

Primary Antibody Deficiency Panel, Sequencing and Deletion/Duplication

Primary antibody deficiency (PAD) syndromes are a group of rare genetic disorders affecting antibody (immunoglobulin) production. They include common variable immunodeficiency (CVID) disorders, agammaglobulinemia, and hyper-IgM syndrome. Molecular testing is used to determine the genetic etiology of PAD in affected individuals.

Disease Overview

Symptoms

- Unusual, opportunistic, or severe infections
- Infections typically affect multiple organs/organ systems:
 - Lungs
 - Pneumonia/empyema
 - Gastrointestinal
 - Intermittent or chronic diarrhea
 - Skin (infections)
 - Head and neck
 - Oral ulcers/gingivitis/stomatitis
 - Conjunctivitis
 - Otitis media
 - Lymphadenopathy
 - Central nervous system
 - Meningitis
- Other signs
 - Sepsis
 - Failure to thrive
 - Splenomegaly
 - Autoimmune conditions
 - Neutropenia
 - Granulomatous disease
 - Lymphoid and nonlymphoid malignancies

Age of Onset

- Agammaglobulinemia and hyper-IgM syndrome usually occur within the first 2 years of life.
- CVID manifests at all ages, but most often in the second and third decade.

Incidence

Estimated at 4.6/100,000

Inheritance

X-linked, autosomal dominant, or autosomal recessive, depending on the causative gene

Test Interpretation

See [Genes Tested](#) table for genes included in the panel.

Tests to Consider

[Primary Antibody Deficiency Panel, Sequencing and Deletion/Duplication 2011156](#)

Method: Massively Parallel Sequencing/Exonic Oligonucleotide-based CGH Microarray

Determine the genetic etiology of a primary antibody deficiency in affected individuals.

See [Related Tests](#) for initial screening tests for immunodeficiency and mutation testing for a known familial pathogenic variant.



Clinical Sensitivity

- Agammaglobulinemia, 90%¹
- Hyper-IgM syndrome, 75-80%²
- CVID, 20%³

Indications for Ordering

Determine the genetic etiology of a primary antibody deficiency in affected individuals.

Limitations

- A negative result does not exclude a PAD syndrome.
- Diagnostic errors can occur due to rare sequence variations.
- Interpretation of this test result may be impacted if the individual has had an allogeneic stem cell transplantation.
- The following will not be evaluated:
 - Variants outside the coding regions and intron-exon boundaries of the targeted genes
 - Regulatory region variants and deep intronic variants
 - Breakpoints of large deletions/duplications
 - Deletions/duplications in *ADA2*, *ATP6AP1*, *CARD11*, *CD27*, *CD70*, *CDCA7*, *CTLA4*, *CXCR4*, *DNMT3B*, *GATA2*, *HELLS*, *IKBKG*, *IL21*, *IL21R*, *INO80*, *KDM6A*, *KMT2D*, *LRBA*, *LRRC8A*, *MALT1*, *MAP3K14*, *MOGS*, *NFKB1*, *PIK3CD*, *PIK3CG*, *PIK3R1*, *PLCG2*, *PRKCD*, *RNF168*, *SH2D1A*, *TCF3*, *TNFSF12*, *TRNT1*, *TTC37*, *XIAP*, *ZBTB24*
 - Noncoding transcripts
 - Translocations
 - The following exon is not sequenced due to technical limitations of the assay:
 - *CXCR4* (NM_001348056) 2
- The following may not be detected:
 - Deletions/duplications/insertions of any size by massively parallel sequencing
 - Deletions/duplications less than 1 kb in the targeted genes by array
 - Some variants due to technical limitations in the presence of pseudogenes, repetitive, or homologous regions
 - Low-level somatic variants
 - Single exon deletions/duplications in the following exons:

Gene	Exon(s)
<i>ADA</i>	(NM_000022) 1
<i>AICDA</i>	(NM_020661) 5
<i>BTK</i>	(NM_000061) 11; (NM_001287344) 1
<i>CD79B</i>	(NM_000626) 2
<i>CR2</i>	(NM_001006658) 1
<i>DCLRE1C</i>	(NM_001033855) 4, 6, 8; (NM_001289076) 3
<i>IGLL1</i>	(NM_020070) 2, 3; (NM_152855) 2
<i>NFKB2</i>	(NM_001077494) 21
<i>RAC2</i>	(NM_002872) 2
<i>TNFRSF13B</i>	(NM_012452) 2



Gene	Exon(s)
<i>TNFRSF13C</i>	(NM_052945) 1

Analytical Sensitivity

For massively parallel sequencing:

Variant Class	Analytical Sensitivity (PPA) Estimate ^a (%)	Analytical Sensitivity (PPA) 95% Credibility Region ^a (%)
SNVs	99.2	96.9-99.4
Deletions 1-10 bp	93.8	84.3-98.2
Deletions 11-44 bp	100	87.8-100
Insertions 1-10 bp	94.8	86.8-98.5
Insertions 11-23 bp	100	62.1-100

^aGenes included on this test are a subset of a larger methods-based validation from which the PPA values are derived.

bp, base pairs; PPA, positive percent agreement; SNVs, single nucleotide variants

Genes Tested

Gene	MIM Number	Disorder	Inheritance
<i>ADA</i>	608958	SCID, T-cell/B-cell/NK-cell negative	AR
<i>ADA2</i>	607575	Polyarteritis nodosa Vasculitis Hypogammaglobulinemia	AR
<i>AICDA</i>	605257	Immunodeficiency with hyper-IgM, type 2	AR
<i>ATM</i>	607585	Ataxia-telangiectasia	AR
<i>ATP6AP1</i>	300197	Immunodeficiency 47	XL
<i>BLNK</i>	604515	Agammaglobulinemia	AR
<i>BTK</i>	300300	XL agammaglobulinemia	XL
<i>CARD11</i>	607210	Immunodeficiency 11	AR
		B-cell expansion with NFKB and T-cell energy Immunodeficiency 11B with atopic dermatitis	AD
<i>CD19</i>	107265	CVID	AR

AD, autosomal dominant; AR, autosomal recessive; SCID, severe combined immunodeficiency; XL, X-linked



Gene	MIM Number	Disorder	Inheritance
<i>CD27</i>	186711	Lymphoproliferative syndrome 2	AR
<i>CD40</i>	109535	Immunodeficiency with hyper-IgM, type 3	AR
<i>CD40LG</i>	300386	Immunodeficiency with hyper-IgM, type 1	XL
<i>CD70</i>	602840	Hypogammaglobulinemia EBV susceptibility	AR
<i>CD79A</i>	112205	Agammaglobulinemia 3	AR
<i>CD79B</i>	147245	Agammaglobulinemia 6	AR
<i>CD81</i>	186845	CVID	AR
<i>CDCA7</i>	609937	Immunodeficiency-centromeric instability-facial anomalies syndrome	AR
<i>CR2</i>	120650	CVID	AR
<i>CTLA4</i>	123890	Autoimmune lymphoproliferative syndrome, type V	AD
<i>CXCR4</i>	162643	Whim syndrome	AD
<i>DCLRE1C</i>	605988	SCID with sensitivity to ionizing Omenn syndrome	AR
<i>DNMT3B</i>	602900	Immunodeficiency-centromeric instability-facial anomalies syndrome	AR
<i>GATA2</i>	137295	Immunodeficiency 21	AD
<i>HELLS</i>	603946	Immunodeficiency-centromeric instability-facial anomalies syndrome	AR
<i>ICOS</i>	604558	CVID	AR
<i>IGHM</i>	147020	Agammaglobulinemia	AR
<i>IGLL1</i>	146770	Agammaglobulinemia	AR
<i>IKBKG</i>	300248	Ectodermal dysplasia, hypohidrotic, with immune deficiency Ectodermal dysplasia, anhidrotic, with immunodeficiency, Immunodeficiency without anhidrotic ectodermal dysplasia Immunodeficiency 33 Invasive pneumococcal disease, recurrent isolated, 2	XL
<i>IKZF1</i>	603023	CVID	AD
<i>IL21</i>	605384	CVID	AR



Gene	MIM Number	Disorder	Inheritance
<i>IL21R</i>	605383	IgE responsiveness, atopic	AD
		Immunodeficiency 56	AR
<i>INO80</i>	610169	Hyper-IgM	AR
<i>IRF2BP2</i>	615332	CVID	AD
<i>KDM6A</i>	300128	Kabuki syndrome 1	AD
		Kabuki syndrome 2	XL
<i>KMT2D</i>	602113	Kabuki syndrome 1	AD
<i>LRBA</i>	606453	CVID with autoimmunity	AR
<i>LRRC8A</i>	608360	Agammaglobulinemia	AD
<i>MALT1</i>	604860	Immunodeficiency 12	AR
<i>MAP3K14</i>	604655	Hypogammaglobulinemia	AR
<i>MOGS</i>	601336	Congenital disorder of glycosylation, type IIB	AR
<i>MS4A1</i>	112210	CVID	AR
<i>NBN</i>	602667	Nijmegen breakage syndrome	AR
<i>NFKB1</i>	164011	CVID	AD
<i>NFKB2</i>	164012	CVID	AD
<i>NFKBIA</i>	164008	Ectodermal dysplasia, anhidrotic, with T-cell immunodeficiency	AD
<i>PIK3CD</i>	602839	Immunodeficiency 14	AD
<i>PIK3CG</i>	601232	Hyper-IgM	AD
<i>PIK3R1</i>	171833	Agammaglobulinemia	AR AD
<i>PLCG2</i>	600220	Autoinflammation, antibody deficiency, and immune dysregulation, <i>PLCG2</i> associated	AD
<i>PRKCD</i>	176977	Autoimmune lymphoproliferative syndrome, type III	AR
<i>RAC2</i>	602049	Neutrophil immunodeficiency syndrome	AD



Gene	MIM Number	Disorder	Inheritance
RAG1	179615	Combined cellular and humoral immune defects with granulomas SCID, T-cell negative, B-cell negative, NK-cell positive Omenn syndrome Alpha/beta T-cell lymphopenia with gamma/delta T-cell expansion, severe	AR
RAG2	179616	Combined cellular and humoral immune defects with granulomas SCID, T-cell/B-cell negative, NK-cell positive Omenn syndrome	AR
RNF168	612688	Riddle syndrome	AR
SH2D1A	300490	Lymphoproliferative syndrome, X-linked, 1	XL
STAT3	102582	Hyper-IgE recurrent infection syndrome	AD
TCF3	147141	Agammaglobulinemia 8	AD
TNFRSF13B	604907	CVID Immunoglobulin A deficiency	AR
TNFRSF13C	606269	CVID	AR
TNFSF12	602695	Hypogammaglobulinemia	AD
TRNT1	612907	Sideroblastic anemia with B-cell immunodeficiency, periodic fevers	AR
TTC37	614589	Trichohepatoenteric syndrome 1	AR
UNG	191525	Immunodeficiency with hyper-IgM syndrome	AR
VAV1	164875	CVID	AD
XIAP	300079	X-linked lymphoproliferative syndrome	XL
ZBTB24	614064	Immunodeficiency-centromeric instability-facial anomalies syndrome 2	AR

AD, autosomal dominant; AR, autosomal recessive; SCID, severe combined immunodeficiency; XL, X-linked

References

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Additional Resources



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Related Information

[Agammaglobulinemia](#)

[Common Variable Immune Deficiency Syndromes - CVID](#)

[Primary Immunodeficiency Diseases - Immunoglobulin Disorders](#)

[Inherited T-Cell Deficiency Disorders](#)

[Neutropenia](#)

[Severe Combined Immunodeficiencies - SCID](#)

[Immunodeficiency Evaluation for Chronic Infections in Infants and Children Testing Algorithm](#)

[Immunodeficiency Evaluation for Chronic Infections in Adults and Older Children Testing Algorithm](#)

Related Tests

[CBC with Platelet Count and Automated Differential 0040003](#)

Method: Automated Cell Count/Differential

[Lymphocyte Subset Panel 7 - Congenital Immunodeficiencies 0095899](#)

Method: Quantitative Flow Cytometry

[B Cell Subset Analysis 3002216](#)

Method: Flow Cytometry

[Lymphocyte Antigen and Mitogen Proliferation Panel 0096056](#)

Method: Cell Culture

[Lymphocyte Proliferation, Antigen-Mitogen Panel by Flow Cytometry \(24-Hr Critical Room Temp\) 3001319](#)

Method: Cell Culture/Flow Cytometry

[Immunoglobulins \(IgA, IgG, IgM\), Quantitative 0050630](#)

Method: Quantitative Immunoturbidimetry

[Immunoglobulin G Subclasses \(1, 2, 3, 4\) 0050577](#)

Method: Quantitative Immunoturbidimetry

[Familial Mutation, Targeted Sequencing 2001961](#)

Method: Polymerase Chain Reaction/Sequencing

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