## Variant Confirmation Report

| Sample ID:                        | EO_21_18  |
|-----------------------------------|---|
| Description:                      |   |
| Gene                              | FBN1  |
| Exon                              | 38/66   |
| Sequence Variant                  | ENST00000316623.5:c.4640C>T   |
|                                   | ENSP00000325527.5:p.Thr1547lle  |
| Variant Location (GRCh37)         | 15:48760242   |
| Allele Balance                    | 0.73  |
| Allele Depth (REF,ALT)            | 25,66   |
| Allele Frequency (ESP,ExAC,dbSNP) | - 0.00331 0.00040   |
| Variant Found                     | Υ   |
| Comment                           | This mutation has been asserted as a likely disease-causing muatation in the HGMD database  HGMD Accession: CM145768 HGMD Classification: DM? Lerner-Ellis (2014) Mol Genet Metab 112: 171 PubMed: 24793577  Date of Variant Class Change From DM to DM?: 2014-06-09 00:00:00 Lerner-Ellis (2014) Mol Genet Metab 112: 171 PubMed: 24793577 |

Validation:

Fluidigm Image:

| The following information is for research purpose only. Any decisions made on the information sh<br>by an appropriate responsible clinician who may require further confirmation within a clinical labo |
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