

# Autism and Intellectual Disability Comprehensive Panel

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Autism spectrum disorder (ASD) and intellectual disability (ID) represent a neurodevelopmental continuum with significant comorbidity and overlapping etiologies. ASD is classified by varying degrees of social impairment, communication limitations, repetitive behaviors, and/or restricted interests. Symptoms typically present by age 3. ID is classified by broad impairment in cognitive and adaptive functioning, typically with an IQ below 70, and presents before age 18. A global developmental delay (DD) diagnosis often precedes a diagnosis of ID, as cognitive skill or IQ cannot be reliably assessed prior to age 6. Those with severe DD diagnosed before age 6 are most likely to develop ID.

ASD and IDs are often comorbid and may be a feature in various genetic syndromes associated with chromosomal copy number variants (CNVs), fragile X syndrome, and inborn errors of metabolism (IEMs). Literature suggests ~10-15% of individuals with ASD and/or ID have detectable chromosomal abnormalities, CNVs, and fragile X syndrome.

## Disease Overview

### Indications for Ordering (Testing)

- Comprehensive evaluation of an individual with ASD and/or ID (with or without comorbidities)
- Assist with decisions about treatment and management of an individual with ASD and/or ID
- Confirmation of a clinical diagnosis of ASD and/or ID

### Heritability

- ASD: ~90%
- ID: ~50%

### Prevalence

- ASD: ~1/68 (1/42 males, 1/189 females)
- ID: ~1/100
- ASD and ID combined incidence: ~1/250

## Test Limitations

Autism and intellectual disability panel testing limitations:

- Other etiologies of ASD and ID, such as single gene disorders, may not be identified
- Uninformative results may occur due to variants of uncertain clinical significance
- Results may not predict disorder severity

Cytogenomic SNP microarray limitations:

- The genome-wide resolution is approximately 25-50 kb for copy number changes and approximately 3 Mb for regions of homozygosity
- Testing does not detect
  - CNVs below the limit of resolution of this platform
  - Sequence-level variants (mutations) including point mutations and indels
  - Low-level mosaicism (generally, less than 20-30%)
  - Balanced chromosomal rearrangements (translocations, inversions, and insertions)
  - Genomic imbalance in repetitive DNA regions (centromeres, telomeres, segmental duplications, and acrocentric chromosome short arms)
  - Mitochondrial DNA alterations

## Featured ARUP Testing

### [Autism and Intellectual Disability Comprehensive Panel 2014314](#)

**Method:** Tandem Mass Spectrometry/Electrophoresis/Spectrophotometry/ Gas Chromatography-Mass Spectrometry/Liquid Chromatography-Tandem Mass Spectrometry/Quantitative Liquid Chromatography-Tandem Mass Spectrometry, Genomic Microarray (Oligo-SNP Array), Polymerase Chain Reaction/Capillary Electrophoresis

- Comprehensive genetic evaluation of individuals with ASD and/or ID
- Includes cytogenomic microarray, fragile X methylation analysis, and comprehensive metabolic disorders analysis

### Test Components

May be ordered separately

### [Cytogenomic SNP Microarray 2003414](#)

**Method:** Genomic Microarray (Oligo-SNP Array)

- Preferred first-tier test for DD, ID, ASD, and multiple anomalies
- Testing is performed on peripheral blood
- CytoScan HD platform (Thermo Fisher Scientific) to detect copy number changes, including deletions and duplications as well as copy-neutral regions of homozygosity

### [Cytogenomic SNP Microarray Buccal Swab 2006267](#)

**Method:** Genomic Microarray (Oligo-SNP Array)

- Preferred first-tier test for DD, ID, ASD, and multiple anomalies
- Testing is performed on buccal sample

### [Fragile X \(FMR1\) with Reflex to Methylation Analysis 2009033](#)

**Method:** Polymerase Chain Reaction (PCR)/Capillary Electrophoresis

- Preferred test to diagnose fragile X syndrome and carrier screening in individuals with a positive family history
- If fragile X testing detects a CGG repeat of 100 or greater by polymerase chain reaction and capillary electrophoresis, methylation analysis will be added

### [Autism and Intellectual Disability Metabolic Panel 2014312](#)

**Method:** Tandem Mass Spectrometry/Electrophoresis/Spectrophotometry/ Gas Chromatography-Mass

#### Fragile X testing limitations:

- Estimated size is not provided for full mutations with >200 repeats
- Rare mutations in *FMR1* unrelated to trinucleotide expansion will not be detected
- Diagnostic errors can occur due to rare sequence variations

Spectrometry/Liquid Chromatography-Tandem Mass Spectrometry  
Evaluate for ASD, DD, and/or ID in individuals who have had negative *FMR1* (fragile X) and cytogenomic SNP microarray testing

## Additional Resources

American Psychiatric Association. *Diagnostic and Statistical Manual of Mental Disorders*. 5th ed. American Psychiatric Publishing; 2013.

U.S. Department of Health and Human Services, Centers for Disease Control and Prevention. [Autism and Developmental Disabilities Monitoring \(ADDM\) Network](#). [Last reviewed: Aug 2019; Accessed: Feb 2020]

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Karam SM, Riegel M, Segal SL, et al. [Genetic causes of intellectual disability in a birth cohort: a population-based study](#). *Am J Med Genet A*. 2015;167(6):1204-1214.

Kaufman L, Ayub M, Vincent JB. [The genetic basis of non-syndromic intellectual disability: a review](#). *J Neurodev Disord*. 2010;2(4):182-209.

Michelson DJ, Shevell MI, Sherr EH, et al. [Evidence report: genetic and metabolic testing on children with global developmental delay: report of the Quality Standards Subcommittee of the American Academy of Neurology and the Practice Committee of the Child Neurology Society](#). *Neurology*. 2011;77(17):1629-1635.

Rauch A, Hoyer J, Guth S, et al. [Diagnostic yield of various genetic approaches in patients with unexplained developmental delay or mental retardation](#). *Am J Med Genet A*. 2006;140(19):2063-2074.

Sandin S, Lichtenstein P, Kuja-Halkola R, et al. [The familial risk of autism](#). *JAMA*. 2014;311(17):1770-1777.

Sempere A, Arias A, Farré G, et al. [Study of inborn errors of metabolism in urine from patients with unexplained mental retardation](#). *J Inherit Metab Dis*. 2010;33(1):1-7.

South ST, Lee C, Lamb AN, et al. [ACMG standards and guidelines for constitutional cytogenomic microarray analysis, including postnatal and prenatal applications: revision 2013](#). *Genet Med*. 2013;15(11):901-909.

van Karnebeek CDM, Stockler S. [Treatable inborn errors of metabolism causing intellectual disability: a systematic literature review](#). *Mol Genet Metab*. 2012;105(3):368-381.

## Related Information

[Alpha-Iduronidase Enzyme Activity in Leukocytes](#)

[Creatine Disorders Panel Testing](#)

[Laboratory Testing for Developmental Delay, Intellectual Disability, and Autism Spectrum Disorder](#)

[Testing for Genetic Syndromes Related to Developmental Delay \(DD\), Intellectual Disability \(ID\), and Autism Spectrum Disorder \(ASD\)](#)

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