

severity	code	details	expression
32 error	structure	DiagnosticReport.resultsInterpreter.identifier: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/ StructureDefinition/DiagnosticReport 0.1.0)	[Bundle.entry[6].resource.resultsInterpreter[0]]
31 error	structure	DiagnosticReport.resultsInterpreter.type: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/ StructureDefinition/DiagnosticReport 0.1.0)	[Bundle.entry[6].resource.resultsInterpreter[0]]
19 error	invalid	Wrong Display Name 'PANEL : R125.1, R125.2 - Thoracic aortic aneurysm and dissection v1.2' for https:// fhir.nwgenomics.nhs.uk/CodeSystem/ IGEAP#R125_Cardiology. Valid display is 'PANEL: R125.1, R125.2 - Thoracic aortic aneurysm and dissection v1.2' (for the language(s) 'en')	[Bundle.entry[6].resource/*DiagnosticReport/ null*.code.coding[0].display]
16 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]]
0 information informational		A definition could not be found for Canonical URL 'https:// fhir.nwgenomics.nhs.uk/Endpoint/IGENE' but a definition can be found for 'http:// fhir.nwgenomics.nhs.uk/Endpoint/IGENE' - change from https: to http: Entry 'urn:uuid:36e99b75-3a66-43d5- a35c-7916ce5a2c33' isn't reachable by traversing forwards from the	[Bundle.entry[0].resource/*MessageHeader/ null*.source.endpoint]
27 information informational		MessageHeader. Check that this is meant to be included (needed to process the message) 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]]
28 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.basedOn[0]]
29 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]]
30 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]]
21 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)	[Bundle.entry[6].resource/*DiagnosticReport/ null*.code]
22 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (<a href="http://hl7.org/fhir/
ValueSet/identifier-type 4.0.1">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]
17 warning	code-invalid		

severity	code	details	expression
		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]
15 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]
14 warning	code-invalid	Unknown Code 'R125.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory' 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*/.code.coding[0].code]
12 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]
7 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]
2 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]
20 warning	code-invalid	Unknown Code 'R125.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory' 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[6].resource/*DiagnosticReport/ null*/.code.coding[1].code]