

## Autism and Intellectual Disability Comprehensive Panel

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

### Tests to Consider

#### Autism and Intellectual Disability Comprehensive Panel 2014314

**Method:** Tandem Mass Spectrometry, Electrophoresis/Spectrophotometry, Gas Chromatography/Mass Spectrometry, Liquid Chromatography/Tandem Mass Spectrometry, and 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/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

- 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
- For further review, see [Additional Technical Information](#) for Cytogenomic SNP Microarray 2003414

### Autism and Intellectual Disability Metabolic Panel 2014312

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

See [Related Tests](#)

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
- For further review, see [Additional Technical Information](#) for Fragile X (*FMR1*) with Reflex to Methylation Analysis 2009033

## Additional Resources

American Psychiatric Association. Diagnostic and Statistical Manual of Mental Disorders. 5th ed. Arlington, VA: 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]

Campistol J, Díez-Juan M, Callejón L, et al. [Inborn error metabolic screening in individuals with nonsyndromic autism spectrum disorders](#). *Dev Med Child Neurol*. 2016;58(8):842-847. PubMed

Ghaziuddin M, Al-Owain M. [Autism spectrum disorders and inborn errors of metabolism: an update](#). *Pediatr Neurol*. 2013;49(4):232-236. PubMed

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. PubMed

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

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. PubMed

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. PubMed

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

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

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. PubMed

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. PubMed

## Related Information

[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 \(ASD\)](#)

## Related Tests

**Amino Acids Quantitative by LC-MS/MS, Plasma 2009389**

**Method:** Quantitative Liquid Chromatography/Tandem Mass Spectrometry

**Creatine Disorders Panel, Urine 2002333**

**Method:** Liquid Chromatography-Tandem Mass Spectrometry (LC-MS/MS)

**Creatine Disorders Panel, Serum or Plasma 2002328**

**Method:** Liquid Chromatography-Tandem Mass Spectrometry (LC-MS/MS)

**Organic Acids, Urine 0098389**

**Method:** Gas Chromatography-Mass Spectrometry (GC-MS)

**Mucopolysaccharides Screen - Electrophoresis and Quantitation, Urine 0081352**

**Method:** Electrophoresis/Spectrophotometry

**Acylcarnitine Quantitative Profile, Plasma 0040033**

**Method:** Tandem Mass Spectrometry

ARUP Laboratories is a nonprofit enterprise of the University of Utah and its Department of Pathology, 500 Chipeta Way, Salt Lake City, UT 84108  
(800) 522-2787 | (801) 583-2787 | aruplab.com | arupconsult.com  
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