



Bo's Case

NOTES

Phenotype	<p>A laceration on left index finger with prolonged bleeding previous episodes of prolonged bleeding which hadn't "risen to the level of an ER visit but were concerning."</p> <p>Family history: • No "genetic" family history is available as Bo was adopted from China at the age of 3 years old</p>
Preliminary Diagnosis	<p>Hemophilia (sub-type not determined yet)</p>
Genetic Variation(s)	<p>F9 p.Asp110Gly</p>
Laboratory Assertion(s)	<p>pathogenic</p>
Variant Information: <ul style="list-style-type: none"> • Asserted interpretation listed in ClinVar • HGVS names from ClinVar • Is population data available in dbSNP? 	<p>pathogenic</p> <p>NG_007994.1(F9): g.15392A>G NP_000124.1(F9): p.Asp110Gly</p> <p>rs137852234 Yes! And it is pretty darn rate.</p>
Gene Information in NCBI Gene: <ul style="list-style-type: none"> • Symbol and Name • Gene Summary • Tissue Expression information • Gene Ontology information 	<p>F9 & Coagulation factor IX</p> <p>F9 ...vitamin K-dependent coagulation factor IX that circulates in the blood as an inactive zymogen....converted to an active form by factor XIa,...activates factor X ... through interactions with Ca+2 ions, membrane phospholipids, and factor VIII. Alterations of this gene...cause factor IX deficiency, which is a recessive X-linked disorder.... [provided by RefSeq, Sep 2015]</p> <p>Pretty much just expressed in the liver</p> <p>Extracellular Blood coagulation Ca+2-binding & endopeptidase</p>
Ultimate Impacted Biomolecule based on: <ul style="list-style-type: none"> • GDV to view the chromosome and gene region • RefSeqGene Graphics view of gene region and transcript(s) • RefSeq Protein Graphics view of protein and domains • CDD or iCn3D to view a structure, <i>as needed</i> 	<p>Located in the coding region within exon 4.</p> <p>Located in the coding region within exon 4.</p> <p>The protein is made, but with a change in amino acid 110 from an acidic Aspartate to a neutral Glycine.</p> <p>The variant is identified as one of 3 residues annotated as critical for binding to Ca+2. The change from acidic Aspartate residue to neutral Glycine likely prevents its participation.</p>
Proposed Molecular Mechanism of Variant Impact	<p>This changes an acidic residue which is needed for binding a critical calcium ion which is required for F9 function.</p>
How does this relate back to the phenotype (symptoms/clinical features & diagnosis)?	<p>With the loss of one of 3 coordinating residues for a critical calcium ion, the F9 protein is not fully functional and may not effectively activate the next clotting factor. This correlates with a less severe phenotype than other children might experience.</p>