

**INDIVIDUAL LABORATORY REPORT (ILR)\***  
**2024 Exome Sequencing (trio) Postnatal**  
**(neonatal and paediatric)**  
**Version: Final**  
**Participant**



Member of UK NEQAS consortium

Mean Overall Score	
Genotyping	2.00
Interpretation	2.00
Clerical Accuracy	2.00

Conclusions	
Recommendations	None
Performance Statement <sup>2</sup>	Satisfactory

**Thank you for your participation**

\* To be read in conjunction with the EQA Summary Report

1 The maximum score for each category is 2.00

2 There are only two categories of performance for any EQA; "Satisfactory" and "Poor"

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