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Genome-wide Sequencing Ontario

CHEO Genetics Diagnostic Laboratory
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Patient name:

Date of Birth (DD-MM-YYYY):

Gender: ☐ *Male* ☐ *Female* *MRN:*

Address:

Telephone #:

Ontario health card #: *Version:*

GENOME-WIDE SEQUENCING REANALYSIS

Ordering physician:

Name: _____

Institution: _____

Address: _____

Phone: _____

Fax: _____

Email address: _____

Copy report to:

Name: _____

Institution: _____

Address: _____

Phone _____

Fax _____

Updated Clinical Information:

Please note any phenotype progression or new/emerging phenotype(s) below

For laboratory use only:

Date (DD-MM-YYYY) | **Time Received**:

_____-_____-_____ | _____h

Order #: _____

Specimen type, amt & # of tubes: _____

Comments:

_____

_____

Reanalysis and Reinterpretation Request:

Previous Analysis Information

Date of previous report (DD-MM-YYYY): _____-_____-_____

Ordering provider: _____

Full Reanalysis:

☐ Routine Reanalysis (≥ 18 months since previous report date)

☐ Priority Reanalysis (<18 months since previous report date)

Must meet at least one of the below criteria:

☐ New and ongoing pregnancy

EDD (DD-MM-YYYY): _____

Relationship to pregnant person: _____

☐ Phenotype progression where results are expected to impact management

☐ New or emerging phenotype(s) expected to impact data analysis

Single Gene Reinterpretation or Analysis:

☐ Variant Reinterpretation

☐ Targeted Gene Analysis - Gene(s): _____

GWS Reanalysis and Reinterpretation requirements:

Eligibility:

Reanalysis is available upon request for all patients ≥ 18 months after the previous report date. If there is a new and ongoing pregnancy, progression of a previously reported phenotype where results are expected to impact management, or new phenotype expected to impact data analysis, reanalysis can be requested ≤ 18 months from the previous report date.

Reinterpretation of a previously reported variant is available upon request ≥ 12 months after the previous report date, when relevant to patient management. If there is evidence supporting a different variant classification than previously reported by CHEO/SickKids, this can be shared with the laboratory at any time and a reinterpretation may be performed.

Targeted gene(s) analysis of the previously generated sequencing data for a gene(s) that was not associated with a disease at time of previous analysis can be performed at any time. Please provide evidence of new disease-gene association.

Clinical information and Supporting Documents:

The following information has been provided for the proband and family:

- Updated phenotypic information
- Updated clinic note(s) and/or letter(s)
- Segregation results (Reinterpretation only)
- Publications supporting a change in variant classification (Reinterpretation only)
- Publications supporting a new disease-gene association (Targeted gene(s) analysis only)