

10, Feb 2026

severity	code	details	expression
14 information	business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/ CodeSystem/ GenomicClinicalIndication 0.1.0	[Bundle.entry[5].resource/*ServiceRequest/ null*/.reasonCode[0]]
17 information	code-invalid	None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (http://hl7.org/fhir/ValueSet/ report-codes 4.0.1), and a coding is recommended to come from this value set (codes = https:// fhir.nwgenomics.nhs.uk/ CodeSystem/ IGEAP#R125_Cardiology, https://fhir.nhs.uk/CodeSystem/ England- GenomicTestDirectory#R125.1) Entry 'urn:uuid:fb15ba1e-2d4d-47ab- bd05-450a1b063873' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	[Bundle.entry[6].resource/*DiagnosticReport/ null*/.code]
23 information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[8]]
24 information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[6].resource.resultsInterpreter[0]]
25 information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[6].resource.code.coding[0]]
18 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/ fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http:// terminology.hl7.org/ CodeSystem/v2-0203#NH)	[Bundle.entry[6].resource/*DiagnosticReport/ null*/.subject.identifier.type]
13 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/ fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http:// terminology.hl7.org/ CodeSystem/v2-0203#NH)	[Bundle.entry[5].resource/*ServiceRequest/ null*/.subject.identifier.type]

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15 warning	code-invalid	Unknown Code 'R125' in the CodeSystem 'https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication' version '0.1.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment	[Bundle.entry[5].resource/*ServiceRequest/null*/.reasonCode[0].coding[0].code]
11 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)	[Bundle.entry[4].resource/*Specimen/null*/.subject.identifier.type]
6 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)	[Bundle.entry[3].resource/*DocumentReference/null*/.subject.identifier.type]
1 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)	[Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]