

severity	code	details
17 error	structure	The type 'DocumentReference' is not a valid Target for this element (must be one of [CanonicalType[http://hl7.org/fhir/StructureDefinition/Procedure], CanonicalType[http://hl7.org/fhir/StructureDefinition/Observation], CanonicalType[http://hl7.org/fhir/StructureDefinition/DiagnosticReport]]) [Bundle.entry[6].resource/*DiagnosticReport/null*.extension[0].extension[0].value.ofType(Reference)]
18 error	structure	Invalid Resource target type. Found DocumentReference, but expected one of ([Observation, Procedure, DiagnosticReport]) [Bundle.entry[6].resource/*DiagnosticReport/null*.extension[0].extension[0].value.ofType(Reference)]
27 error	processing	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice DiagnosticReport.category:Genetics in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[6].resource.category[0]]
28 error	invalid	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0, Element matches more than one slice - organization, organisation [Bundle.entry[6].resource.performer[0]]
29 error	invalid	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0, Element matches more than one slice - organization, operator [Bundle.entry[6].resource.performer[0]]
20 error	invalid	No definition could be found for URL value 'http://hl7.org/fhir/diagnosticreport-relevant-information-types' Slice 'DiagnosticReport.performer:organisation': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0). Note that other slices are allowed in addition to this required slice The extension http://hl7.org/fhir/5.0/StructureDefinition/extension-DiagnosticReport.supportingInfo is not allowed to be used at this point (this element is [Bundle.entry.resource]; allowed for this version = e:DiagnosticReport) DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required [Bundle.entry[6].resource/*DiagnosticReport/null*.extension[0].extension[1].value.ofType(CodeableConcept).coding[0]]
31 error	structure	[Bundle.entry[6].resource]
32 error	structure	[Bundle.entry[6].resource]
30 information	not-supported	(DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0) [Bundle.entry[6].resource]
14 information	business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.0. This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0 (this may not be a problem, but you should check that it's not intended to match a slice) None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (http://hl7.org/fhir/ValueSet/report-
33 information	informational	[Bundle.entry[6].resource.code.coding[0]]
21 information	code-invalid	[Bundle.entry[6].resource/*DiagnosticReport/null*.code]

severity	code	details
19 warning	not-found	<p>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)</p> <p>A definition for CodeSystem 'http://hl7.org/fhir/diagnosticreport-relevant-information-types' could not be found, so the code cannot be validated</p> <p>Unknown Code 'R125' in the CodeSystem 'https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication' version '0.2.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>
15 warning	code-invalid	[Bundle.entry[5].resource/*ServiceRequest/null*.reasonCode[0].coding]
13 warning	code-invalid	<p>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)</p> <p>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)</p>
11 warning	code-invalid	[Bundle.entry[4].resource/*Specimen/null*.subject.identifier.type]
6 warning	code-invalid	<p>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)</p> <p>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)</p>
1 warning	code-invalid	[Bundle.entry[1].resource/*Patient/null*.identifier[0].type]
22 warning	code-invalid	[Bundle.entry[6].resource/*DiagnosticReport/null*.subject.identifier.type]