

	severity	code	details
18 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)]
19 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)]
29 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]]
30 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]]
21 error	invalid	No definition could be found for URL value 'http://hl7.org/fhir/diagnosticreport-relevant-information-types'	[Bundle.entry[6].resource/*DiagnosticReport/null*.extension[0].extension[1].value.ofType(CodeableConcept).coding[
31 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]]
33 error	structure	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	[Bundle.entry[6].resource]
34 error	structure	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)	[Bundle.entry[6].resource]
32 information	not-supported	DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required (DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0)	[Bundle.entry[6].resource]
15 information	business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.0	[Bundle.entry[5].resource/*ServiceRequest/null*.reasonCode[0]]
35 information	informational	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)	[Bundle.entry[6].resource.code.coding[0]]
23 information	code-invalid	None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (http://hl7.org/fhir/ValueSet/report-	[Bundle.entry[6].resource/*DiagnosticReport/null*.code]

severity	code	details
24 warning	code-invalid	<p>codes 4.0.1), and a coding is recommended to come from this value set (codes = <a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/IGEAP#CCP19v1_R112">https://fhir.nwgenomics.nhs.uk/CodeSystem/IGEAP#CCP19v1_R112</a>, <a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R112.1">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R112.1</a>)</p> <p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
20 warning	not-found	<p>A definition for CodeSystem '<a href="http://hl7.org/fhir/diagnosticreport-relevant-information-types">http://hl7.org/fhir/diagnosticreport-relevant-information-types</a>' could not be found, so the code cannot be validated</p> <p>Unknown Code 'R112' in the CodeSystem '<a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication">https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication</a>' 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>
16 warning	code-invalid	<p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
14 warning	code-invalid	<p>Unknown Code 'R112.1' in the CodeSystem '<a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory</a>' 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>
13 warning	code-invalid	<p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
11 warning	code-invalid	<p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
6 warning	code-invalid	<p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
1 warning	code-invalid	<p>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</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>
22 warning	code-invalid	<p>[Bundle.entry[6].resource/*DiagnosticReport/null*/.subject.identifier.type]</p>

severity	code	details
		Unknown Code 'R112.1' in the CodeSystem ' <a href="https://fhir.nhs.uk/">https://fhir.nhs.uk/</a> CodeSystem/England- GenomicTestDirectory' version '0.2.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment