

20, Feb 2026

	severity	code	details
0	error	invalid	Object must have some content [Bundle.entry[16].resource.value[x].coding[0]]
153	error	structure	Unable to find a profile match for urn:uuid:61b2ceec-3fa7-4d26-bda9-6805228a50f3 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[7]]
149	error	structure	Unable to find a profile match for urn:uuid:a8d7657e- fc43-4b2e-9f33-ddca3ae1b7ff among choices: https:// fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation [Bundle.entry[25].resource.result[5]]
147	error	structure	Unable to find a profile match for urn:uuid:86987d94-6a02-4732-99dd-2419c22929a0 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[4]]
145	error	structure	Unable to find a profile match for urn:uuid:003a69ff-e8e6-4284-aaf7-3ae2a65b7528 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation [Bundle.entry[25].resource.result[3]]
143	error	structure	Unable to find a profile match for urn:uuid:6e4748f0-b39c-4fd7-be44-c9ae770db4fe among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation [Bundle.entry[25].resource.result[2]]
141	error	structure	Unable to find a profile match for urn:uuid:75dd95b5-1a44-4a0d-8387-6974ea634f32 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[1]]
139	error	structure	Unable to find a profile match for urn:uuid:43f44242-2429-438b-8a3f-89261d7a017e among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[0]]
155	error	structure	Unable to find a profile match for urn:uuid:830670a5-4f57-4606-9b13-e632de305b55 among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[8]]
137	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 [Bundle.entry[25].resource.category[0]]
119	error	structure	Unable to find a profile match for urn:uuid:7b4bf27c1fe4-423e-b148-9ebce7e6ce76 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[23].resource/ *Observation/ null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[23].resource/*Observation/null*/.derivedFrom]
121	error	invalid	Unable to find a profile match for urn:uuid:c45e83e8-56ce-49c4-897e-a65dbe1d3f2b among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[23].resource/*Observation/ null*/.component[0].value.ofType(CodeableConcept)]
157	error	structure	Unable to find a profile match for urn:uuid:c45e83e8-56ce-49c4-897e-a65dbe1d3f2b among choices: https://fhir.nwgenomics.nhs.uk/ StructureDefinition/Observation [Bundle.entry[25].resource.result[9]]
151	error	structure	Unable to find a profile match for urn:uuid:54acbe54-a402-499a-91ed-20938a813692 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/ Observation [Bundle.entry[25].resource.result[6]]

	severity	code	details
159	error	structure	Unable to find a profile match for urn:uuid:f70dd662-f225-472a-bfed-0b28a5d71d8a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[10]]
177	error	structure	Unable to find a profile match for urn:uuid:3cd7cee2-821a-478e-a1df-520d96bcae5b among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[19]]
175	error	structure	Unable to find a profile match for urn:uuid:7b4bf27c-1fe4-423e-b148-9ebce7e6ce76 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[18]]
93	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[22].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[22].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
173	error	structure	Unable to find a profile match for urn:uuid:a909e4ce-17aa-4a36-89e6-758bc920dfb3 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[17]]
171	error	structure	Unable to find a profile match for urn:uuid:4aae243f-19bb-416c-9d58-4e7f5c6d3f79 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[16]]
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[22].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[22].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
169	error	structure	Unable to find a profile match for urn:uuid:867d30f2-443d-4b26-9aad-43dc9eb435a8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[15]]
167	error	structure	Unable to find a profile match for urn:uuid:e701f1e7-c042-4d25-afef-68ed694afa6a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[14]]
165	error	structure	Unable to find a profile match for urn:uuid:97ed03ab-bc1a-40bb-b689-046bf607d805 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[13]]
163	error	structure	Unable to find a profile match for urn:uuid:457abb89-dbe1-43aa-b2eb-36ed8650ed70 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[12]]
161	error	structure	Unable to find a profile match for urn:uuid:7ee8f4ee-7744-4e7e-8a08-7d0456da8e83 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[25].resource.result[11]]
108	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[22].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept)]
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) [Bundle.entry[22].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept)]
103	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[22].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)]

	severity	code	details
		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)	[Bundle.entry[22].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
101	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[22].resource/*Observation/null*.component[5].value.ofType(CodeableConcept)]
104	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[22].resource/*Observation/null*.component[8].value.ofType(string)]
120	information structure	Details for urn:uuid:7b4bf27c-1fe4-423e-b148-9ebce7e6ce76 matching against profile http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core 2.0.0-ballot	[Bundle.entry[23].resource/*Observation/null*.derivedFrom]
116	information business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1	[Bundle.entry[23].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
115	information structure	Details for urn:uuid:7b4bf27c-1fe4-423e-b148-9ebce7e6ce76 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[23].resource/*Observation/null*.derivedFrom]
95	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[22].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]
105	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[22].resource/*Observation/null*.component[9].value.ofType(Range)]
100	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[22].resource/*Observation/null*.component[4].value.ofType(CodeableConcept)]
113	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[23].resource/*Observation/null*.category[1]]
106	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[22].resource/*Observation/null*.component[10].value.ofType(string)]
112	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[22].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)]
97	information code-invalid	Binding for path <code>Bundle.entry[22].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)</code> in profile <code>StructureDefinition[https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1]</code> has no source, so can't be checked	[Bundle.entry[22].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
102	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[22].resource/*Observation/null*.component[6].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)
110	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[14].value.ofType(CodeableConcept)]
		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)
98	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
		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)
109	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept)]
		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)
107	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept)]
		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)
99	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept)]
		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)
111	information informational	[Bundle.entry[22].resource/*Observation/null*/.component[15].value.ofType(CodeableConcept)]
136	information informational	[Bundle.entry[27]]
		Entry 'urn:uuid:01a42e01-d2c0-41af-a595-9f7a139556a2' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)
123	information informational	[Bundle.entry[23].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]
		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)
178	information structure	[Bundle.entry[25].resource.result[19]]
		Details for urn:uuid:3cd7cee2-821a-478e-a1df-520d96bcac5b matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
176	information structure	[Bundle.entry[25].resource.result[18]]
		Details for urn:uuid:7b4bf27c-1fe4-423e-b148-9ebce7e6ce76 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
174	information structure	[Bundle.entry[25].resource.result[17]]
		Details for urn:uuid:a909e4ce-17aa-4a36-89e6-758bc920dfb3 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
172	information structure	[Bundle.entry[25].resource.result[16]]
		Details for urn:uuid:4aae243f-19bb-416c-9d58-4e7f5c6d3f79 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
170	information structure	[Bundle.entry[25].resource.result[15]]
		Details for urn:uuid:867d30f2-443d-4b26-9aad-43dc9eb435a8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
168	information structure	[Bundle.entry[25].resource.result[14]]
		Details for urn:uuid:e701f1e7-c042-4d25-afef-68ed694afa6a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1

severity	code	details
166 information structure	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 Details for urn:uuid:97ed03ab-bc1a-40bb-b689-046bf607d805 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[13]]
164 information structure	Details for urn:uuid:457abb89-dbe1-43aa-b2eb-36ed8650ed70 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[12]]
162 information structure	Details for urn:uuid:7ee8f4ee-7744-4e7e-8a08-7d0456da8e83 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[11]]
160 information structure	Details for urn:uuid:f70dd662-f225-472a-bfed-0b28a5d71d8a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[10]]
158 information structure	Details for urn:uuid:e45e83e8-56ce-49c4-897e-a65dbe1d3f2b matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[9]]
156 information structure	Details for urn:uuid:830670a5-4f57-4606-9b13-e632de305b55 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[8]]
154 information structure	Details for urn:uuid:61b2ceec-3fa7-4d26-bda9-6805228a50f3 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[7]]
152 information structure	Details for urn:uuid:54acbe54-a402-499a-91ed-20938a813692 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[6]]
150 information structure	Details for urn:uuid:a8d7657e-fc43-4b2e-9f33-ddca3ae1b7ff matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[5]]
148 information structure	Details for urn:uuid:86987d94-6a02-4732-99dd-2419c22929a0 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[4]]
146 information structure	Details for urn:uuid:003a69ff-e8e6-4284-aaf7-3ae2a65b7528 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[3]]
144 information structure	Details for urn:uuid:6e4748f0-b39c-4fd7-be44-c9ae770db4fe matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[2]]
142 information structure	Details for urn:uuid:75dd95b5-1a44-4a0d-8387-6974ea634f32 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[1]]
140 information structure	Details for urn:uuid:43f44242-2429-438b-8a3f-89261d7a017e matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[25].resource.result[0]]
138 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[25].resource]

	severity	code	details
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)	[Bundle.entry[22].resource/*Observation/null*/.component[
131	information structure	Details for urn:uuid:3cd7cee2-821a-478e-a1df-520d96bcae5b matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[25].resource/*DiagnosticReport/null*/.result[
130	information structure	Details for urn:uuid:7b4bf27c-1fe4-423e-b148-9ebce7e6ce76 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[25].resource/*DiagnosticReport/null*/.result[
128	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#R185.1)	[Bundle.entry[25].resource/*DiagnosticReport/null*/.code[
122	information code-invalid	Binding for path Bundle.entry[23].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept) in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 has no source, so can't be checked	[Bundle.entry[23].resource/*Observation/null*/.component[0].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[22].resource/*Observation/null*/.component[
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)	[Bundle.entry[22].resource/*Observation/null*/.component[
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[22].resource/*Observation/null*/.component[
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[22].resource/*Observation/null*/.component[
8	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)]
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[22].resource/*Observation/null*/.category[2]]
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[22].resource/*Observation/null*/.component[
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[22].resource/*Observation/null*/.component[
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[22].resource/*Observation/null*/.component[

	severity	code	details
80	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>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>
63	information	code-invalid	<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>
82	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>
83	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>
84	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>
85	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>
1	information	informational	<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>
81	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>
23	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	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p>
26	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>
27	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p>
29	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>
32	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</p>

	severity	code	details
33	warning	not-found	<p>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> <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> <p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p> <p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p> <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> <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> <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> <p>Unknown Code 'R185' 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</p> <p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p> <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> <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> <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> <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>
35	warning	code-invalid	[Bundle.entry[12].resource/*Observation/null*.value.ofType(CodeableConcept).coding[0].system]
36	warning	not-found	[Bundle.entry[12].resource/*Observation/null*.value.ofType(CodeableConcept).coding[0].system]
30	warning	not-found	[Bundle.entry[10].resource/*Observation/null*.value.ofType(CodeableConcept).coding[0].system]
20	warning	code-invalid	[Bundle.entry[7].resource/*Observation/null*.subject.identifier]
3	warning	code-invalid	[Bundle.entry[1].resource/*Patient/null*.identifier[0].type]
5	warning	code-invalid	[Bundle.entry[2].resource/*Observation/null*.subject.identifier]
7	warning	code-invalid	[Bundle.entry[3].resource/*Observation/null*.subject.identifier]
9	warning	code-invalid	[Bundle.entry[3].resource/*Observation/null*.value.ofType(CodeableConcept).coding[0].code]
21	warning	not-found	[Bundle.entry[7].resource/*Observation/null*.value.ofType(CodeableConcept).coding[0].system]
11	warning	code-invalid	[Bundle.entry[4].resource/*Observation/null*.subject.identifier]
14	warning	code-invalid	[Bundle.entry[5].resource/*Observation/null*.subject.identifier]
38	warning	code-invalid	[Bundle.entry[13].resource/*Observation/null*.subject.identifier]

	severity	code	details
15	warning	not-found	<p>validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)</p> <p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> <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>
17	warning	code-invalid	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p> <p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated</p>
18	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p>
12	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>
39	warning	not-found	<p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p>
44	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>
41	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>
114	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>
64	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>
65	warning	not-found	<p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p>
66	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p>
67	warning	not-found	<p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p>
68	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>
61	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>
69	warning	not-found	<p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p>
71	warning	not-found	

severity	code	details
	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated	[Bundle.entry[22].resource/*Observation/null*.component[6].value.ofType(CodeableConcept).coding]
72 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
73 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
74 warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
94 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
70 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
117 warning	code-invalid	Unknown Code 'R185' 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://varnomen.hgvs.org' 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
47 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)
129 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)
49 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)
127 warning	code-invalid	Unknown Code 'R185.1' 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
125 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)
124 warning	code-invalid	Unknown Code 'R185.1' 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
54 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)

	severity	code	details
59	warning	code-invalid	validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH) 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[20].resource/*Observation/null*.subject.ident