

	severity	code	details
72	error	invalid	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[14].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
115	error	processing	DiagnosticReport.category:Genetics in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
118	error	structure	Slice 'DiagnosticReport.code.coding:GenomicTestDirectory': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1). Note that other slices are allowed in addition to this required slice
119	error	structure	Unable to find a profile match for urn:uuid:ed9bee63-1665-466b-8252-c46a049bb838 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
100	error	structure	Unable to find a profile match for urn:uuid:3662b42bd29f-4dd1-8ef5-4197193ebbd0 among choices: http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core, http://hl7.org/fhir/StructureDefinition/ImagingStudy
121	error	structure	Unable to find a profile match for urn:uuid:b0d89d0f-0763-4c71-8168-11e82bda5447 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
123	error	structure	Unable to find a profile match for urn:uuid:7e90bc58-19a2-4f6d-a2e8-9a80872b08cf among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
125	error	structure	Unable to find a profile match for urn:uuid:9c0bcf16-b0eb-4013-aeb5-e0b7c6e2e3bc among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
127	error	structure	Unable to find a profile match for urn:uuid:1f4236bef2ff-47e2-9c41-de1d6e95c836 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
129	error	structure	Unable to find a profile match for urn:uuid:bb8d8d55-1335-4e34-a95e-f43ace95a9b9 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
75	error	invalid	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
141	error	structure	Unable to find a profile match for urn:uuid:6196a7b5-e5ed-47b1-9e66-cbea3681616e among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
139	error	structure	Unable to find a profile match for urn:uuid:3662b42bd29f-4dd1-8ef5-4197193ebbd0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
102	error	invalid	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/

severity	code	details
137 error	structure	StructureDefinition/DiagnosticImplication 0.2.1 and slicing is CLOSED: Bundle.entry[15].resource/*Observation/*Observation/null*.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) Unable to find a profile match for urn:uuid:8c4065b0-9dfb-4bfa-a9e6-e866a3e9c8b1 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
133 error	structure	Unable to find a profile match for urn:uuid:a4b7540a-99bc-4f97-b122-4ba5668780fb among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
52 error	invariant	Constraint failed: obs-7: 'If Observation.code is the same as an Observation.component.code then the value element associated with the code SHALL NOT be present'
131 error	structure	Unable to find a profile match for urn:uuid:bed12a54-1866-4ba8-814e-c66057c6641c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
135 error	structure	Unable to find a profile match for urn:uuid:b509b23a-252f-41ec-abec-95bbc4b59fdc among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
91 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
89 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
93 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
90 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
92 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
86 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
87 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
85 information informational		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
84 information informational		

	severity	code	details
		This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept)]
83	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)]
82	information code-invalid	None of the codings provided are in the value set 'Laboratory Code' (http://hl7.eu/fhir/base/ValueSet/lab-obsCode-eu-lab 2.0.0-ballot), and a coding is recommended to come from this value set (codes = null#48006-1)	[Bundle.entry[14].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept)]
81	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept)]
80	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]
79	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept)]
78	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
77	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
76	information code-invalid	Binding for path <code>Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 has no source, so can't be checked	[Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
88	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept)]
94	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[18].value.ofType(CodeableConcept)]
114	information informational	Entry 'urn:uuid:5966713b-286f-41c0-93eb-f85f8201fc93' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	[Bundle.entry[19]]
97	information structure	Details for urn:uuid:3662b42bf29f-4dd1-8ef5-4197193ebbd0 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[15].resource/*Observation/null*/.derivedFrom[11]]
142	information structure	Details for urn:uuid:6196a7b5-e5ed-47b1-9e66-cbea3681616e matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1	[Bundle.entry[17].resource.result[11]]

severity	code	details
140 information structure	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 Details for urn:uuid:3662b42bd29f-4dd1-8ef5-4197193ebbd0 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[10]]
138 information structure	Details for urn:uuid:8c4065b0-9dfb-4bfa-a9e6-e866a3e9c8b1 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[9]]
136 information structure	Details for urn:uuid:b509b23a-252f-41ec-abec-95bbc4b59fdc matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[8]]
134 information structure	Details for urn:uuid:a4b7540a-99bc-4f97-b122-4ba5668780fb matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[7]]
132 information structure	Details for urn:uuid:bed12a54-1866-4ba8-814ec66057c6641c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[6]]
130 information structure	Details for urn:uuid:bb8d8d55-1335-4e34-a95ef43ace95a9b9 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[5]]
128 information structure	Details for urn:uuid:1f4236be-f2ff-47e2-9c41-de1d6e95c836 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[4]]
126 information structure	Details for urn:uuid:9c0bcf16-b0eb-4013-aeb5-e0b7c6e2e3bc matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[3]]
95 information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:7e90bc58-19a2-4f6da2e8-9a80872b08cf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[15].resource/*Observation/null*/.category[1]] [Bundle.entry[17].resource.result[2]]
124 information structure	Details for urn:uuid:ed9bee63-1665-466b-8252-c46a049bb838 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[17].resource.result[0]]
117 information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required	[Bundle.entry[17].resource.code.coding[0]]
116 information not-supported	(DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[17].resource]
74 information informational	Details for urn:uuid:6196a7b5-e5ed-47b1-9e66-cbea3681616e matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
109 information structure	Details for urn:uuid:6196a7b5-e5ed-47b1-9e66-cbea3681616e matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[17].resource/*DiagnosticReport/null*/.result[1]]
108 information structure	Details for urn:uuid:3662b42bd29f-4dd1-8ef5-4197193ebbd0 matching against	[Bundle.entry[17].resource/*DiagnosticReport/null*/.result[2]]

	severity	code	details
		profile http://hl7.org/fhir/StructureDefinition/ Observation 4.0.1	
104	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/DiagnosticImplication 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[15].resource/*Observation/ null*.component[0].value.ofType(CodeableConcept)]
103	information code-invalid	Binding for path Bundle.entry[15].resource/ *Observation/ null*.component[0].value.ofType(CodeableConcept) in profile StructureDefinition[https:// fhir.nwgenomics.nhs.uk/StructureDefinition/ DiagnosticImplication 0.2.1] has no source, so can't be checked	[Bundle.entry[15].resource/*Observation/ null*.component[0].value.ofType(CodeableConcept)]
101	information structure	Details for urn:uuid:3662b42b- d29f-4dd1-8ef5-4197193ebbd0 matching against profile http://hl7.eu/fhir/base/StructureDefinition/ medicalTestResult-eu-core 2.0.0-ballot	[Bundle.entry[15].resource/*Observation/null*.derivedFrom]
122	information structure	Details for urn:uuid:b0d89d0f-0763-4c71-8168-11e82bda5447 matching against profile https:// fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation 0.2.1	[Bundle.entry[17].resource.result[1]]
71	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/ null*.component[0].value.ofType(CodeableConcept)]
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:	[Bundle.entry[0].resource/*MessageHeader/null*.source.e
57	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
70	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
53	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.category[2].value.ofType(CodeableConcept)]
54	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
55	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
56	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
58	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Variant 0.2.1 (this may not be a	[Bundle.entry[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]

	severity	code	details
16	warning	not-found	<p>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>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p>
21	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>
22	warning	not-found	<p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p>
15	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>
12	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>
10	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated</p>
9	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>
7	warning	not-found	<p>A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated</p>
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>
24	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>
4	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>
2	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>
13	warning	not-found	<p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p>
40	warning	not-found	

severity	code	details
		A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
73 warning	not-found	A definition for CodeSystem 'http://www.genenames.org' version 'null' could not be found, so the code cannot be validated. Valid versions: 3.0.0 or 3.1.0
50 warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
49 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/projects/SNP' could not be found, so the code cannot be validated
48 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
47 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
96 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)
46 warning	invalid	Coding has no system. A code with no system has no defined meaning, and it cannot be validated. A system should be provided
98 warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
45 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
29 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)
44 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
42 warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
41 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
105 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)
107 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)
39 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)
36 warning	code-invalid	[Bundle.entry[13].resource/*Observation/null*.subject.identifier.type]

severity	code	details
		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)
34 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)
43 warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
		[Bundle.entry[12].resource/*Observation/null*/.subject.id]
		[Bundle.entry[14].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding[0].code]