blood advances

Baronciani L, Peake I, Schneppenheim R, et al. Genotypes of European and Iranian patients with type 3 von Willebrand disease enrolled in 3WINTERS-IPS. *Blood Adv*. 2021;5(15):2987-3001.

Page 2987: Author affiliation no. 28 (Flora Peyvandi) was incorrect. The affiliation should be the Department of Pathophysiology and Transplantation, Università degli Studi di Milano, Milan, Italy.

Page 2991: Figure 2 has the following corrections:

- The word "different" was misspelled in the top heading.
- The variant c.7729-7C>T should appear as c.7729 + 7C>T.
- The following variants were not clearly readable: p.S1699F, p.S2469P, and p.C2739Y.

The corrected Figure 2 is shown below.

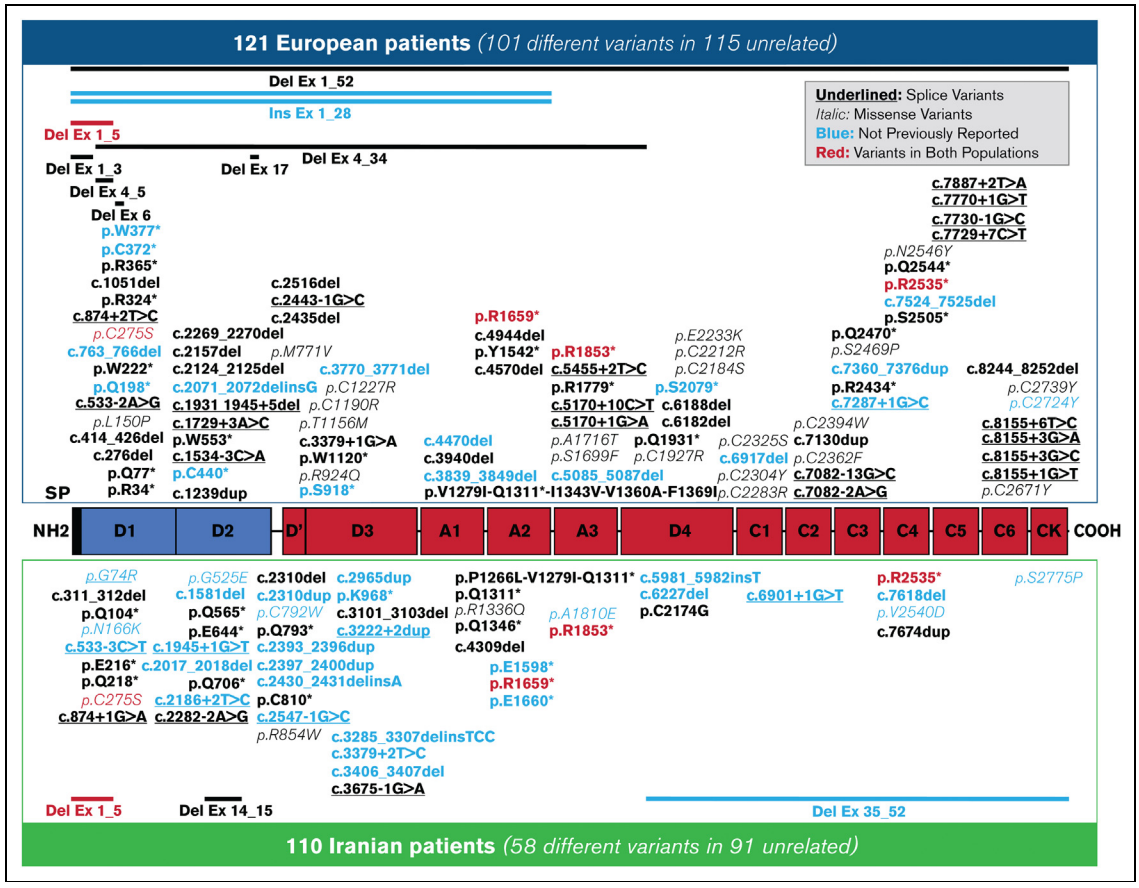

Figure 2. Distribution of the 154 different unique variants identified in the 3WINTERS-IPS cohort. A schmatic representation of the pro-VWF polypeptide with the homologous repeated domain structure. Variants identified in the European population are reported at the top; those found in the Iranian population are reported at the bottom. The candidate missense mutations are reported in italics, and the potential splice site mutations are undefined. The 5 variants identified in both populations are reported in red. The variants not previously reported in the EAHAD, HGMD, or Ensembl databases are reported in blue.

Page 2996: In Table 3, the reference number in the "First reported" column for variant NM_000552.3:c.3379 + 1G>A should be 80, not 81.

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