWD:pp are released from storage into circulation; once released, VWF multimers are cleaved at specific site in the A2 domain into multimers of variable sizes by a metalloprotease, ADAMTS13 [36–38]. Under normal physiological condition, VWF exists as large, intermediate and low molecular weight multimers in a balanced distribution. However, when this normal size distribution is disturbed, it will result in disease conditions such as thrombotic thrombocytopenic purpura when there is ultra large multimers [39] or bleeding when there is absence or decrease in large multimers  $[40]$ .

 Circulating VWF plays an important role in both primary and secondary hemostasis. In primary hemostasis, VWF serves to support platelet adhesion to the site of vascular injury via binding to sub-endothelial collagen and to glycoprotein Ib-V-X complex (GPIb) on platelet surface. This interaction is important in recruiting and activating platelets at site of vascular injury  $[41]$ . In terms of secondary hemostasis, VWF serves as a carrier protein for FVIII, which both protects FVIII from proteolysis and localizes FVIII to platelet surface  $[42, 43]$  $[42, 43]$  $[42, 43]$ .

#### **von Willebrand Disease**

 As discussed previously, VWD can present with widely different bleeding phenotypes depending on the underlying pathophysiology. In general, VWD (Table [7.1](#page-3-0)) can be broadly divided into two types of VWF defects, quantitative (type 1 and type 3) and qualitative (type 2). It is important to

<span id="page-2-0"></span>

 **Fig. 7.1** demonstrates the production of VWF multimers in endothelial cells. Matured VWF and propeptide are stored within Weibel-Palade bodies ready for release upon activations. From

Haberichter SL, Regulated release of VWF and FVII and the biologic implications, Pediatr Blood Cancer, 2006 May 1: 46(5):547–53



**Fig. 7.2** showed the various known VWF domains and their respective contributions to the VWF function

Condition	VWF:Act	VWF:Ag	<b>FVIII</b>	VWF:Act/Ag	VWF:MA
Type 1	$30$	$30$	$L \text{ to } N$	>0.7	Normal but light
Type 2A	$30$	$<30-200$	$L \text{ to } N$	< 0.7	Missing large
Type 2B/PT-VWD	$30$	$<30-200$	$L \text{ to } N$	< 0.7	multimer
Type 2M	$30$	$<30-200$	$L \text{ to } N$	< 0.7	Normal
Type 2N	$30 - 200$	$30 - 200$	Significantly lower	>0.7	Normal
Type 3	$\leq$ 3	$\leq$ 3	<10	$\overline{\phantom{0}}$	-
"Low VWF"	$30 - 50$	$30 - 50$	Normal	>0.7	Normal
Normal	$50 - 200$	$50 - 200$	Normal	>0.7	Normal

<span id="page-3-0"></span>**Table 7.1** Various VWD subtypes and their expected laboratory findings

distinguish the various subtypes of VWD via an algorithmic laboratory approach, as it can greatly impact the management of the patient. Various laboratory workups and algorithm is found in details in Chapter \*\*.

## **Type 1 VWD**

Type 1 VWD account for the majority of VWD  $(80\%)$ . It is a quantitative defect defined as either VWF antigen (VWF:Ag) or VWF activity (VWF:Act) between 1 and 30 % without any observable VWF function defects that VWF activity to antigen ratio (VWF:Act/Ag) should be  $>0.5-0.7$  [44]. In addition, VWF multimer analysis (VWF:MA) should show a normal size distribution with decreased intensity. Mechanism for type 1 VWD is due to decreased synthesis of VWF; however, type 1 variants (type 1C and type 1 Vicenza) have been shown to have increased clearance and decreased VWF half-life. It is important to rule out such type 1 variants as desmopressin treatment will not be an effective treatment as the therapy yield only short-lasting effect  $[45]$ . In both type 1C VWD and type 1 Vicenza, desmopressin challenge is expected to show good 1 h post-administration response with a decreased 4 h post-administration response [46]. Compared to other type 1 VWD, VWF propeptide to antigen ratio (VWF:pp/Ag) is increased, which is the defining characteristic of these type 1 variants  $[47]$ . It is important to note that FVIII in type 1 VWD is proportionally decreased as VWF is a carrier protein for FVIII, thus VWD workup should be performed in initial diagnosis of Hemophilia A. Furthermore, Hemophilia A can coexist with other subtypes of VWD. Lastly, laboratory diagnosis of VWD is proven challenging as VWF is an acute phase protein. Since the level can be increased several folds from baseline, a one-time normal VWF:Ag and VWF:Act cannot definitively rule out type 1 VWD [48]. Furthermore, FVIII is not an effective marker for acute phase as its level is directly affected by VWF:Ag level. Concurrent fibrinogen level or C reactive protein level may be used as potential acute phase markers, but neither of them has been universally validated. Until a better marker can be established, the most effective method to distinguish acute phase from baseline study remains to be repeat testing.

## **Low VWF**

 It is important to discuss "Low VWF" in the discussion of quantitative type 1 and type 3 VWD. As normal VWF:Ag and VWF: Act level is usually defined as  $>50\%$  and type 1 VWD is <30 %, the gray-zone area between 30-50 % can be difficult to define [44]. This "in-between" VWF:Ag and VWF:Act can fall within type 1 VWD in European region as type 1 VWD is defined as  $\lt 45\%$  [49]. However, in the United States, type 1 VWD is strictly reserved for patient with VWF:Ag or VWF:Act <30 % without functional defects. Hence, individuals with repeat VWF level between 30 and 50 % should be considered as having "low VWF" and not VWD. Of note, blood group O individuals are more likely to have "low VWF" than other blood groups  $[50]$ , which may be related to post-translational modification  $[51]$ . Individuals with "low VWF" should be made aware of increased risk for bleeding, but should not be considered as having true VWD.

# **Type 3 VWD**

Type 3 VWD accounts for less than 1% of VWD. It is a severe quantitative defect defined as absence  $\left($ <1 %) of both VWF:Ag and VWF:Act [44]. As FVIII level can fall within moderate Hemophilia A range, it is important to rule out type 3 VWD.

#### **Type 2A VWD**

Type 2A VWD accounts for approximately 10% of all VWD. It is characterized by the absence of both high and intermediate molecular weight VWF secondary to decreased synthesis or increased proteolysis by ADAMTS13 [44]. Therefore, laboratory workup demonstrates a qualitative defect of decreased VWF:Act but relatively normal VWF:Ag, which results in decreased VWF:Act/Ag ratio at <0.5-0.7. Of note, unlike type 2B VWD, mild thrombocytopenia is not an expected finding. Type 2A VWD is often considered as a diagnosis of exclusion.

#### **Type 2B VWD**

 Type 2B VWD accounts for approximately 3-5 % of all VWD. Although its laboratory finding is similar to type 2A VWD with decreased VWF:Act/Ag ratio and loss of high molecular weight multimer, the mechanism for disease is entirely different  $[44]$ . Type 2B VWD is due to gain of function mutation in the platelet GPIb binding to A1 domain of the VWF protein, which results in increased and spontaneous binding of VWF to platelets without shear stress [52]. This abnormal interaction results in loss of both high molecular weight VWF and platelets, which explains the pathognomonic findings of thrombocytopenia in type 2B VWD. This gain of function mutation makes desmopressin a contraindication for type 2B VWD as it may result in thrombotic complications. Therefore, it is important to distinguish type 2B from other type 2 VWD; ristocetin-induced platelet aggregation study (RIPA) is only abnormal in gain-of-function VWD which includes Type 2B VWD.

#### **Pseudo-VWD/Platelet-type VWD**

 As in type 2B VWD, the pathogenesis of platelet-type VWD (PT-VWD) is due to abnormal spontaneous interaction between platelet GPIb and VWF [44]. However, in contrast to type 2B VWD, the gain of function mutation is in the platelet GPIb receptor [53]. Overall, initial laboratory workup is indistinguishable from type 2B VWD, including decreased VWF:Act/Ag, loss of high molecular VWF, thrombocytopenia, and even abnormal RIPA. Specialized laboratory test, 2B binding assay can be used to differentiate PT-VWD from type 2B VWD. As in type 2B VWD, desompressin is contraindicated for treatment of PT-VWD.

# **Type 2M VWD**

 Type 2M VWD accounts for only 1–2 % of VWD. The initial laboratory workup, similar to type 2A, 2B or PT-VWD, showed decreased VWF:Act/Ag [44]. Like type 2B VWD, pathogenesis for type 2M VWD also lies in the A1 domain of VWF, but it is a loss-of-function mutation where the interaction between platelet GPIb receptor and VWF is decreased [54]. Therefore, multimer analysis for type 2M VWD is normal and does not demonstrate loss of high or intermediate molecular weight VWF. It is the presence of normal multimer distribution with decreased VWF:Act/Ag that makes up the defining laboratory characteristic of type 2M VWD.

#### **Type 2N VWD**

 Type 2N VWD is qualitative VWF disorder that accounts for  $1-2\%$  of all VWD [44]. However, its functional defect lies not in VWF function as a coagulation protein, but its FVIII carrier function. Mutations within the D' and D3 domain of VWF molecule render the binding of VWF to FVIII defective [55]. As the coagulation function of VWF is unaffected, the VWF laboratory workup is unremarkable at first glance; normal VWF:Ag, VWF:Act, VWF:Act/Ag, and even normal multimer analysis. However, FVIII activity can be decreased to as low as  $5-15\%$ , making 2N VWD sometimes difficult to differentiate from Hemophilia A. The inheritance pattern of 2N VWD is autosomal in contrast to X-linked in Hemophilia A. FVIII binding assay (Discuss in Chapter \*\*) can be used to distinguish type 2N VWD from Hemophilia A. It is important to note that like other VWD subtypes, 2N VWD can coexist in patients with Hemophilia A. Therefore, concurrent 2N VWD should always be considered and ruled out as it can affect patient's response to recombinant FVIII infusion.

## **Acquired von Willebrand Syndrome**

 Acquired von Willebrand Syndrome (aVWS) is a collection of acquired bleeding disorders (Table [7.2 \)](#page-5-0), secondary of loss of VWF quantitative or qualitative functions [56]. Dozens of diseases have been associated with aVWS; nonetheless, laboratory findings often mimic subtype of congenital VWD, especially type 2A VWD with decreased VWF:Act/Ag and loss of high to intermediate molecular weight VWF. The loss of high molecular weight VWF can be secondary to either pathological high shear stress as in aortic stenosis [57], presence of autoantibodies against VWF [58], or even direct absorption by tumor cells [59]. Less commonly, aVWS may result from decreased overall VWF production as opposed to selective loss of high molecular weight VWF as in the case of hypothyroidism  $[60]$ . Bleeding diathesis of aVWS may vary, but bleeding symptoms and VWF laboratory abnormalities usually resolve upon resolution of underlying disorders.

# **Management of Hemophilia A, Hemophilia B, and von Willebrand Disease for Invasive Procedure, Surgery, and Pregnancy**

 The management of patients with Hemophilia A, B, and VWD can be complex; however, there have been established recommended guidelines (Table 7.3) that can provide some important standard of care guidance in managing these patients

<span id="page-5-0"></span> **Table 7.2** Shows the various disorders that have been reported to be associated with aVWS

Underlying disorders	Previous literature, ISTH-SSC and German registry $1968-2011$ ( $n=1292$ )		
Cardiovascular	414	$32\%$	
Aortic stenosis	201	$16\%$	
Cardiac assist device	110	9%	
AV septal defects	21	$2\%$	
Myeloproliferative	350	27%	
Essential thrombocythemia	212	$16\%$	
Polycythemia vera	88	7%	
CML and myelofibrosis	64	$5\%$	
Lymphoproliferative	321	25%	
<b>MGUS</b>	193	$15\%$	
MM and WMg	80	6%	
NHL, HCL, and ALCL	28	$2\%$	
Systemic diseases (hepatitis C, cirrhosis, hypo-thyroid, hemoglobinopathies, uremia, diabetes)	87	7%	
Drugs (valproate, hydroxystarch, etc.)	40	3%	
Neoplasia	32	$2\%$	
<i>Immune</i>	21	$2\%$	

**Table 7.3** Published recommendations for peri-operative management of Hemophilia A, B, and VWD patients



 around time of procedures, surgeries, or deliveries. It is important to note that factor concentrates should be used in place of plasma products as replacement of choice since the concentration is much higher and infectious risk is significantly less. DDAVP may be used in patients with mild Hemophilia A, mild VWD, or Hemophilia A carrier in place of factor replacements; however, a trial should be performed to ensure effectiveness prior to use in surgerical settings. Antifibrinolytic may be used in conjunction with standard factor replacement [61]; however, this practice has not been well-standardized beyond dental procedure, but should be considered if risk is high or if replacement therapy alone is ineffective.

## **References**

 1. Schneppenheim R, Budde U, Krey S, Drewke E, Bergmann F, Lechler E, Oldenburg J, Schwaab R. Results of a screening for von Willebrand disease type 2N in patients with suspected haemophilia A or von Willebrand disease type 1. Thromb Haemost. 1996; 76(4):598–602.

- 2. Lindsay H, Bergstrom K, Srivaths L. Co-inheritance of mild Hemophilia A and heterozygosity for type 2N von Willebrand disease: a diagnostic and therapeutic challenge. Pediatr Blood Cancer. 2014;61(10):1888–90.
- 3. Chai-Adisaksopha C, Hillis C, Thabane L, Iorio A. A systematic review of definitions and reporting of bleeding outcome measures in haemophilia. Haemophilia. 2015;21(6):731–5.
- 4. Di Michele DM, Gibb C, Lefkowitz JM, Ni Q, Gerber LM, Ganguly A. Severe and moderate haemophilia A and B in US females. Haemophilia. 2014;20(2):e136–43.
- 5. Radic CP, Rossetti LC, Abelleyro MM, Tetzlaff T, Candela M, Neme D, Sciuccati G, Bonduel M, Medina-Acosta E, Larripa IB, de Tezanos PM, De Brasi CD. Phenotype-genotype correlations in Hemophilia A carriers are consistent with the binary role of the phase between F8 and X-chromosome inactivation. J Thromb Haemost. 2015;13(4):530–9. doi[:10.1111/jth.12854.](http://dx.doi.org/10.1111/jth.12854) Epub 2015 Mar 14.
- 6. Renault NK, Dyack S, Dobson MJ, Costa T, Lam WL, Greer WL. Heritable skewed X-chromosome inactivation leads to haemophilia A expression in heterozygous females. Eur J Hum Genet. 2007;15(6):628–37.
- <span id="page-6-0"></span> 7. Kentsis A, Anewalt R, Ganguly A, Allen JB, Neufeld EJ. Discordant haemophilia A in male siblings due to a de novo mutation on a familial missense mutant allele. Haemophilia. 2009;15(4):971–2.
- 8. Kapsimali Z, Pavlova A, Pergantou H, Adamtziki E, Oldenburg J, Platokouki H. Two de novo factor VIII gene mutations in the family of an isolated severe haemophilia A patient. Haemophilia. 2012;18(1):e3–4.
- 9. Castaldo G, D'Argenio V, Nardiello P, Zarrilli F, Sanna V, Rocino A, Coppola A, Di Minno G, Salvatore F. Haemophilia A: molecular insights. Clin Chem Lab Med. 2007;45(4):450–61.
- 10. Stel HV, van der Kwast TH, Veerman EC. Detection of factor VIII/coagulant antigen in human liver tissue. Nature. 1983; 303(5917):530–2.
- 11. Zelechowska MG, van Mourik JA, Brodniewicz-Proba T. Ultrastructural localization of factor VIII procoagulant antigen in human liver hepatocytes. Nature. 1985;317(6039):729–30.
- 12. Choi SJ, Jang KJ, Lim JA, Kim HS. Human coagulation factor VIII domain-specific recombinant polypeptide expression. Blood Res. 2015;50(2):103–8.
- 13. Venkateswarlu D. Structural investigation of zymogenic and activated forms of human blood coagulation factor VIII: a computational molecular dynamics study. BMC Struct Biol. 2010;10:7.
- 14. Fang H, Wang L, Wang H. The protein structure and effect of factor VIII. Thromb Res. 2007;119(1):1–13.
- 15. Thompson AR. Structure and function of the factor VIII gene and protein. Semin Thromb Hemost. 2003;29(1):11–22.
- 16. Fay PJ. Factor VIII structure and function. Int J Hematol. 2006;83(2):103–8.
- 17. Regan LM, Fay PJ. Cleavage of factor VIII light chain is required for maximal generation of factor VIIIa activity. J Biol Chem. 1995;270(15):8546–52.
- 18. Fay PJ, Haidaris PJ, Smudzin TM. Human factor VIIIa subunit structure. Reconstruction of factor VIIIa from the isolated A1/A3-C1-C2 dimer and A2 subunit. J Biol Chem. 1991;266(14):8957–62.
- 19. Rydz N, Leggo J, Tinlin S, James P, Lillicrap D. The Canadian "National Program for Hemophilia mutation testing" database: a ten-year review. Am J Hematol. 2013;88(12):1030–4.
- 20. Miller CH, Benson J, Ellingsen D, Driggers J, Payne A, Kelly FM, Soucie JM. Craig Hooper W; Hemophilia Inhibitor Research Study Investigators. F8 and F9 mutations in US haemophilia patients: correlation with history of inhibitor and race/ethnicity. Haemophilia. 2012;18(3):375–82.
- 21. Gouw SC, Van Der Bom JG, Van Den Berg HM, Zewald RA, Ploos Van Amstel JK, Mauser-Bunschoten EP. Influence of the type of F8 gene mutation on inhibitor development in a single centre cohort of severe haemophilia A patients. Haemophilia. 2011;17(2):275–81.
- 22. Oldenburg J, El-Maarri O, Schwaab R. Inhibitor development in correlation to factor VIII genotypes. Haemophilia. 2002;8 Suppl  $2:23-9$ .
- 23. Goodeve AC, Williams I, Bray GL, Peake IR. Relationship between factor VIII mutation type and inhibitor development in a cohort of previously untreated patients treated with recombinant factor VIII (Recombinate). Recombinate PUP Study Group. Thromb Haemost. 2000;83(6):844–8.
- 24. Goodeve AC, Peake IR. The molecular basis of Hemophilia A: genotype- phenotype relationships and inhibitor development. Semin Thromb Hemost. 2003;29(1):23–30.
- 25. Favaloro EJ, Meijer P, Jennings I, Sioufi J, Bonar RA, Kitchen DP, Kershaw G, Lippi G. Problems and solutions in laboratory testing for Hemophilia. Semin Thromb Hemost. 2013;39(7):816–33.
- 26. Bolton-Maggs PH, Favaloro EJ, Hillarp A, Jennings I, Kohler HP. Difficulties and pitfalls in the laboratory diagnosis of bleeding disorders. Haemophilia. 2012;18 Suppl 4:66–72.
- 27. Jennings I, Kitchen DP, Kitchen S, Woods TA, Walker ID. Investigation of a prolonged APTT. Different approaches taken

by laboratories to achieve the same diagnosis. Int J Lab Hematol. 2013;35(2):177–82.

- 28. Armstrong E, Hillarp A. Assay discrepancy in mild haemophilia A. Eur J Haematol Suppl. 2014;76:48–50.
- 29. Duncan EM, Rodgers SE, McRae SJ. Diagnostic testing for mild Hemophilia a in patients with discrepant one-stage, two-stage, and chromogenic factor VIII:C assays. Semin Thromb Hemost. 2013;39(3):272–82.
- 30. Goodeve AC. Hemophilia B: molecular pathogenesis and mutation analysis. J Thromb Haemost. 2015;13(7):1184–95.
- 31. Gailani D, Geng Y, Verhamme I, Sun MF, Bajaj SP, Messer A, Emsley J. The mechanism underlying activation of factor IX by factor XIa. Thromb Res. 2014;133 Suppl 1:S48–51.
- 32. Girolami A, Scandellari R, Scapin M, Vettore S. Congenital bleeding disorders of the vitamin K-dependent clotting factors. Vitam Horm. 2008;78:281–374.
- 33. Federici AB. Diagnosis of inherited von Willebrand disease: a clinical perspective. Semin Thromb Hemost. 2006;32(6):555–65.
- 34. Michiels JJ, Gadisseur A, Budde U, Berneman Z, van der Planken M, Schroyens W, van de Velde A, van Vliet H. Characterization, classification, and treatment of von Willebrand diseases: a critical appraisal of the literature and personal experiences. Semin Thromb Hemost. 2005;31(5):577–601.
- 35. von Lillicrap D. Willebrand disease: advances in pathogenetic understanding, diagnosis, and therapy. Hematology Am Soc Hematol Educ Program. 2013;2013:254–60.
- 36. Turner NA, Nolasco L, Ruggeri ZM, Moake JL. Endothelial cell ADAMTS-13 and VWF: production, release, and VWF string cleavage. Blood. 2009;114(24):5102–11.
- 37. López JA, Dong JF. Cleavage of von Willebrand factor by ADAMTS-13 on endothelial cells. Semin Hematol. 2004; 41(1):15–23.
- 38. Pearson JD. The control of production and release of haemostatic factors in the endothelial cell. Baillieres Clin Haematol. 1993;6(3):629–51.
- 39. Budde U, Schneppenheim R. Interactions of von Willebrand factor and ADAMTS13 in von Willebrand disease and thrombotic thrombocytopenic purpura. Hamostaseologie. 2014;34(3):215–25.
- 40. Morrison KA, Jorde UP, Garan AR, Takayama H, Naka Y, Uriel N. Acquired von Willebrand disease during CentriMag support is associated with high prevalence of bleeding during support and after transition to heart replacement therapy. ASAIO J. 2014;60(2):241–2.
- 41. Huck V, Schneider MF, Gorzelanny C, Schneider SW. The various states of von Willebrand factor and their function in physiology and pathophysiology. Thromb Haemost. 2014;111(4):598–609.
- 42. Terraube V, O'Donnell JS, Jenkins PV. Factor VIII and von Willebrand factor interaction: biological, clinical and therapeutic importance. Haemophilia. 2010;16(1):3–13.
- 43. Federici AB. The factor VIII/von Willebrand factor complex: basic and clinical issues. Haematologica. 2003;88(6):EREP02.
- 44. The Diagnosis. Evaluation, and Management of von Willebrand Disease. 2007. NHLBI von Willebrand Disease Expert Panel.
- 45. Millar CM, Riddell AF, Brown SA, Starke R, Mackie I, Bowen DJ, Jenkins PV, van Mourik JA. Survival of von Willebrand factor released following DDAVP in a type 1 von Willebrand disease cohort: influence of glycosylation, proteolysis and gene mutations. Thromb Haemost. 2008;99(5):916–24.
- 46. Haberichter SL, Castaman G, Budde U, Peake I, Goodeve A, Rodeghiero F, Federici AB, Batlle J, Meyer D, Mazurier C, Goudemand J, Eikenboom J, Schneppenheim R, Ingerslev J, Vorlova Z, Habart D, Holmberg L, Lethagen S, Pasi J, Hill FG, Montgomery RR. Identification of type 1 von Willebrand disease patients with reduced von Willebrand factor survival by assay of the VWF propeptide in the European study: molecular and clinical

<span id="page-7-0"></span>markers for the diagnosis and management of type 1 VWD (MCMDM-1VWD). Blood. 2008;111(10):4979–85.

- 47. Haberichter SL, Balistreri M, Christopherson P, Morateck P, Gavazova S, Bellissimo DB, Manco-Johnson MJ, Gill JC, Montgomery RR. Assay of the von Willebrand factor (VWF) propeptide to identify patients with type 1 von Willebrand disease with decreased VWF survival. Blood. 2006;108(10):3344–51.
- 48. Blann AD. von Willebrand factor antigen as an acute phase reactant and marker of endothelial cell injury in connective tissue diseases: a comparison with CRP, rheumatoid factor, and erythrocyte sedimentation rate. Z Rheumatol. 1991;50(5):320–2.
- 49. Knol HM, Mulder AB, Bogchelman DH, Kluin-Nelemans HC, van der Zee AG, Meijer K. The prevalence of underlying bleeding disorders in patients with heavy menstrual bleeding with and without gynecologic abnormalities. Am J Obstet Gynecol. 2013;209(3):202. e1–7. doi:[10.1016/j.ajog.2013.05.059](http://dx.doi.org/10.1016/j.ajog.2013.05.059).
- 50. Klarmann D, Eggert C, Geisen C, Becker S, Seifried E, Klingebiel T, Kreuz W. Association of ABO(H) and I blood group system development with von Willebrand factor and Factor VIII plasma levels in children and adolescents. Transfusion. 2010;50(7): 1571–80.
- 51. O'Donnell J, Boulton FE, Manning RA, Laffan MA. Amount of H antigen expressed on circulating von Willebrand factor is modified by ABO blood group genotype and is a major determinant of plasma von Willebrand factor antigen levels. Arterioscler Thromb Vasc Biol. 2002;22(2):335–41.
- 52. Blenner MA, Dong X, Springer TA. Structural basis of regulation of von Willebrand factor binding to glycoprotein Ib. J Biol Chem. 2014;289(9):5565–79.
- 53. Miller JL, Kupinski JM, Castella A, Ruggeri ZM. von Willebrand factor binds to platelets and induces aggregation in platelet-type but not type IIB von Willebrand disease. J Clin Invest. 1983;72(5): 1532–42.
- 54. Gadisseur A, van der Planken M, Schroyens W, Berneman Z, Michiels JJ. Dominant von Willebrand disease type 2M and 2U are

variable expressions of one distinct disease entity caused by lossof- function mutations in the A1 domain of the von Willebrand factor gene. Acta Haematol. 2009;121(2-3):145–53.

- 55. Miller CH, Kelley L, Green D. Diagnosis of von Willebrand disease type 2N: a simplified method for measurement of factor VIII binding to von Willebrand factor. Am J Hematol. 1998;58(4):311–8.
- 56. Federici AB, Budde U, Castaman G, Rand JH, Tiede A. Current diagnostic and therapeutic approaches to patients with acquired von Willebrand syndrome: a 2013 update. Semin Thromb Hemost. 2013;39(2):191–201.
- 57. Warkentin TE, Moore JC, Morgan DG. Aortic stenosis and bleeding gastrointestinal angiodysplasia: is acquired von Willebrand's disease the link? Lancet. 1992;340(8810):35–7.
- 58. Koyama T, Fujimoto K, Shima M. Acquired von Willebrand syndrome associated with Hashimoto's thyroiditis and subcutaneous mucosa-associated lymphoid tissue lymphoma. Intern Med. 2013;52(23):2661–3.
- 59. Facon T, Caron C, Courtin P, Wurtz A, Deghaye M, Bauters F, Mazurier C, Goudemand J. Acquired type II von Willebrand's disease associated with adrenal cortical carcinoma. Br J Haematol. 1992;80(4):488–94.
- 60. Federici AB. Acquired von Willebrand syndrome associated with hypothyroidism: a mild bleeding disorder to be further investigated. Semin Thromb Hemost. 2011;37(1):35–40.
- 61. World Federation of Hemophilia. Protocols for the treatment of Hemophilia and von Willebrand disease. In: Hemophilia of Georgia, 3rd edn, Vol. 14; 2008, Montreal: World Federation of Hemophilia.
- 62. Mensah PK, Gooding R. Surgery in patients with inherited bleeding disorders. Anaesthesia. 2015;70 Suppl 1:112–20. e39-40.
- 63. Kouides PA. An update on the management of bleeding disorders during pregnancy. Curr Opin Hematol. 2015;22(5):397–405.
- 64. Kadir RA, Economides DL, Braithwaite J, Goldman E, Lee CA. The obstetric experience of carriers of haemophilia. Br J Obstet Gynaecol. 1997;104(7):803–10.