

Agammaglobulinemia Panel

Indications for Ordering

Confirm suspected agammaglobulinemia in individual with clinical symptoms

Test Description

- Targeted capture of all coding exons and intron/exon boundaries followed by massively parallel sequencing
 - Reported variants are confirmed by Sanger sequencing
- Deletion/duplication analysis by tiled, custom-designed comparative genomic hybridization (CGH) array

Tests to Consider

Primary test

[Agammaglobulinemia Panel, Sequencing \(9 Genes\) and Deletion/Duplication \(6 Genes\) 2011151](#)

- Preferred test for individuals with clinical phenotype of agammaglobulinemia

Related tests

Initial screening for immunodeficiency

- [CBC with Platelet Count and Automated Differential 0040003](#)
- [Lymphocyte Subset Panel 7 – Congenital Immunodeficiencies 0095899](#)
- [Lymphocyte Antigen and Mitogen Proliferation Panel with Cytokine Response 2013117](#)
- [Immunoglobulins \(IgA, IgG, IgM\), Quantitative 0050630](#)
- [Familial Mutation, Targeted Sequencing 2001961](#)

Initial screening for suspected primary agammaglobulinemia

- [Bruton Tyrosine Kinase \(BTK\) Protein Expression by Flow Cytometry 2012002](#)

Disease Overview

Prevalence

- X-linked agammaglobulinemia – ~1/250,000-700,000
- Autosomal inherited agammaglobulinemia – rare

Age of onset – usually within the first 2 years of life

Symptoms

- Recurrent or chronic infections
 - Encapsulated pyogenic bacteria
 - Enteroviruses
- Neutropenia
- Pneumonia/empyema
- Gastroenteritis
- Otitis media
- Meningitis
- Sepsis

Genetics

Genes – see table for genes tested and for gene-specific information

Mutations

- Mutations in multiple genes appear to cause overlapping phenotypes for agammaglobulinemia
- Other genetic and/or environmental factors may influence severity of clinical phenotype

Test Interpretation

Clinical sensitivity – ~90%

Results

- Positive
 - Two pathogenic mutations on opposite chromosomes detected in a gene with autosomal recessive (AR) inheritance
 - Confirms diagnosis of agammaglobulinemia
 - Single pathogenic mutation detected in an X-linked gene in males or in an autosomal dominant (AD) gene in males or females
 - Confirms diagnosis of agammaglobulinemia
 - Detection of one pathogenic mutation in gene with AR inheritance
 - Individual is a carrier
 - Detection of one pathogenic mutation in an X-linked gene in females
 - Individual is a carrier
- Negative – no pathogenic mutation detected
 - Reduces, but does not exclude, a diagnosis of agammaglobulinemia
- Inconclusive – variants of uncertain clinical significance may be identified

Limitations

- Not determined or evaluated
 - Mutations in genes not included on the panel
 - Deep intronic and regulatory region mutations
 - Breakpoints for large deletions/duplications
 - Translocations
- Deletions/duplications will not be detected in
 - *LRRC8A* gene
 - *PIK3R1* gene
 - *SH2D1A* gene
- Small deletions or insertions may not be detected

- Diagnostic errors can occur due to rare sequence variations
- Lack of detectable gene mutation does not exclude diagnosis of agammaglobulinemia

Reference

Conley ME, et al. Primary B cell immunodeficiencies: comparisons and contrasts. *Annu Rev Immunol.* 2009;27:199-227

Gene Symbol	Gene Name	NM #	OMIM #	Phenotype/Disorder	Inh.*
<i>BLNK</i>	B-cell linker	013314	604515	Agammaglobulinemia	AR
<i>BTK</i>	Bruton agammaglobulinemia tyrosine kinase	000061	300300	Agammaglobulinemia and isolated growth hormone deficiency X-linked agammaglobulinemia	XL
<i>CD79A</i>	CD79A antigen, Iga	001783	112205	Agammaglobulinemia	AR
<i>CD79B</i>	CD79B molecule, Igb	000626	147245	Agammaglobulinemia	AR
<i>IGHM</i>	Immunoglobulin heavy constant mu	cDNA:X17115	147020	Agammaglobulinemia	AR
<i>IGLL1</i>	Immunoglobulin lambda-like polypeptide 1, 15	020070	146770	Agammaglobulinemia	AR
<i>LRRC8A</i>	Leucine rich repeat containing 8 family, member A	019594	608360	Agammaglobulinemia	AD
<i>PIK3R1</i>	Phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	181523	171833	Agammaglobulinemia	AR
<i>SH2D1A</i>	SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	002351	300490	X-linked lymphoproliferative syndrome	XL

*Inh. = inheritance, AD = autosomal dominant, AR = autosomal recessive, XL = X-linked