

Exome Reanalysis Request [Internal Test Code: 9270 / 9271]

Complimentary reanalysis of exome sequencing data may be requested once in the absence of new clinical information or at any time when a patient presents with a new phenotype. Reanalysis or amended report(s) will be issued for all reanalysis requests; the report type will depend on whether any new findings are observed. Since reanalysis relies on new data, it is recommended that requests are submitted at least one year after the original report was issued. Turnaround time for a reanalysis request is 4-5 weeks.

Please note that a lab-initiated reclassification report will be issued if a new relevant gene-disease association is identified through Ambry's ongoing gene curation process, even in the absence of a reanalysis request.

To request reanalysis of your patient's exome sequencing data, please complete the information below.

Proband's Name: _____ Date of Birth: _____

Ambry Accession Number: _____ Date of Request: _____

If you would like to include (re)analysis of ACMG Secondary Findings for individual(s) who underwent exome sequencing (i.e. any member of the trio), please provide their information below.

NAME	DATE OF BIRTH

The undersigned person (or representative thereof) ensures he/she is a licensed medical professional authorized to order genetic testing and confirms that the patient has given appropriate consent. I confirm that receipt of these results is medically necessary and that these results may impact medical management for the patient(s).

Medical Professional Name* and Credential: _____

Medical Professional's Organization: _____

Medical Professional Signature: _____

Phone: _____ Fax: _____

**If you are not the original ordering clinician, you will be added to the order as an additional recipient.*

Have there been any changes to the patient's clinical phenotype since initial exome analysis or previous reanalysis request? No Yes

If yes, please list any **new** clinical features or changes to the patient's history since the initial analysis. Please also list any additional genes of interest to be considered for differential diagnoses.