



ExomeNext[®] ExomeReveal[™]

Genetic Testing for
Rare Disease and
Neurodevelopmental
Disorders



Help families end the diagnostic odyssey.

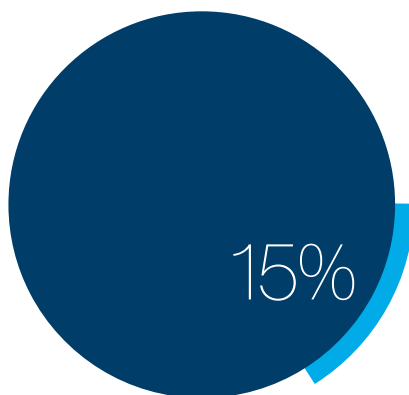
Whole exome sequencing (WES) is a comprehensive testing approach recommended for the evaluation of:¹⁻⁴

- Multiple congenital anomalies (MCA)
- Autism spectrum disorder (ASD)
- Developmental delay (DD)
- Unexplained epilepsy
- Intellectual disability (ID)
- Cerebral palsy (CP)

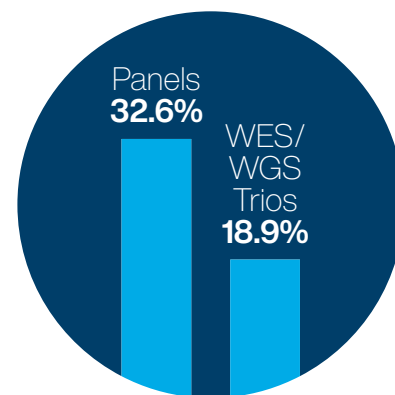
Exome first: More answers. Clearer results.



Up to 1 in 3 patients will have diagnostic results from exome testing compared to 15–20% from chromosomal microarray³



Additional findings on exome that would have been missed by panel testing⁵



Lower rates of variants of uncertain significance (VUS) in exome/genome⁶

Proactive Reanalysis at No Additional Cost

All exome sequencing tests depend on our current understanding of gene-disease relationships and variant classifications, and new discoveries are made every month. Through our Patient for Life™ program, our team proactively identifies new discoveries, reviews exome results, and notifies clinicians about new diagnostic findings—indefinitely and at no additional cost.

How Patient for Life Works:



1. Research Review

Ambry's clinical scientists review the latest findings on gene-disease relationships and updates to variant classifications.



2. Continuous Reanalysis

Patients' data are continuously analyzed based on new scientific findings.



3. Revised Reporting

Ordering providers receive a fully updated report detailing the reclassification and outreach from a genetic counselor.

Increased diagnostic yield

→ 1 in 20

5% of patients who initially test negative on exome will have a diagnosis identified later through Patient for Life⁷

We do the work for you

With Patient for Life, reanalysis is always on for new gene-disease associations and variant classifications. Therefore, you don't need to request a reanalysis in most cases.

Manual reanalysis requests are helpful if there are updates to the family history or major phenotypic changes in the patient—new signs and symptoms that could provide clues to a diagnosis.

Equitable access to genetic discovery



Healthcare disparities because of race and ethnicity impact genetic testing. Our data show that African American/Black patients were consistently left behind in exome testing, with lower diagnostic yields and provider-initiated reanalysis rates. However, they were most likely to have a reclassification when reanalysis was performed.⁸ Patient for Life ensures equitable access to new diagnostic information.

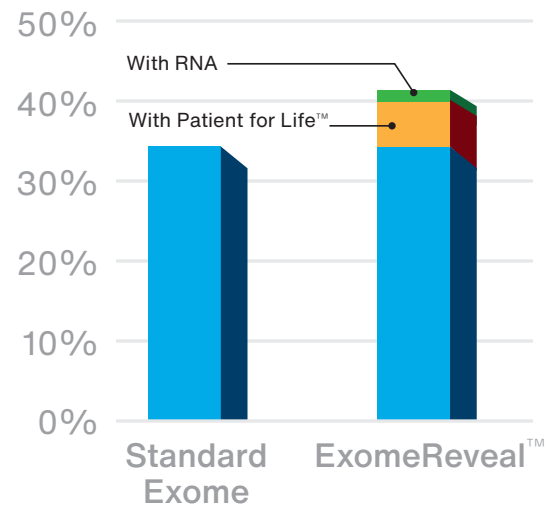
Maximize the Diagnostic Yield Over Standard Exomes

Introducing

ExomeReveal™

ExomeReveal adds the power of RNA analysis to exome testing.

- Provides an innovative, multi-omic approach to rare disease
- Expected to impact 1 in 50 patients¹⁰
- No additional cost to the patient



+20%

Increase in diagnostic yield compared to standard exomes when combined with Patient for Life^{7, 9, 10}

RNA Experience that Matters



8+ years performing RNA analysis and interpreting splice variants



30+ RNA experts in-house at Ambry



Our RNA experts serve on multiple independent variant working groups

Let us be your trusted partner

All exome tests utilize Ambry's Classifi™ program, a proprietary, knowledge-driven engine for gene classification, variant analysis & interpretation, and reporting. The Classifi program delivers the highest quality test results and ensures we leave no stone unturned in getting answers for you and your patients.

Ambry
Classifi™

Advanced Diagnostic Testing for Potentially Life-Changing Answers

ExomeNext®

PREFERRED

Trio
(with both parents)

ACCEPTABLE

Duo
(with one parent)

Proband Only

6–8 weeks turnaround

Blood, saliva and
buccal samples accepted

ExomeReveal™

RNA analysis performed for eligible DNA variants expected to impact splicing.



- Requires EDTA (DNA) and PAXgene (RNA) specimens.
- Reported 3–4 weeks after initial DNA results.

Did you know?



Ambry was the first commercial lab to offer whole exome sequencing in 2011.

Technical Test Performance

- >97% of the exome covered with a minimum depth of coverage of 20X
- Detects gross deletions and duplications ≥ 5 exons
- Within mitochondrial DNA, >5% heteroplasmy is detected

References:

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