

INDIVIDUAL LABORATORY REPORT (ILR) - Lab 1678

SCHEME: LYNCH

SEASON: 2023

Case 1		
Assessment Category	Score ¹	Comments (& deductions ²)
Genotyping	2.00	Version number missing / incorrect / inconsistent for reference sequence (0.00)
Interpretation	2.00	All essential interpretative elements provided (0.00)
Patient Identifiers and Clerical Accuracy	1.30	Failure to provide patient identifiers on each page of the report (0.20) Errors in sample batch number or no sample batch number (0.50) Spelling and typographic error in the body of the text that do not change the meaning of the report (0.00) Incorrect or no pagination (eg X of Y) (0.00) Date of the report is in 2021.

Case 2		
Assessment Category	Score ¹	Comments (& deductions ²)
Genotyping	2.00	Version number missing / incorrect / inconsistent for reference sequence (0.00)
Interpretation	2.00	All essential interpretative elements provided (0.00)
Patient Identifiers and Clerical Accuracy	1.30	Failure to provide patient identifiers on each page of the report (0.20) Errors in sample batch number or no sample batch number (0.50) Incorrect or no pagination (eg X of Y) (0.00)

Case 3		
Assessment Category	Score ¹	Comments (& deductions ²)
Genotyping	2.00	Correct result reported (0.00)
Interpretation	1.80	Failure to provide adequate details of assay limitations eg. Limit of detection (LOD), sensitivity, specificity (0.20) sensitivity of the method is missing
Patient Identifiers and Clerical Accuracy	1.30	Failure to provide patient identifiers on each page of the report (0.20) Errors in sample batch number or no sample batch number (0.50) Incorrect or no pagination (eg X of Y) (0.00)

¹ Maximum score is 2.00

² Deductions from the maximum score

³ Green >= Scheme mean, Orange < Scheme mean, Red Poor performance.

NRS no results submitted. WDS withdrew from scheme

General Comments	Thank you for participating in this year's EQA scheme. We look forward to you joining us in the 2024 scheme.
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SUMMARY OF YOUR PERFORMANCE IN THIS SCHEME

Assessment Category	Performance ³ (mean score)
Genotyping	2.00
Interpretation	1.93
Patient Identifiers and Clerical Accuracy	1.30
Scheme result (SATISFACTORY or POOR)	SATISFACTORY

Report approved and authorised by Simon Patton (13 September 2023) on behalf of EMQN.
Signed:



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² Deductions from the maximum score

³ **Green** >= Scheme mean, **Orange** < Scheme mean, **Red** Poor performance.

NRS no results submitted. WDS withdrew from scheme