

severity	code	details	expression
29	information informational	<p>This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/">https://fhir.nwgenomics.nhs.uk/</a></p> <p>StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/">https://fhir.nwgenomics.nhs.uk/</a></p>	[Bundle.entry[6].resource.code.coding[0]]
28	information informational	<p>StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>Entry 'urn:uuid:dff0b8e5-a42d-4ad0-952e-b7c20c143877' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p>	[Bundle.entry[6].resource.resultsInterpreter[0]]
27	information informational	<p>None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (<a href="http://hl7.org/fhir/ValueSet/report-codes 4.0.1">http://hl7.org/fhir/ValueSet/report-codes 4.0.1</a>), 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>Reference to draft CodeSystem <a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0">https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0</a></p>	[Bundle.entry[8]]
20	information code-invalid		[Bundle.entry[6].resource/*DiagnosticReport/null*.code]
16	information business-rule		[Bundle.entry[5].resource/*ServiceRequest/null*.reasonCode[0]]
21	warning invalid	<p>value should not start or finish with whitespace ' Factor II deficiency (Single gene sequencing &gt;=10 amplicons)'</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>	[Bundle.entry[6].resource/*DiagnosticReport/null*.code.coding[1].display]
22	warning code-invalid		[Bundle.entry[6].resource/*DiagnosticReport/null*.subject.identifier.type]
19	warning code-invalid	<p>Unknown Code 'R112.1' in the CodeSystem '<a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory 0.1.0">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory 0.1.0</a>' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>	[Bundle.entry[6].resource/*DiagnosticReport/null*.code.coding[1].code]

<b>severity</b>	<b>code</b>	<b>details</b>	<b>expression</b>
15 warning	code-invalid	<p>None of the codings provided are in the value set 'IdentifierType' (<a href="http://hl7.org/fhir/ValueSet/identifier-type">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p> <p>Unknown Code 'R112' in the CodeSystem '<a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/">https://fhir.nwgenomics.nhs.uk/CodeSystem/</a>'</p>	[Bundle.entry[5].resource/*ServiceRequest/null*./subject.identifier.type]
17 warning	code-invalid	<p>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</p> <p>value should not start or finish with whitespace ' Factor II deficiency (Single gene sequencing &gt;=10 amplicons)'</p> <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 '0.1.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>	[Bundle.entry[5].resource/*ServiceRequest/null*./reasonCode[0].coding[0].code]
14 warning	invalid	<p>None of the codings provided are in the value set 'IdentifierType' (<a href="http://hl7.org/fhir/ValueSet/identifier-type">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>	[Bundle.entry[5].resource/*ServiceRequest/null*./code.coding[0].display]
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">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>	[Bundle.entry[5].resource/*ServiceRequest/null*./code.coding[0].code]
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">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>	[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' (<a href="http://hl7.org/fhir/ValueSet/identifier-type">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>	[Bundle.entry[3].resource/*DocumentReference/null*./subject.identifier.type]
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">http://hl7.org/fhir/ValueSet/identifier-type</a> 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p>	[Bundle.entry[1].resource/*Patient/null*./identifier[0].type]