Comprehensive Platelet Disorder Panel

Versiti offers comprehensive genetic analysis to detect sequence variants and large deletions and duplications in 62 genes plus one targeted variant known to cause platelet function disorders and/or inherited thrombocytopenia. This panel can be ordered as:

- Next generation sequencing (NGS) only*;
- NGS with reflex to array Comparative Genomic Hybridization (aCGH) deletion/ duplication analysis if clinically significant variants explaining the patient's phenotype are not detected by sequencing;
- NGS with concurrent aCGH deletion/duplication analysis (both testing methodologies performed simultaneously); or
- Deletion/duplication by aCGH only.
- * Includes PLAU performed by aCGH

Disorders of platelet number and/or function are a heterogeneous group of disorders with overlapping clinical phenotypes, generally differentiated by platelet counts. Inherited thrombocytopenia disorders are typically characterized by platelet counts less than 150,000/uL. but often can vary with age, sex and ethnic background, while platelet function disorders are usually, but not always, associated with normal platelet counts and are caused by defects in platelet adhesion, glycoprotein expression, receptor function, signaling pathways, aggregation, cytoskeleton proteins, secretion, granular contents, or abnormalities in procoagulant activity. Symptoms of platelet disorders may include purpura, petechiae, prolonged bleeding from cuts, epistaxis, gum bleeding, excessive bleeding after surgery, hemoptysis, hematuria, and menorrhagia in women. Severe platelet disorders

can present in the newborn period or early childhood, while mild thrombocytopenia may remain undiagnosed until incidentally detected on routine blood testing in adulthood. Some inherited platelet disorders have only hematologic manifestations, such as differences in platelet size or distinctive granulocyte inclusions, while other syndromic types present with additional non-hematologic manifestations. Certain types of platelet disorders cause predisposition to acute myelogenous or lymphoid neoplasms.

Misdiagnosis of platelet disorders can result in inappropriate therapies and inadequate surveillance for additional medical complications, underscoring the importance of accurate diagnosis. The diagnosis may be difficult to establish based solely on functional studies, as platelet function assays are not widely available due to their technical complexity and need for immediate testing on fresh patient platelets due to limited sample stability. Advances in genetic testing through next-generation sequencing allows for identification of underlying genetic defects and for distinguishing inherited platelet disorder cases from other acquired disorders of platelet number, such as immune thrombocytopenia, and platelet function, such as those induced by medications or systemic disorders. Accurate diagnosis provides information about the phenotype and prognosis, guides medical management decisions, assists with the identification of affected family members, and allows for accurate genetic recurrence risk assessment. Variants in genes known to be associated with syndromic or non-syndromic platelet disorders may be inherited in an autosomal recessive, autosomal dominant or X-linked manner. More common and rare types of platelet disorders will be identified with this panel. While thrombocytopenia may be the presenting symptom in type 2B von Willebrand disease (VWD) and congenital thrombotic thrombocytopenic purpura (TTP), these disorders are usually suspected based on specific plasma studies and genetic testing of VWF and ADAMTS13 for these conditions is offered by our laboratory separately (individually or as part of other genetic panels).



The NGS panel evaluates for single nucleotide variants and small deletions and duplications, which are most commonly responsible for genetic disease. However, large deletions and duplications, also referred to as copy number variations (CNVs), are a known cause of genetic disorders, but can escape detection by next generation sequence analysis. Additional testing with aCGH deletion/duplication analysis is available for all genes on this panel to evaluate for large deletions and duplications encompassing one or more exons, or affecting an entire gene. Note that the Quebec Platelet Disorder (QPD) is associated with a heterozygous 77.9-kb tandem duplication of the *PLAU* gene which will be detected by aCGH and not by NGS; analysis of *PLAU* by aCGH is included in the otherwise NGS-only version of this panel.

Refer to the table for further information about each gene in the Comprehensive Platelet Disorder Panel, including the clinical phenotype and inheritance pattern.

Gene	Clinical Phenotype	Disorder of platelet number and/or platelet function	Inheritance
ABCG5	Sitosterolemia with macrothrombocytopenia: xanthomas, premature atherosclerosis, arthritis, hepatic dysfunction and hematologic abnormalities including stomatocytosis leading to hemolytic anemia and macrothrombocytopenia with mild to moderate bleeding and large platelets surrounded by vacuoles on peripheral smear, as well as splenomegaly	Thrombocytopenia	Autosomal Recessive
ABCG8	Sitosterolemia with macrothrombocytopenia: xanthomas, premature atherosclerosis, arthritis, hepatic dysfunction and hematologic abnormalities including stomatocytosis leading to hemolytic anemia and macrothrombocytopenia with mild to moderate bleeding and large platelets surrounded by vacuoles on peripheral smear, as well as splenomegaly	Thrombocytopenia	Autosomal Recessive
ACTB	ACTB-related thrombocytopenia: mild developmental disability, non-specific minor facial abnormalities, microcephaly and thrombocytopenia with platelet anisocytosis	Thrombocytopenia	Autosomal Dominant
ACTN1	<i>ACTN1</i> -related thrombocytopenia (platelet-type bleeding disorder 15): mild macrothrombocytopenia with minimal or absent bleeding tendency	Thrombocytopenia	Autosomal Dominant
ANKRD26	ANKRD26-related thrombocytopenia (thrombocytopenia 2): thrombocytopenia with normal platelet size, mild/absent bleeding and an increased predisposition to hematologic myeloid malignancies	Thrombocytopenia	Autosomal Dominant
ANO6	Scott syndrome: platelet dysfunction with mild to moderate bleeding phenotype with normal platelet aggregation and platelet counts, and decreased platelet procoagulant activity with characteristic flow cytometry findings	Platelet dysfunction	Autosomal Recessive
AP3B1	Hermansky-Pudlak syndrome type 2 (HPS2): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder, as well as pulmonary fibrosis and neutropenia	Platelet dysfunction	Autosomal Recessive
AP3D1	Hermansky-Pudlak syndrome type 10 (HPS10): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder, as well as neutropenia, seizures and developmental delay	Platelet dysfunction	Autosomal Recessive
ARPC1B	<i>ARPC1B</i> -related thrombocytopenia: microthrombocytopenia, decreased platelet dense granules, allergic and inflammatory disease	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
BLOC1S3	Hermansky-Pudlak syndrome type 8 (HPS8): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
BLOC1S6	Hermansky-Pudlak syndrome type 9 (HPS9): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
CDC42	<i>CDC42</i> -related thrombocytopenia (Takenouchi-Kosaki syndrome): macrothrombocytopenia, variable intellectual disability, distinct facial features, lymphedema, camptodactyly, and variable involvement of other organ systems	Thrombocytopenia	Autosomal Dominant
CYCS	CYCS-related thrombocytopenia (thrombocytopenia 4): non-syndromic thrombocytopenia with normal platelet size	Thrombocytopenia	Autosomal Dominant
DIAPH1	<i>DIAPH1-related thrombocytopenia:</i> macrothrombocytopenia and sensorineural hearing loss	Thrombocytopenia	Autosomal Dominant
DTNBP1	Hermansky-Pudlak syndrome type 7 (HPS7): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
ETV6	<i>ETV6</i> -related thrombocytopenia (thrombocytopenia 5): thrombocytopenia with normal platelet size, red cell macrocytosis, mild to moderate bleeding and predisposition to both myeloid and lymphoid malignancies	Thrombocytopenia	Autosomal Dominant
FERMT3	Leukocyte adhesion deficiency-III (LAD-III): severe bleeding phenotype with a Glanzmann thrombasthenia-like phenotype on platelet aggregation studies and associated immunodeficiency	Platelet dysfunction	Autosomal Recessive

FLI1	<i>FLI1</i> -related thrombocytopenia (platelet-type bleeding disorder-21): macrothrombocytopenia with moderate bleeding from platelet dysfunction due to alpha granule deficiency (large/fused platelet alpha granules on platelet electron microscopy), with or without delta granule deficiency	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
FLNA	FLNA-related thrombocytopenia: macrothrombocytopenia with or without associated periventricular heterotopia	Thrombocytopenia with platelet dysfunction	X-linked
FYB1	<i>FYB1</i> -related thrombocytopenia (thrombocytopenia 3): non-syndromic microthrombocytopenia and platelet dysfunction leading to increased bleeding	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
GATA1	GATA1-related X-linked cytopenia: characterized by macrothrombocytopenia and/ or anemia with moderate bleeding due to platelet alpha granule deficiency	Thrombocytopenia with platelet dysfunction	X-linked
GFI1B	<i>GFI1B-related thrombocytopenia</i> (platelet-type bleeding disorder-17): macrothrombocytopenia with platelet alpha granule deficiency leading to variable bleeding tendency, red cell anisopoikilocytosis, increased numbers of dysplastic megakaryocytes and increased platelet CD34 expression	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
GNE	<i>GNE-related thrombocytopenia:</i> macrothrombocytopenia with mild to moderate bleeding with or without myopathy	Thrombocytopenia	Autosomal Recessive
	Bernard Soulier syndrome (BSS): macrothrombocytopenia with normal platelet granularity and moderate to severe bleeding due to decreased/absent/ dysfunctional platelet GPIb/IX expression with decreased/absent platelet aggregation with ristocetin	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
GP1BA	<i>GP1BA-related macrothrombocytopenia:</i> mild to moderate thrombocytopenia with absent/mild bleeding	Thrombocytopenia	Autosomal Dominant
	Platelet type von Willebrand disease: thrombocytopenia with mild bleeding due to loss of VWF high molecular weight multimers from increased binding of platelets and VWF	Thrombocytopenia	Autosomal Dominant
GP1BB	Bernard Soulier syndrome (BSS): macrothrombocytopenia with normal platelet granularity and moderate to severe bleeding due to decreased/absent/ dysfunctional platelet GPIb/IX expression with decreased/absent platelet aggregation with ristocetin	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
	<i>GP1BB-related macrothrombocytopenia:</i> mild to moderate thrombocytopenia with absent/mild bleeding	Thrombocytopenia	Autosomal Dominant
GP6	<i>GP6-related platelet dysfunction</i> (platelet- type bleeding disorder 11): mild bleeding and decreased aggregation response to collagen on platelet aggregation studies due to deficiency of platelet glycoprotein VI	Platelet dysfunction	Autosomal Recessive
GP9	Bernard Soulier syndrome (BSS): macrothrombocytopenia with normal platelet granularity and moderate to severe bleeding due to decreased/absent/ dysfunctional platelet GPIb/IX expression with decreased/absent platelet aggregation with ristocetin	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
	GP9-related macrothrombocytopenia: mild to moderate thrombocytopenia with absent/mild bleeding	Thrombocytopenia	Autosomal Dominant
HOXA11	Radioulnar synostosis with amegakaryocytic thrombocytopenia (RUSAT1): thrombocytopenia with normal platelet size and radial abnormalities	Thrombocytopenia	Autosomal Dominant
HPS1	Hermansky-Pudlak syndrome type 1 (HPS1): oculocutaneous albinism of variable severity and mild bleeding due to a platelet storage pool disorder, as well as pulmonary fibrosis and granulomatous colitis	Platelet dysfunction	Autosomal Recessive
HPS3	Hermansky-Pudlak syndrome type 3 (HPS3): mild ocular albinism and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
HPS4	Hermansky-Pudlak syndrome type 4 (HPS4): oculocutaneous albinism and mild bleeding due to a platelet storage pool disorder, as well as pulmonary fibrosis and granulomatous colitis	Platelet dysfunction	Autosomal Recessive
HPS5	Hermansky-Pudlak syndrome type 5 (HPS5): mild ocular albinism and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
HPS6	Hermansky-Pudlak syndrome type 6 (HPS6): mild ocular albinism and mild bleeding due to a platelet storage pool disorder	Platelet dysfunction	Autosomal Recessive
ITGA2B	Glanzmann thrombasthenia: normal platelet count with severe bleeding and decreased/absent platelet aggregation with all agonists except ristocetin due to decreased/absent/dysfunctional expression of platelet glycoprotein (GP) IIb/IIIa	Platelet dysfunction	Autosomal Recessive
	<i>ITGA2B</i> -related macrothrombocytopenia: mild to moderate thrombocytopenia with absent/mild bleeding	Thrombocytopenia	Autosomal Dominant
ITGB3	Glanzmann thrombasthenia: normal platelet count with severe bleeding and decreased/absent platelet aggregation with all agonists except ristocetin due to decreased/absent/dysfunctional expression of platelet glycoprotein (GP) IIb/IIIa	Platelet dysfunction	Autosomal Recessive
	<i>ITGB3</i> -related macrothrombocytopenia: mild to moderate thrombocytopenia with absent/mild bleeding	Thrombocytopenia	Autosomal Dominant

KDSR			
ND ON	<i>KDSR-related thrombocytopenia</i> (Erythrokeratodermia variabilis et progressiva 4): thrombocytopenia with normal platelet size and platelet dysfunction with or without skin hyperkeratosis and ichthyosis	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
LYST	Chediak-Higashi syndrome: partial oculocutaneous albinism, immunodeficiency, and a mild bleeding from platelet delta granule deficiency	Platelet dysfunction	Autosomal Recessive
MECOM	<i>MECOM-associated syndrome</i> (Radioulnar synostosis with amegakaryocytic thrombocytopenia 2): bone marrow failure with hypomegakaryocytic thrombocytopenia with normal platelet size, radioulnar synostosis, clinodactyly, cardiac and renal malformations, B-cell deficiency and hearing loss	Thrombocytopenia	Autosomal Dominant
МҮН9	MYH9-related disorders (MYH9-RD) characterized by macrothrombocytopenia with or without extra hematologic manifestations including renal dysfunction, hearing loss, cataracts and liver enzyme elevation	Thrombocytopenia	Autosomal Dominant
MPIG6B	<i>MPIG6B</i> -related thrombocytopenia: macrothrombocytopenia with focal myelofibrosis	Thrombocytopenia	Autosomal Recessive
MPL	Congenital amegakaryocytic thrombocytopenia (CAMT): thrombocytopenia with normal platelet size and progression to bone marrow failure	Thrombocytopenia	Autosomal Recessive
NBEA	NBEA-related platelet dysfunction: neurodevelopmental disorders, including autism and seizures and moderate bleeding due to platelet delta storage pool disorder	Platelet dysfunction	Autosomal Dominant
NBEAL2	Gray platelet syndrome (GPS): macrothrombocytopenia with mild to moderate bleeding due to alpha granule deficiency, splenomegaly and bone marrow fibrosis	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
P2RY12	<i>P2RY12-related platelet dysfunction</i> (platelet-type bleeding disorder 8): mild- moderate mucocutaneous bleeding and excessive bleeding in response to trauma or surgery due to impaired platelet aggregation responses to ADP	Platelet dysfunction	Autosomal Recessive
PLA2G4A	<i>PLA2G4A</i> -related platelet dysfunction (cytosolic phospholipase-A2 alpha deficiency): platelet dysfunction from a metabolic defect and small bowel ulcers caused by decreased production of eicosanoids	Platelet dysfunction	Autosomal Recessive
PLAU*	Quebec Platelet Disorder (QPD): delayed onset bleeding, large trauma induced hematomas, hemarthrosis, muscle bleeds and hematuria from hyperfibrinolysis due to increased platelet urokinase plasminogen activator from a tandem 77.9kb duplication encompassing the PLAU gene	Platelet dysfunction	Autosomal Dominant
PRKACG †	<i>PRKACG</i> -related thrombocytopenia (platelet-type bleeding disorder 19): severe macrothrombocytopenia with associated platelet dysfunction leading to moderate to severe bleeding attributed to the <i>PRKACG</i> p.lle74Met variant.	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
RASGRP2	RASGRP2-related platelet dysfunction (platelet-type bleeding disorder 18): moderate to severe bleeding and decreased platelet aggregation with ADP and epinephrine and in some cases arachidonic acid, collagen and thrombin	Platelet dysfunction	Autosomal Recessive
RBM8A	Thrombocytopenia absence radius (TAR) syndrome: bilateral absence of the radii and severe thrombocytopenia with normal platelet size that is usually transient	Thrombocytopenia	Autosomal Recessive
RNU4ATAC	Roifman syndrome: growth retardation, distinctive facial features, microcephaly, cognitive delay, retinal dystrophy, spondyloepiphyseal dysplasia, hypogammaglobulinemia and in some cases thrombocytopenia	Thrombocytopenia	Autosomal Recessive
RUNX1	Familial platelet disorder with predisposition to myeloid leukemia (FPD/AML): mild to moderate thrombocytopenia with normal platelet size, platelet delta storage pool disorder and a predisposition to development of myeloid malignancies	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
SLFN14	<i>SLFN14-related thrombocytopenia</i> (platelet-type bleeding disorder 20): mild to moderate macrothrombocytopenia with associated platelet dysfunction from dense granule deficiency leading to variable bleeding	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
SRC	<i>SRC</i> -related thrombocytopenia (thrombocytopenia 6): thrombocytopenia and bleeding with associated myelofibrosis and bone pathology	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
STIM1	STIM1-related thrombocytopenia (Tubular aggregate myopathy and Stormorken syndrome): variable muscle weakness, miosis, thrombocytopenia with normal platelet size, hyposplenism, ichthyosis, dyslexia and short stature. Electron dense platelet inclusions and target-like organelles are characteristic	Thrombocytopenia with platelet dysfunction	Autosomal Dominant
STXBP2	Familial hemophagocytic lymphohistiocytosis type 5 (FLH5): prolonged fever, cytopenias and hepatosplenomegaly due to proliferation and infiltration of hyperactivated macrophages and T-lymphocytes	Thrombocytopenia	Autosomal Recessive
TBXA2R	Thromboxane receptor defect: pathogenic variants in <i>TBXA2R</i> have been proposed as contributing to a bleeding phenotype in the presence of additional pathogenic variants in genes affecting platelet function; these variants cause impaired platelet response to arachidonic acid and U46619 in vitro, but have not been shown to consistently correlate with a clinical phenotype	Platelet dysfunction	Risk allele
TBXAS1	<i>TBXAS1</i> -related platelet dysfunction (Ghosal syndrome; platelet-type bleeding disorder 14): increased bone density and platelet dysfunction due to impaired	Platelet dysfunction	Autosomal Recessive

THPO	<i>THPO</i> -related thrombocytopenia: characterized by severe thrombocytopenia with normal platelet size progressing to bone marrow failure	Thrombocytopenia	Autosomal Recessive
TUBB1	<i>TUBB1</i> -related thrombocytopenia: mild macrothrombocytopenia and minimal/ absent bleeding	Thrombocytopenia	Autosomal Dominant
VIPAS39	Arthrogryposis, renal dysfunction, and cholestasis syndrome type 2 (ARCS2): macrothrombocytopenia with platelet dysfunction from alpha granule deficiency with associated arthrogryposis, renal dysfunction, and cholestasis	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
VPS33B	Arthrogryposis, renal dysfunction, and cholestasis syndrome type 1 (ARCS1): macrothrombocytopenia with platelet dysfunction from alpha granule deficiency with associated arthrogryposis, renal dysfunction, and cholestasis	Thrombocytopenia with platelet dysfunction	Autosomal Recessive
WAS	<i>WAS-related disorders:</i> spectrum of disorders including Wiskott-Aldrich syndrome characterized by microthrombocytopenia, eczema and recurrent infections, X-linked thrombocytopenia and X-linked neutropenia	Thrombocytopenia	X-linked
WIPF1	Wiskott-Aldrich syndrome type 2 (WAS2): recurrent infections, eczema, thrombocytopenia with normal platelet size, defective T cell proliferation and impaired natural killer cell function	Thrombocytopenia	Autosomal Recessive

* Available only via aCGH.

+ PRKACG NGS includes only a region to cover the p.lle74Met variant.

Indications for testing:

Comprehensive Platelet Disorder Panel (NGS and/or aCGH), order code 4830:

The Comprehensive Platelet Disorder Panel should be considered:

- In patients with thrombocytopenia with bleeding out of proportion to the degree of decrease in platelet counts in which congenital platelet dysfunction is suspected
- In patients with lifelong thrombocytopenia and bleeding in which platelet function testing cannot be obtained or is unreliable due to the severity of the thrombocytopenia
- In patients with suspected platelet dysfunction and intermittent thrombocytopenia fluctuating with low normal platelet counts

Single Gene Analysis (order code 4855) or Custom Blood Disorder Panel (Order Code 4850), (NGS and/or aCGH):

 Analysis of genes included in this panel may also be ordered as a standalone Single Gene Analysis or as a Custom Blood Disorder Panel (2-10 genes), by NGS and/or by aCGH, as dictated by the patient's clinical and laboratory phenotype, as well as their ancestry, or to supplement previous genetic testing.

Targeted Familial Variant Analysis (order code 4970):

Targeted variant analysis for clinical diagnosis, carrier identification, or prenatal diagnosis can also be performed on any gene in the panel when the pathogenic variant(s) is known in the family. If the proband was not tested at Versiti, a control sample is preferred and may be required (please call the laboratory to discuss). If the familial variant is a large deletion or duplication, aCGH for the involved gene is required.

For clinical questions about laboratory tests and test utilization support, contact Versiti Client Services: (414) 937-6396 or 800-245-3117, Option 1, to be directed to our genetic counselors and clinical support team.

Informed Consent

It is recommended that healthcare providers obtain informed consent from the patient when genetic testing is ordered, consistent with any applicable state laws and regulations, documenting that the patient has been advised of understands the indications for and implications of the genetic test. This panel is designed for clinical detection of germline genetic variants in genes with strong or definitive evidence for causality of platelet function disorders and/or inherited thrombocytopenia. Test results may nonetheless yield genetic findings that may be unrelated to the current clinical presentation, and/ or may carry individual or familial implications such as risk for syndromic manifestation, predisposition to malignancy, and/or reproductive implications (such as carrier status). If needed, an informed consent form for Versiti Hematology Genetics testing can be found at http://www.versiti.org/hg under forms.

Test method:

NGS: This next-generation sequencing assay analyzes the complete coding region of 62 genes (excluding PLAU and *PRKACG*) plus a minimum 30bp of non-coding DNA, including intron-exon boundaries, and is compared to the build GRCh37.p13 reference sequence. ANKRD26 analysis also includes approximately 200bp upstream of coding region to identify clinically significant variants in the 5'UTR. Targeted sequencing of *PRKACG* c.222C>G (p.IIe74Met) is also included. These targeted regions are captured by hybridization, amplified, and sequenced by massively parallel sequencing. Regions will have a minimum coverage of 50x and those regions with less than 50 sequencing reads or low quality coverage are supplemented with Sanger sequencing. All regions are covered by bidirectional analysis. Variants are identified by a customized bioinformatics pipeline, analyzed and comprehensively interpreted by our team of practicing hematologists with

expertise in non-malignant hematology and laboratory diagnostics, scientists, and genetic counselors. All reported variants, including pathogenic, likely pathogenic, and variants of uncertain significance, are confirmed by Sanger sequencing. For prenatal testing, analysis of variable number tandem repeats (VNTR) is used to confirm results are not affected by maternal cell contamination.

aCGH: The specific genes are analyzed for copy number variations due to deletion or duplication by high density gene-focused array Comparative Genomic Hybridization. Probes are approximately 60bp in length and density of coverage in exonic regions is a minimum of 4 probes per 500 bp. Genomic DNA for the samples and sex-matched references are denatured, labeled with fluorescent dye and hybridized, the array is washed and scanned, and analysis is performed for the specific genes requested.

Assay sensitivity and limitations:

NGS: The analytical sensitivity of the NGS test is >99% for single nucleotide changes and insertions and deletions of less than 20 bp. NGS analysis is not designed to detect large deletions or duplications (>20 bp), or variants that are outside the regions sequenced. Low level mosaicism will not be detected by this sequencing methodology.

aCGH: Balanced chromosomal rearrangements (i.e., translocations, inversions) or point mutations that may be the cause of the clinical phenotype cannot be detected via aCGH. Any exonic deletion or duplication smaller than 500bp may not be detected. Low level of mosaicism will not be detected by aCGH. Probe performance could be affected by multiple SNPs in a given region. Breakpoints occurring outside the targeted gene(s) will not be defined.

Clinical Sensitivity

The clinical sensitivity of comprehensive genetic testing (NGS and aCGH) of the 63 genes in this panel is highest in patients with lifelong thrombocytopenia and/or platelet dysfunction in whom acquired causes of a platelet disorder have been ruled out. A family history of a platelet disorder with a similar phenotype increases the pre-test probability of a congenital platelet disorder.

Reporting of Results

Results are classified and reported in accordance with ACMG next-generation sequencing and copy number variation standards and guidelines. Sequence variants and large deletions and duplications predicted to be pathogenic, likely pathogenic, and of uncertain significance will be reported; variants classified as likely benign or benign are typically not reported but such data are available upon request. Sequence variants are described using standard Human Genome Variation Society (HGVS) nomenclature (http://hgvs.org); copy number variants are described in accordance with the International System for Human Cytogenomic Nomenclature (ISCN).

Specimen Requirements

Parental/Patient/Pediatric: 3-5 mL Whole blood (EDTA tube, lavender top), 2-5 mL Bone marrow (EDTA tube, lavender top), 3-4 Buccal swabs, or ≥1ug of DNA at ≥50ng/ uL of High Quality DNA.

Fetal: 7-15 mL amniotic fluid, 5-10 mg chorionic villi; back up culture of amniocytes or chorionic villi is highly recommended. Cultured: Two T25 flasks cultured amniocytes or chorionic villi (2x10⁶ minimum). Maternal blood sample of 3-5 mL Whole blood (EDTA tube, lavender top) is requested for all prenatal samples for maternal cell contamination studies. For questions please contact the laboratory to discuss sample requirements.



Shipping Requirements

Ship on an ice pack at room temperature. Protect from freezing. Place the specimen and the requisition into plastic bags and seal. Insert into a Styrofoam container, seal and place into a sturdy cardboard box, and tape securely. Ship the package in compliance with your overnight carrier guidelines. Label with the following address:

Client Services/Diagnostic Laboratory Versiti 638 N. 18th St Milwaukee, WI, 53233



Required Forms

Please complete all pages of the requisition form. Clinical history (including patient-reported ancestry, clinical diagnosis, family history, and relevant laboratory findings) is necessary for optimal interpretation of genetic test results and recommendations. Clinical and laboratory history can either be recorded on the

ORDER

requisition form or clinical and laboratory reports can be submitted with the sample.

CPT Codes/Billing/Turnaround Time

Test code: 4830

For suggested CPT codes, visit Versiti.org/test-catalog

Turnaround time: NGS only, aCGH only, or NGS and aCGH concurrently: 21 days

NGS reflex to aCGH: 21 days (if NGS only, aCGH not needed) or 42 days (with reflex to aCGH)

The CPT codes provided are subject to change as more

information becomes available. CPT codes are provided only as guidance to assist clients with billing.

For additional information related to shipping, billing or pricing, please contact Versiti Client Services: (414) 937-6396 or 800-245-3117, Option 1, or LabInfo@versiti.org

References

Inherited platelet disorder references

- 1. Balduini CL, Melazzini F. Research at the heart of hematology: thrombocytopenias and platelet function disorders. Haematologica. 2017;102(2):203-205. doi:10.3324/haematol.2016.158055
- 2. Bastida JM, Lozano ML, Benito R, et al. Introducing high-throughput sequencing into mainstream genetic diagnosis practice in inherited platelet disorders. Haematologica. 2018;103(1):148-162. doi:10.3324/haematol.2017.171132
- 3. Johnson B, Lowe GC, Futterer J, et al. Whole exome sequencing identifies genetic variants in inherited thrombocytopenia with secondary qualitative function defects. Haematologica. 2016;101(10):1170-1179. doi:10.3324/haematol.2016.146316
- Rabbolini D, Connor D, Morel-Kopp MC, et al. An integrated approach to inherited platelet disorders: results from a research collaborative, the Sydney Platelet Group. Pathology. 2020;52(2):243-255. doi:10.1016/j.pathol.2019.10.005
- Lambert MP. Inherited Platelet Disorders: A Modern Approach to Evaluation and Treatment. Hematol Oncol Clin North Am. 2019;33(3):471-487. doi:10.1016/j.hoc.2019.01.008
- Greinacher A, Pecci A, Kunishima S, et al. Diagnosis of inherited platelet disorders on a blood smear: a tool to facilitate worldwide diagnosis of platelet disorders. J Thromb Haemost. 2017;15(7):1511-1521. doi:10.1111/jth.13729
- Lentaigne C, Freson K, Laffan MA, Turro E, Ouwehand WH; BRIDGE-BPD Consortium and the ThromboGenomics Consortium. Inherited platelet disorders: toward DNA-based diagnosis. Blood. 2016;127(23):2814-2823. doi:10.1182/blood-2016-03-378588
- 8. Maclachlan A, Watson SP, Morgan NV. Inherited platelet disorders: Insight from platelet genomics using next-generation sequencing. Platelets. 2017;28(1):14-19. doi:10.1080/09537104.2016.1195492
- 9. Megy K, Downes K, Simeoni I, et al. Curated disease-causing genes for bleeding, thrombotic, and platelet disorders: Communication from the SSC of the ISTH. J Thromb Haemost. 2019;17(8):1253-1260. doi:10.1111/jth.14479
- Sharma R, Perez Botero J, Jobe SM. Congenital Disorders of Platelet Function and Number. Pediatr Clin North Am. 2018;65(3):561-578. doi:10.1016/j.pcl.2018.02.009

Variant interpretation references

- 11. Bean LJH, Funke B, Carlston CM, et al. Diagnostic gene sequencing panels: from design to report-a technical standard of the American College of Medical Genetics and Genomics (ACMG). Genet Med. 2020;22(3):453-461. doi:10.1038/s41436-019-0666-z
- 12. Rehm HL, Bale SJ, Bayrak-Toydemir P, et al. ACMG clinical laboratory standards for next-generation sequencing. Genet Med. 2013;15(9):733-747. doi:10.1038/gim.2013.92
- Richards S, Aziz N, Bale S, et al. Standards and guidelines for the interpretation of sequence variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics and the Association for Molecular Pathology. Genet Med. 2015;17(5):405-424. doi:10.1038/gim.2015.30
- 14. Riggs ER, Andersen EF, Cherry AM, et al. Technical standards for the interpretation and reporting of constitutional copy-number variants: a joint consensus recommendation of the American College of Medical Genetics and Genomics (ACMG) and the Clinical Genome Resource (ClinGen). Genet Med. 2020;22(2):245-257. doi:10.1038/s41436-019-0686-8.

