

	severity	code	details
94	error	structure	Unable to find a profile match for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d among choices: http://hl7.eu/fhir/base/ StructureDefinition/medicalTestResult-eu-core, http:// hl7.org/fhir/StructureDefinition/ImagingStudy This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ StructureDefinition/DiagnosticImplication 0.2.1 and slicing is CLOSED: Bundle.entry[14].resource/ *Observation/ null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
96	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))
69	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[13].resource/*Observation/ null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
112	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 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
114	error	structure	Unable to find a profile match for urn:uuid:7b9b85cb-39c7-42c2-a0c5-1b35b5b62e01 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation
116	error	structure	Unable to find a profile match for urn:uuid:4345aa81-414f-4fed-98be-474f4cc9d503 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation
118	error	structure	Unable to find a profile match for urn:uuid:1d219ff3- feb2-4af2-b437-cde2ed8434c4 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation
66	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[13].resource/*Observation/ null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
130	error	structure	Unable to find a profile match for urn:uuid:ef6fd243-5856-46cd-b084-60b8d4b3f2e8 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation
120	error	structure	Unable to find a profile match for urn:uuid:586483cd-51fb-4e86-a996-53f2cf5e8610 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation
122	error	structure	Unable to find a profile match for urn:uuid:f6fa269a-9cc7-4aff-bb74-076fab5e7e38 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation
134	error	structure	Unable to find a profile match for urn:uuid:79f6868b- f94b-4887-8910-439d64482640 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation
124	error	structure	Unable to find a profile match for urn:uuid:e393d45d-0d96-4d62-a7a8-71f96e652158

	severity	code	details
		among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	
126 error	structure	Unable to find a profile match for urn:uuid:98fdd1bd-d654-4bb1-bb6b-d4a2adec2321 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[16].resource.result[6]]
128 error	structure	Unable to find a profile match for urn:uuid:c401e33a-80e0-467c-bf2b-10878904dde2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[16].resource.result[7]]
132 error	structure	Unable to find a profile match for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[16].resource.result[9]]
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)	[Bundle.entry[13].resource/*Observation/null*.component[17].value.ofType(CodeableConcept)]
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)	[Bundle.entry[13].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)]
84 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[13].resource/*Observation/null*.component[15].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[13].resource/*Observation/null*.component[14].value.ofType(CodeableConcept)]
82 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[13].resource/*Observation/null*.component[13].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[13].resource/*Observation/null*.component[12].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[13].resource/*Observation/null*.component[11].value.ofType(string)]
119 information structure		Details for urn:uuid:1d219ff3-feb2-4af2-b437-cde2ed8434c4 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[16].resource.result[2]]
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[13].resource/*Observation/null*.component[9].value.ofType(string)]
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)	[Bundle.entry[13].resource/*Observation/null*.component[18].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	[Bundle.entry[13].resource/*Observation/null*.component[8].value.ofType(CodeableConcept)]

	severity	code	details
		problem, but you should check that it's not intended to match a slice)	
		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)	
76	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[13].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)]
75	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[13].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept)]
74	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[13].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept)]
73	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[13].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]
72	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[13].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept)]
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[13].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
70	information code-invalid	Binding for path Bundle.entry[13].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept) in profile StructureDefinition[https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1] has no source, so can't be checked	[Bundle.entry[13].resource/*Observation/null*/.component[2].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[13].resource/*Observation/null*/.component[10].value.ofType(Range)]
88	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)	[Bundle.entry[14].resource/*Observation/null*/.category[1]]
95	information structure	Details for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d matching against profile http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core 2.0.0-ballot	[Bundle.entry[14].resource/*Observation/null*/.derivedFrom[0]]
91	information business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1	[Bundle.entry[14].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]
121	information structure	Details for urn:uuid:586483cd-51fb-4e86-a996-53f2cf5e8610 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[16].resource.result[3]]
117	information structure	Details for urn:uuid:4345aa81-414f-4fed-98be-474f4cc9d503 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[16].resource.result[1]]
115	information structure	Details for urn:uuid:7b9b85cb-39c7-42c2-a0c5-1b35b5b62e01 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[16].resource.result[0]]

severity	code	details
123 information structure	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	Details for urn:uuid:f6fa269a-9cc7-4aff-bb74-076fab5e7e38 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[4]]
125 information structure	urn:uuid:e393d45d-0d96-4d62-a7a8-71f96e652158	Details for urn:uuid:e393d45d-0d96-4d62-a7a8-71f96e652158 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[5]]
127 information structure	urn:uuid:98fdd1bd-d654-4bb1-bb6b-d4a2adee2321	Details for urn:uuid:98fdd1bd-d654-4bb1-bb6b-d4a2adee2321 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[6]]
129 information structure	urn:uuid:c401e33a-80e0-467c-bf2b-10878904dde2	Details for urn:uuid:c401e33a-80e0-467c-bf2b-10878904dde2 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[7]]
113 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 0.2.1)	[Bundle.entry[16].resource]
111 information informational	urn:uuid:51a7427c-12ba-468a-9051-5450c8341c52' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	Entry [Bundle.entry[18]]
131 information structure	urn:uuid:ef6fd243-5856-46cd-b084-60b8d4b3f2e8	Details for urn:uuid:ef6fd243-5856-46cd-b084-60b8d4b3f2e8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[8]]
133 information structure	urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d	Details for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[9]]
106 information structure	urn:uuid:f94b-4887-8910-439d-64482640	Details for urn:uuid:f94b-4887-8910-439d-64482640 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[16].resource/*DiagnosticReport/null*.result[10]]
105 information structure	urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d	Details for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[16].resource/*DiagnosticReport/null*.result[11]]
103 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.nhs.uk/CodeSystem/England-GenomicTestDirectory R210.2)	[Bundle.entry[16].resource/*DiagnosticReport/null*.code]
98 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[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
97 information code-invalid	Binding for path Bundle.entry[14].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[14].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
135 information structure	urn:uuid:79f6868b-f94b-4887-8910-439d-64482640	Details for urn:uuid:79f6868b-f94b-4887-8910-439d-64482640 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[16].resource.result[10]]
90 information structure		[Bundle.entry[14].resource/*Observation/null*.derivedFrom]

	severity	code	details
0	information informational	Details for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	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:
68	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[0].resource/*MessageHeader/null*/.source.en
51	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[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
34	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)	[Bundle.entry[13].resource/*Observation/null*/.category[1]]
47	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[13].resource/*Observation/null*/.category[2]]
48	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[13].resource/*Observation/null*/.component[1]
49	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[13].resource/*Observation/null*/.component[2]
50	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[13].resource/*Observation/null*/.component[3]
65	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[13].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]
52	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[13].resource/*Observation/null*/.component[4]
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[13].resource/*Observation/null*/.component[5]
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[13].resource/*Observation/null*/.component[6]
7	information business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1	[Bundle.entry[3].resource/*Observation/null*/.value.ofType(CodeableConcept)]
55	information informational		[Bundle.entry[13].resource/*Observation/null*/.component[7]]

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)
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)
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 problem, but you should check that it's not intended to match a slice)
59	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)
60	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)
61	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)
62	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)
63	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)
64	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)
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)
16	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)
18	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)
67	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
20	warning code-invalid	[Bundle.entry[8].resource/*Observation/null*/.subject.identifi

	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)
25	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)
14	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
4	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)
11	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
10	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)
8	warning	code-invalid	Unknown Code 'R210' in the CodeSystem ' https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication ' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
6	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)
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)
13	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)
30	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)
102	warning	code-invalid	Unknown Code 'R210.2' in the CodeSystem ' https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory ' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
45	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
89	warning	code-invalid	

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)
44	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
43	warning not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
92	warning code-invalid	Unknown Code 'R210' in the CodeSystem ' https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication ' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
42	warning not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
41	warning not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
40	warning not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
38	warning not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
99	warning code-invalid	Unknown Code 'R210.2' in the CodeSystem ' https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory ' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
100	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)
37	warning not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
104	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 not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
35	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)
32	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 not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated