Genomic Unity® Case Study

Shifting the diagnostic paradigm with a whole genome platform

Compound heterozygous variants in an sncRNA explains multisystemic symptoms in 4-year-old male

Clinical presentation

A 4-year-old male presented with a complex medical history including:

- Mitochondrial complex I deficiency
- Global developmental delay
- Spasticity
- Congenital hypotonia
- Persistent left superior vena cava
- Mild oropharyngeal dysphagia

Results and interpretation

Variantyx Genomic Unity[®] Whole Genome Analysis identified a likely pathogenic, paternally inherited SNV and a pathogenic, maternally inherited indel, both within the *RNU7-1* gene.

Both variants have been experimentally demonstrated to alter U7 snRNP function.

Diagnosis: Aicardi-Goutieres syndrome 9

Previous genetic testing

Multiple tests were performed including:

- Chromosomal microarray (CMA)
- SMN1/2 testing
- Cerebral palsy panel
- Mitochondrial sequencing & del/dup analysis identified non-diagnostic, heteroplasmic variants
- Whole exome sequencing (WES), reanalysis

Genomic Unity® Whole Genome Analysis was ordered because of its ability to identify all major variant types in a single test.

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Uniform data from WGS clearly shows the inherited SNV and indel.



The Variantyx difference

Why were these compound heterozygous sequence variants detected by Genomic Unity® Whole Genome Analysis, and not detected by other tests?

The variants fall within an sncRNA gene. Other tests focus predominantly, if not exclusively, on protein-coding genes.

While exomes have the potential to include sncRNA genes, a survey of commercial exome probe sets shows that only 1 in 7 targets RNU7-1.

> IDT - xGen Exome Research Panel V2 Probes IDT - xGen Exome Research Panel V2 Target Regions Roche - KAPA HyoerExome Capture Probe Footprint Roche - KAPA HyoerExome Primary Target Regions Roche - SeqCap EZ MedExome + Mito Capture Probe Footprint Roche - SeqCap EZ MedExome + Mito Empirical Target Regions Agilent - SureSelect Clinical Research Exome V2 Covered by Probes Agilent - SureSelect Clinical Research Exome V2 Target Regions Agilent - SureSelect All Exon V6 +UTR r2 Covered by Probes

> > Agilent - SureSelect All Exon V7 Covered by Probes

Agilent SureSelevt All Exon V7 Target Regions Twist - Comprehensive Exome Panel Target Regions Twist - Exome 2.0 Panel Target Regions GENCODE V43 (3 items filtered out)

RNU7-1

Variantyx genome analysis provides truly comprehensive mitochondrial and nuclear gene coverage, including analysis of sncRNAs. Ordering Variantyx genome analysis as a first line test would have shortened the diagnostic odyssey while significantly saving healthcare costs by eliminating multiple step-wise tests.

Variantyx tests that would have identified this variant

Genomic Unity[®] Whole Genome Analysis | Genomic Unity[®] Exome Plus Analysis | Genomic Unity[®] Exome Analysis



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