

severity	code	details
137 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice [Bundle.entry[8].resource/*Observation/null*/]
105 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice [Bundle.entry[7].resource/*Observation/null*.category[1]]]
104 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[7].resource/*Observation/null*.category[0]]]
232 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) 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[11].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[11].resource/*Observation/null*/]
233 error	invalid	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[11].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[11].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]]
90 error	invalid	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[6].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[6].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]]
86 error	invalid	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[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]]
85 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice [Bundle.entry[6].resource/*Observation/null*/]
83 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice [Bundle.entry[6].resource/*Observation/null*.category[1]]]
106 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice [Bundle.entry[7].resource/*Observation/null*.category[2]]]
82 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[6].resource/*Observation/null*.category[0]]]
57 error	invalid	

severity	code	details
55 error	structure	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[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
53 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
52 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
51 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions 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[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
60 error	invalid	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
108 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[7].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
110 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[7].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) 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[7].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
113 error	invalid	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
200 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
199 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
198 error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice

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204 error	invalid	<p>Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>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[10].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>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[10].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>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[9].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p> <p>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[9].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p>	[Bundle.entry[10].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
207 error	invalid	<p>Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[10].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
169 error	invalid	<p>Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[9].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
166 error	invalid	<p>Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[9].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
164 error	structure	<p>Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[9].resource/*Observation/null*/]
162 error	processing	<p>Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[9].resource/*Observation/null*/.category[2]]
161 error	processing	<p>Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>	[Bundle.entry[9].resource/*Observation/null*/.category[1]]
160 error	processing	<p>Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>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[8].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p>	[Bundle.entry[9].resource/*Observation/null*/.category[0]]
146 error	invalid	<p>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[8].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p>	[Bundle.entry[8].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]
143 error	invalid	<p>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[8].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))</p>	[Bundle.entry[8].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
139 error	invalid		

	severity	code	details
			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[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
138 error	structure		[Bundle.entry[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
135 error	processing		Observation.component:clinical-significance: max allowed = 1, but found 2 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
134 error	processing		Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
229 error	processing		Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
230 error	processing		Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
253 error	processing		DiagnosticReport.category:Genetics in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
254 error	invalid		Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 , Element matches more than one slice - organization, operator
202 error	structure		Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)
262 error	structure		Unable to find a profile match for urn:uuid:6d1c937d-5bf2-44a1-be0c-e5579d8b5e40 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
268 error	structure		Unable to find a profile match for urn:uuid:37194eba-c9dc-4c39-a2e3-04281f69f5f4 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
15 error	processing		Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
16 error	processing		[Bundle.entry[2].resource/*Observation/null*.category[1]]

severity	code	details
260 error	structure	<p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions</p> <p>Unable to find a profile match for urn:uuid:8730acb2-151e-4dc6-aaff-c8a80c9a37bc among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
270 error	structure	<p>Unable to find a profile match for urn:uuid:35a7def2-76b8-4d7e-9921-ea291348d280 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
266 error	structure	<p>Unable to find a profile match for urn:uuid:53418515-ac4b-48f3-b474-561434a5dc46 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
18 error	structure	<p>Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1)</p> <p>Slice 'DiagnosticReport.code.coding:GenomicTestDirectory': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1). Note that other slices are allowed in addition to this required slice</p>
258 error	structure	<p>Unable to find a profile match for urn:uuid:4d84c6a3-418c-4da3-8a24-39d6417bd53c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
272 error	structure	<p>Unable to find a profile match for urn:uuid:ed0fbcf5-7874-4d67-b5ff-6a34c20188a2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
264 error	structure	<p>Unable to find a profile match for urn:uuid:970a761f-b7fd-4403-ac92-bf72ec4526ba among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p>
265 information	structure	<p>Details for urn:uuid:970a761f-b7fd-4403-ac92-bf72ec4526ba matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p>
222 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>
216 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>
217 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>
218 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>
215 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</p>

severity	code	details
219 information informational	problem, but you should check that it's not intended to match a slice)	[Bundle.entry[10].resource/*Observation/null*.component[12].value.ofType(CodeableConcept)]
220 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[10].resource/*Observation/null*.component[13].value.ofType(CodeableConcept)]
221 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[10].resource/*Observation/null*.component[14].value.ofType(CodeableConcept)]
223 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[10].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)]
256 information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[13].resource.code.coding[0]]
225 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[11].resource/*Observation/null*.category[1]]
147 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[8].resource/*Observation/null*.component[4].value.ofType(CodeableConcept)]
263 information structure	Details for urn:uuid:6d1c937d-5bf2-44a1-be0ce5579d8b5e40 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[2]]
145 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[8].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]
144 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[8].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
142 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[8].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]
141 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[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
140 information code-invalid	Binding for path Bundle.entry[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept) in profile StructureDefinition[https://fhir.nwgenomics.nhs.uk/StructureDefinition/]	[Bundle.entry[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]

severity	code	details
261 information structure	DiagnosticImplication[0.2.1] has no source, so can't be checked	
136 information not-supported	Details for urn:uuid:8730acb2-151e-4dc6-aaff-c8a80c9a37bc matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation[0.2.1] Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation [Bundle.entry[8].resource/*Observation/null*/] (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication[0.2.1]) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation [Bundle.entry[2].resource/*Observation/null*/] (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel[0.2.1])	[Bundle.entry[13].resource.result[1]]
17 information not-supported	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) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel[0.2.1] (this may not be a 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)	[Bundle.entry[8].resource/*Observation/null*.category[1]]
19 information informational	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) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation [Bundle.entry[2].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]	
128 information informational	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) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation [Bundle.entry[7].resource/*Observation/null*.component[15].value.ofType(CodeableConcept)]	
149 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) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation [Bundle.entry[9].resource/*Observation/null*.category[1]]	
163 information not-supported	Binding for path Bundle.entry[9].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 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[9].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
170 information code-invalid	Details for urn:uuid:ed0fbcf5-7874-4d67-b5ff-6a34c20188a2 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation[0.2.1] Details for urn:uuid:35a7def2-76b8-4d7e-9921-ea291348d280 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation[0.2.1]	[Bundle.entry[9].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
165 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[9].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
273 information structure	Details for urn:uuid:ed0fbcf5-7874-4d67-b5ff-6a34c20188a2 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation[0.2.1]	[Bundle.entry[13].resource.result[7]]
271 information structure	Details for urn:uuid:35a7def2-76b8-4d7e-9921-ea291348d280 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation[0.2.1]	[Bundle.entry[13].resource.result[6]]
203 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[10].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
4 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)	[Bundle.entry[2].resource/*Observation/null*.category[1]]

severity	code	details
206	information informational	<p>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>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>Binding for path Bundle.entry[10].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</p>
208	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>
209	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>
210	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>
211	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>
212	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>
187	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>
213	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>
185	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>
184	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>
183	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>
182	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>
181	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</p>

severity	code	details
214	information informational	<p>problem, but you should check that it's not intended to match a slice)</p> <p>This element does not match any known slice defined in the profile 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>
168	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>
269	information structure	<p>Details for urn:uuid:37194eba-c9dc-4c39-a2e3-04281f69f5f4 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p> <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>
127	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>
171	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>
172	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>
267	information structure	<p>Details for urn:uuid:53418515-ac4b-48f3-b474-561434a5dc46 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p> <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>
173	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>
175	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>
176	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>
177	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>
178	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>
179	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>
180	information informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p>

	severity	code	details
		StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	
174	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[9].resource/*Observation/null*.component[5].value.ofType(CodeableConcept)]
126	information informational	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[7].resource/*Observation/null*.component[13].value.ofType(CodeableConcept)]
123	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[7].resource/*Observation/null*.component[10].value.ofType(CodeableConcept)]
124	information informational	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[7].resource/*Observation/null*.component[11].value.ofType(CodeableConcept)]
73	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[5].resource/*Observation/null*.component[13].value.ofType(CodeableConcept)]
74	information informational	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[5].resource/*Observation/null*.component[14].value.ofType(CodeableConcept)]
75	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[5].resource/*Observation/null*.component[15].value.ofType(CodeableConcept)]
76	information informational	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[5].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)]
246	information structure	Details for urn:uuid:35a7def2-76b8-4d7e-9921-ea291348d280 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*.result]
78	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[6].resource/*Observation/null*.category[1]]
245	information structure	Details for urn:uuid:37194eba-c9dc-4c39-a2e3-04281f69f5f4 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*.result]
244	information structure	Details for urn:uuid:53418515-ac4b-48f3-b474-561434a5dc46 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*.result]
243	information structure	Details for urn:uuid:970a761f-b7fd-4403-ac92-bf72ec4526ba matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*.result]
84	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/) StructureDefinition/DiagnosticImplication 0.2.1)	[Bundle.entry[6].resource/*Observation/null*/]
30	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[2].resource/*Observation/null*.component[12].value.ofType(CodeableConcept)]

	severity	code	details
29	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	
87	information code-invalid	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) Binding for path Bundle.entry[6].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[2].resource/*Observation/null*.component[11].value.ofType(string)]
88	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[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
89	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[6].resource/*Observation/null*.component[0].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[5].resource/*Observation/null*.component[12].value.ofType(CodeableConcept)]
125	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[7].resource/*Observation/null*.component[12].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[5].resource/*Observation/null*.component[11].value.ofType(CodeableConcept)]
69	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[5].resource/*Observation/null*.component[9].value.ofType(Range)]
252	information informational	Entry 'urn:uuid:e25f6270-ceb5-4689-b949-046101febcd' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	[Bundle.entry[15]]
40	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:4d84c6a3-418c-4da3-8a24-39d6417bd53c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[5].resource/*Observation/null*.category[1]]
259	information structure	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)	[Bundle.entry[13].resource.result[0]]
54	information not-supported	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[5].resource/*Observation/null*/]
56	information informational		[Bundle.entry[5].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]

severity	code	details
247	information structure	<p>problem, but you should check that it's not intended to match a slice)</p> <p>Details for urn:uuid:ed0fbcf5-7874-4d67-b5ff-6a34c20188a2 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</p> <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>Binding for path Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)</p> <p>[Bundle.entry[13].resource/*DiagnosticReport/null*/.result]</p>
59	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]</p>
61	information code-invalid	<p>[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]</p>
62	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]</p>
63	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept)]</p>
64	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]</p>
65	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept)]</p>
66	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept)]</p>
67	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)]</p>
68	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[8].value.ofType(string)]</p>
70	information informational	<p>[Bundle.entry[5].resource/*Observation/null*/.component[10].value.ofType(string)]</p>
91	information informational	<p>[Bundle.entry[6].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]</p>
28	information informational	<p>[Bundle.entry[2].resource/*Observation/null*/.component[10].value.ofType(CodeableConcept)]</p>
93	information code-invalid	<p>[Bundle.entry[7].resource/*Observation/null*/.category[1]]</p>

severity	code	details
255 information	not-supported	<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>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)</p>
122 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>
121 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>
120 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>
119 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>
118 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>
117 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>
116 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>
115 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>
114 information	code-invalid	<p>Binding for path <code>Bundle.entry[7].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 has no source, so can't be checked</p>
20 information	informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p>
112 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>
242 information	structure	<code>[Bundle.entry[13].resource/*DiagnosticReport/null*/.result]</code>

	severity	code	details
21	information informational	Details for urn:uuid:6d1c937d-5bf2-44a1-be0c-e5579d8b5e40 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[2].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]
109	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[7].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
231	information not-supported	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) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1)	[Bundle.entry[11].resource/*Observation/null*/]
107	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)	[Bundle.entry[7].resource/*Observation/null*/]
241	information structure	Details for urn:uuid:8730acb2-151e-4dc6-aaff-c8a80c9a37bc matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*/.result]
240	information structure	Details for urn:uuid:4d84c6a3-418c-4da3-8a24-39d6417bd53c matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[13].resource/*DiagnosticReport/null*/.result]
22	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[4].value.ofType(CodeableConcept)]
235	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[11].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
201	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)	[Bundle.entry[10].resource/*Observation/null*/]
236	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[11].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]
26	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[8].value.ofType(CodeableConcept)]
25	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[7].value.ofType(CodeableConcept)]
24	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[6].value.ofType(CodeableConcept)]
23	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[2].resource/*Observation/null*.component[5].value.ofType(CodeableConcept)]

	severity	code	details
27	information	informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a 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/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
234	information	code-invalid	Binding for path Bundle.entry[11].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
226	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)
224	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
227	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
237	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) 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
205	warning	not-found	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)
239	warning	code-invalid	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
196	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
43	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
44	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
46	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
47	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
48	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
42	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
49	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated

	severity	code	details
58	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[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept).coding]
77	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[6].resource/*Observation/null*/]
79	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.identifier.type]
80	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept).coding]
92	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[7].resource/*Observation/null*/]
94	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) 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.identifier.type]
41	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) 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.identifier.type]
1	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]
3	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[2].resource/*Observation/null*/]
5	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.identifier.type]
6	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[1].value.ofType(CodeableConcept).coding]
7	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[2].value.ofType(CodeableConcept).coding]
39	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[5].resource/*Observation/null*/]
8	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[3].value.ofType(CodeableConcept).coding]
10	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[5].value.ofType(CodeableConcept).coding]
11	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[6].value.ofType(CodeableConcept).coding]
12	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[7].value.ofType(CodeableConcept).coding]
13	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*.component[8].value.ofType(CodeableConcept).coding]
34	warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the [Bundle.entry[4].resource/*DocumentReference/null*.subject.identifier.type]

	severity	code	details
9	warning	not-found	validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH) A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
95	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
96	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
97	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
156	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
157	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
158	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
167	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
186	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
155	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
188	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)
190	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
191	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
192	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
193	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
194	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
195	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
189	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
154	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
153	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
152	warning	not-found	

	severity	code	details
		A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated	[Bundle.entry[9].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
98	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
99	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
100	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
101	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
102	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
111	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
129	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
131	warning	code-invalid	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
148	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
151	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated