

20, Feb 2026

	severity	code	details
94	error	structure	<p>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</p> <p>[Bundle.entry[14].resource/*Observation/null*/.derivedFrom</p>
96	error	invalid	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication0.2.1 and slicing is CLOSED: Bundle.entry[14].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>[Bundle.entry[14].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]</p>
69	error	invalid	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant0.2.1 and slicing is CLOSED: Bundle.entry[13].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]</p>
112	error	processing	<p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice DiagnosticReport.category:Genetics in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>[Bundle.entry[16].resource.category[0]]</p>
114	error	structure	<p>Unable to find a profile match for urn:uuid:7b9b85cb-39c7-42c2-a0c5-1b35b5b62e01 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[0]]</p>
116	error	structure	<p>Unable to find a profile match for urn:uuid:4345aa81-414f-4fed-98be-474f4cc9d503 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[1]]</p>
118	error	structure	<p>Unable to find a profile match for urn:uuid:1d219ff3-feb2-4af2-b437-cde2ed8434c4 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[2]]</p>
66	error	invalid	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant0.2.1 and slicing is CLOSED: Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
130	error	structure	<p>Unable to find a profile match for urn:uuid:ef6fd243-5856-46cd-b084-60b8d4b3f2e8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[8]]</p>
120	error	structure	<p>Unable to find a profile match for urn:uuid:586483cd-51fb-4e86-a996-53f2cf5e8610 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[3]]</p>
122	error	structure	<p>Unable to find a profile match for urn:uuid:f6fa269a-9cc7-4aff-bb74-076fab5e7e38 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[4]]</p>
134	error	structure	<p>Unable to find a profile match for urn:uuid:79f6868b-f94b-4887-8910-439d64482640 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>[Bundle.entry[16].resource.result[10]]</p>
124	error	structure	<p>Unable to find a profile match for urn:uuid:c393d45d-0d96-4d62-a7a8-71f96e652158</p> <p>[Bundle.entry[16].resource.result[5]]</p>

severity	code	details
126 error	structure	among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation Unable to find a profile match for urn:uuid:98fdd1bd-d654-4bb1-bb6b-d4a2adee2321 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)
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]
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
	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	
123	information structure	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	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	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	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	Entry '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) [Bundle.entry[18]]
131	information structure	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	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	Details for urn:uuid:79f6868b-f94b-4887-8910-439d64482640 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[16].resource/*DiagnosticReport/null*/.result[10]]
105	information structure	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	Details for urn:uuid:79f6868b-f94b-4887-8910-439d64482640 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	<p>Details for urn:uuid:d4a74eb4-4b44-4465-af88-9a97ed8e002d matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</p> <p>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:</p> <p>[Bundle.entry[0].resource/*MessageHeader/null*/.source.en</p>
68	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
51	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
34	information code-invalid	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.category[1]]</p>
47	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.category[2]]</p>
48	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
49	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
50	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
65	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]</p>
52	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
53	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
54	information informational	<p>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)</p> <p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
7	information business-rule	<p>Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1</p> <p>[Bundle.entry[3].resource/*Observation/null*/.value.ofType(CodeableConcept)]</p>
55	information informational	<p>[Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>

	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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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) [Bundle.entry[13].resource/*Observation/null*/.component[
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[13].resource/*Observation/null*/.component[
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) [Bundle.entry[6].resource/*Observation/null*/.subject.identi
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) [Bundle.entry[7].resource/*Observation/null*/.subject.identi
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 [Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding
20	warning	code-invalid	[Bundle.entry[8].resource/*Observation/null*/.subject.identi

	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)	[Bundle.entry[10].resource/*DocumentReference/null*/.subject.identifier.type]
14	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated	[Bundle.entry[5].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
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)	[Bundle.entry[2].resource/*Observation/null*/.subject.identi
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	[Bundle.entry[4].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
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)	[Bundle.entry[4].resource/*Observation/null*/.subject.identi
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	[Bundle.entry[3].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].code]
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)	[Bundle.entry[3].resource/*Observation/null*/.subject.identi
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)	[Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]
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)	[Bundle.entry[5].resource/*Observation/null*/.subject.identi
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)	[Bundle.entry[11].resource/*Observation/null*/.subject.identi
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	[Bundle.entry[16].resource/*DiagnosticReport/null*/.code.coding[0].code]
45	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated	[Bundle.entry[13].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept).codin
89	warning	code-invalid		[Bundle.entry[14].resource/*Observation/null*/.subject.identi

	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 [Bundle.entry[13].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding]
43	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated [Bundle.entry[13].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[14].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[13].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding]
41	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated [Bundle.entry[13].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
40	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated [Bundle.entry[13].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
38	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[13].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[15].resource/*ServiceRequest/null*/.code.coding]
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) [Bundle.entry[15].resource/*ServiceRequest/null*/.subject.identifier.type]
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 [Bundle.entry[13].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
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) [Bundle.entry[16].resource/*DiagnosticReport/null*/.subject.identifier.type]
36	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[13].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
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) [Bundle.entry[13].resource/*Observation/null*/.subject.identifier.type]
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) [Bundle.entry[12].resource/*Observation/null*/.subject.identifier.type]
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 [Bundle.entry[13].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]