Laboratory-driven Exome Reanalysis Increases Diagnostic Yield and Decreases Burden on Clinicians



Meghan Towne, <u>Melissa Holman</u>, Carolyn Horton, Catherine Schultz, Brooklynn Gasser, Grace VanNoy, Jessica Gage

Contact: mholman@ambrygen.com

Ambry Genetics

BACKGROUND

- Exome sequencing (ES) data can be reanalyzed as understanding of genetic contributions to disease grows
- Reanalysis increases ES diagnostic yield over time
 - Mostly due to gene-disease relationships (GDR)

OBJECTIVE: Assess the outcomes of proactive vs. reactive ES reanalysis processes

STUDY METHODS

Reviewed 10 years of ES for neurological indications (n=8539)







Neurodevelopmental

Epilepsy

Neuromuscular

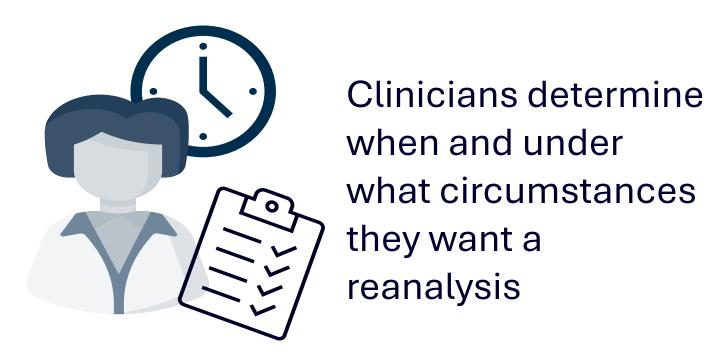
74% of cohort 9% of cohort

17% of cohort

Compared reanalysis outcomes and evidence used for reclassifications based on the reanalysis initiation factor

METHODS FOR EXOME REANALYSIS

Provider requested





Reanalysis is requested from the lab

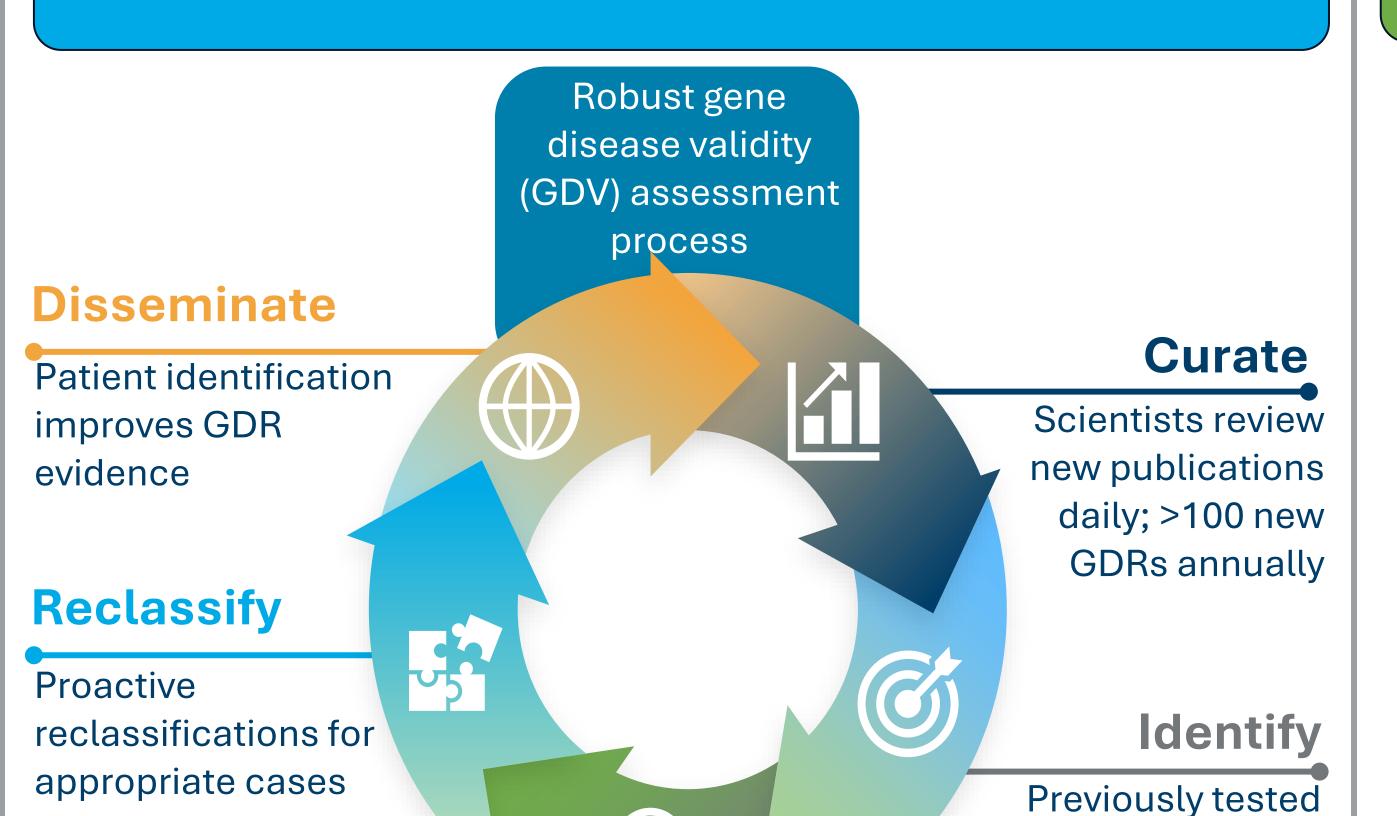
 Labs may have differing criteria, costs, and timeline for reanalysis



Updated report documenting reanalysis issued

- Time-based
- Requires laboratory and clinician resources
- Comprehensive review of data at time of reanalysis

Patient for Life



Scientists review each case for clinical overlap

Reanalyze

- Ongoing assessment
- Requires no clinician resources
- Comprehensive, consistent review of data as evidence is published

Family Studies



Variant of Unknown
Significance
reported on testing



Review likelihood that cosegregation studies in available family members would reclassify a variant



New sample collected from family member(s) with single variant analyzed



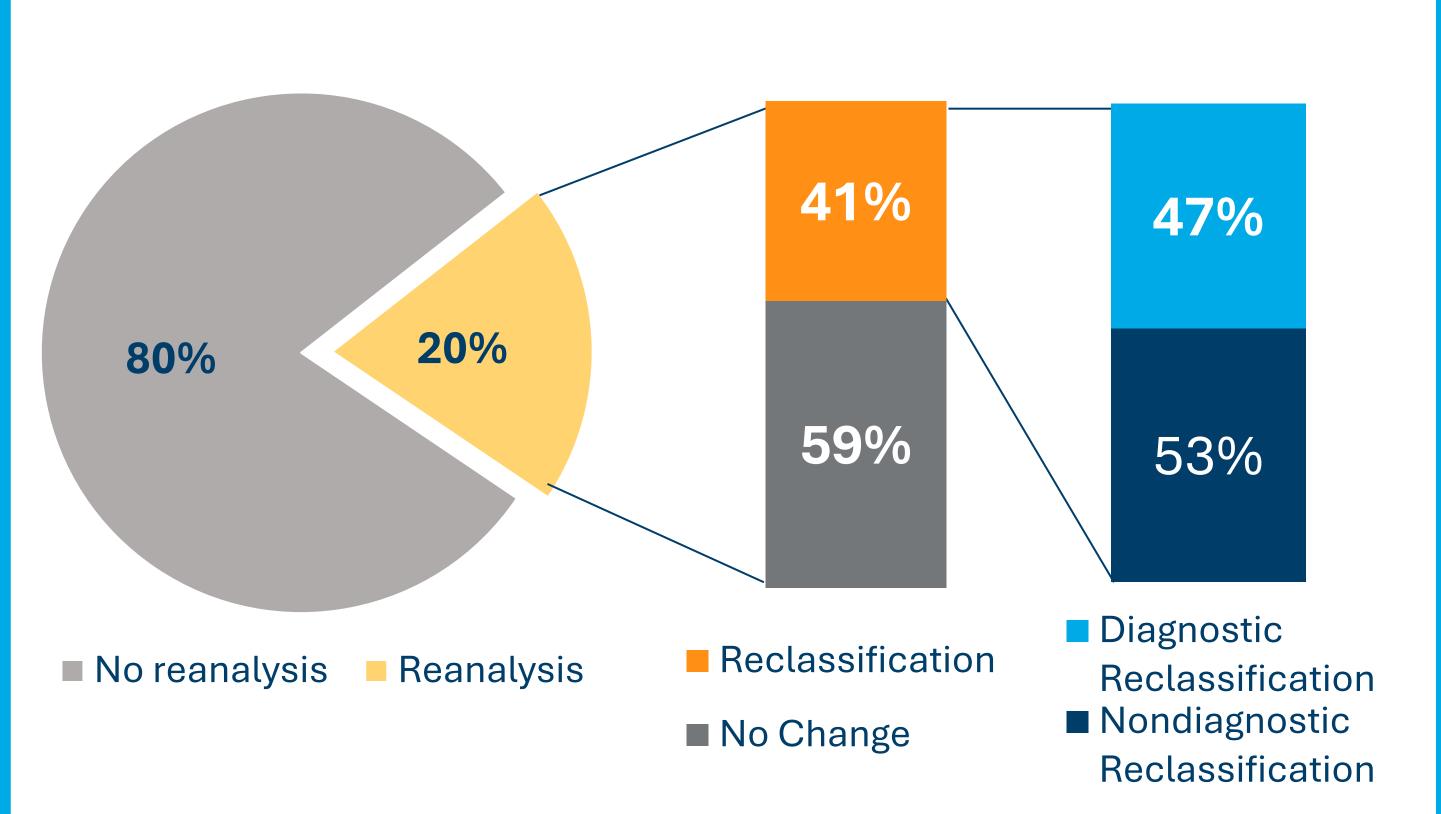
Results incorporated into variant classification and updated report issued

- Limited by available/informative family member(s)
- Requires laboratory and clinician resources
- Limited to just variant in question

RESULTS

1%

FIGURE 1: COHORT REANALYSIS AND RECLASSIFICATION



- 20% of cases (n= 1685) underwent at least one reanalysis during the study period [Figure 1]
 - Of these, 41% (n=694) received a reclassification report
- Increase in overall diagnostic yield (21% vs 25%)
 - 5% of all originally unsolved ES received a diagnostic finding

FIGURE 2: DIAGNOSTIC RECLASSIFICATIONS BY INITIATING FACTOR

15% 84%

exomes with variants in

the new gene of interest

are reassessed

■ Family Studies ■ Provider ■ Patient for Life

- 327 total diagnostic reclassifications (4% of all cases)
- 84% of diagnostic upgrades were through PFL, compared to 15% provider-requested reanalyses, and 1% family studies [Figure 2]

TAKE HOME POINTS

- ES reanalysis increases the diagnostic yield over time
- Patient for Life resulted in higher rates of diagnostic reclassification and initiates when new relevant data is available
- Majority of provider-requested reanalyses resulted in a 'no change' notification, adding work for the provider and laboratory with no clinical benefit