

Individual Laboratory Report - Lab 1643

DNA SEQUENCING - NGS (Somatic SNVs and indels + matched germline sample) : Whole genome sequencing | 2025

SUMMARY OF YOUR PERFORMANCE IN THIS EQA

Assessment Category	Performance (mean score) ¹
Genotyping	2.00
EQA Result (PASS or POOR)	PASS

CASE SPECIFIC FEEDBACK

Case 1

Assessment Category	Score ²	Comments (& deductions) ³
Genotyping	2.00	Results submitted. Please refer to your individual Data Quality Report and Variant Consensus Analysis Report for details of the assessment. (0)

This Individual Laboratory Report should be read in conjunction with scheme summary report.

Thank you for participating in this year's EQA, we look forward to you participating again next year.

Report approved and authorised by Simon Patton (16 April 2026) on behalf of EMQN.



Dr. Simon Patton (CEO)

Key: ¹ **Green** (Pass), **Red** (Poor performance), NRS (no results submitted), WDS (withdrew from scheme)

² Maximum score is 2.00

³ Deductions from the maximum score

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