

| | severity | code | details |
|-----------|-----------|--|---|
| 165 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[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[8].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] |
| 168 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[8].resource/*Observation/null*.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[8].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |
| 277 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)] |
| 204 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[9].resource/*Observation/null*.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[9].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 275 error | structure | Unable to find a profile match for urn:uuid:a271cce9-ed37-409e-99d4-a8932a4ff961 among choices: http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core, http://hl7.org/fhir/StructureDefinition/ImagingStudy | [Bundle.entry[11].resource/*Observation/null*.derivedFrom] |
| 170 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[8].resource/*Observation/null*.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[8].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 201 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[9].resource/*Observation/null*.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[9].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |
| 103 error | structure | Unable to find a profile match for urn:uuid:1cd5764d-354d-4e78-9a25-b324dcb8fd7f among choices: http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core, http://hl7.org/fhir/StructureDefinition/ImagingStudy | [Bundle.entry[6].resource/*Observation/null*.derivedFrom] |
| 81 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[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 250 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 | [Bundle.entry[10].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |

| | severity | code | details |
|-----------|-----------------|---|--|
| 253 error | invalid | CLOSED: Bundle.entry[10].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[10].resource/*Observation/null*.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[10].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 105 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)] |
| 163 error | structure | 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*.derivedFrom[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[8].resource/*Observation/null*.derivedFrom[0].value.ofType(CodeableConcept)] |
| 162 error | structure | Observation.component:clinical-significance: max allowed = 1, but found 2 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) | [Bundle.entry[8].resource/*Observation/null*] |
| 108 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[6].resource/*Observation/null*.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[6].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |
| 298 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[13].resource.category[0]] |
| 306 error | structure | Unable to find a profile match for urn:uuid:8d15e519-01b2-43c7-beea-ab6f0253a05a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation | [Bundle.entry[13].resource.result[4]] |
| 78 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[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) | [Bundle.entry[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |
| 302 error | structure | Unable to find a profile match for urn:uuid:21308855-003f-42ee-ad92-3196f858a63e among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation | [Bundle.entry[13].resource.result[2]] |
| 308 error | structure | Unable to find a profile match for urn:uuid:9fcc7299-d358-4329-b676-84520d5b2a10 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation | [Bundle.entry[13].resource.result[5]] |
| 300 error | structure | Unable to find a profile match for urn:uuid:1cd5764d-354d-4e78-9a25-b324dcb8fd7f among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation | [Bundle.entry[13].resource.result[1]] |
| 138 error | invalid | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ | [Bundle.entry[7].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] |

| | severity | code | details |
|--------------------------------------|-----------------|-------------|--|
| | | | 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)) |
| 304 error | structure | | Unable to find a profile match for urn:uuid:567ffd5e-aef9-4527-90b8-f98f14b66868 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation |
| | | | [Bundle.entry[13].resource.result[3]] |
| 141 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[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) |
| | | | [Bundle.entry[7].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)] |
| 312 error | structure | | Unable to find a profile match for urn:uuid:b6b0de97-b778-4bc8-8629-2ead5a0b326c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation |
| | | | [Bundle.entry[13].resource.result[7]] |
| 310 error | structure | | Unable to find a profile match for urn:uuid:a271cce9-ed37-409c-99d4-a8932a4ff961 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation |
| | | | [Bundle.entry[13].resource.result[6]] |
| 215 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[11].value.ofType(CodeableConcept)] |
| 197 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[1]] |
| 216 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[12].value.ofType(CodeableConcept)] |
| 196 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[1]] |
| 210 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[6].value.ofType(CodeableConcept)] |
| 200 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)] |
| 199 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[1]] |
| 214 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[10].value.ofType(string)] |
| 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 |
| | | | [Bundle.entry[9].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)] |

| severity | code | details |
|-------------------------------|--|--|
| 213 information informational | 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) |
| 212 information informational | 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) |
| 205 information code-invalid | 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 | 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 |
| 206 information informational | 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) |
| 211 information informational | 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) |
| 195 information informational | 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) |
| 208 information informational | 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) |
| 209 information informational | 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) |
| 198 information informational | 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) |
| 207 information informational | A definition could not be found for Canonical URL 'https://fhir.nwgenomics.nhs.uk/Endpoint/IGENE' but a definition can be found for 'http://fhir.nwgenomics.nhs.uk/Endpoint/IGENE' - change from https: to http: | 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) |
| 0 information informational | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) | [Bundle.entry[0].resource/*MessageHeader/null*/.source.endpoint] |
| 193 information informational | 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) |
| 148 information informational | 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) |

| severity | code | details |
|-------------------------------|--|--|
| 149 information informational | problem, but you should check that it's not intended to match a slice) | [Bundle.entry[7].resource/*Observation/null*/.component[8].value.ofType(Range)] |
| 150 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[9].value.ofType(string)] |
| 151 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[10].value.ofType(CodeableConcept)] |
| 152 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[11].value.ofType(CodeableConcept)] |
| 153 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)] |
| 154 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[13].value.ofType(CodeableConcept)] |
| 155 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[14].value.ofType(CodeableConcept)] |
| 156 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[15].value.ofType(CodeableConcept)] |
| 159 information structure | Details for urn:uuid:567ffd5e-aef9-4527-90b8-f98f14b66868 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 | [Bundle.entry[8].resource/*Observation/null*/.derivedFrom] |
| 164 information structure | Details for urn:uuid:567ffd5e-aef9-4527-90b8-f98f14b66868 matching against profile http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eucore 2.0.0-ballot | [Bundle.entry[8].resource/*Observation/null*/.derivedFrom] |
| 166 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/DiagnosticImplication 0.2.1] has no source, so can't be checked | [Bundle.entry[8].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)] |
| 167 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)] |
| 169 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)] |
| 171 information informational | | |

| 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 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 172 | 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) |
| 184 | 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) |
| 185 | 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) |
| 186 | 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) |
| 187 | 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) |
| 188 | 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) |
| 189 | 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) |
| 190 | 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) |
| 191 | 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) |
| 192 | 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) |
| 194 | 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) |
| 217 | 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) |
| 238 | information informational | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ |

| severity | code | details |
|-------------------------------|---|--|
| 219 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) | |
| 265 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[15].value.ofType(CodeableConcept)] |
| 266 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[12].value.ofType(CodeableConcept)] |
| 267 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)] |
| 268 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)] |
| 269 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[15].value.ofType(CodeableConcept)] |
| 270 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]] |
| 272 information structure | Details for urn:uuid:a271cce9-ed37-409e-99d4-a8932a4ff961 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 | [Bundle.entry[11].resource/*Observation/null*.derivedFrom] |
| 276 information structure | Details for urn:uuid:a271cce9-ed37-409e-99d4-a8932a4ff961 matching against profile http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core 2.0.0-ballot | [Bundle.entry[11].resource/*Observation/null*.derivedFrom] |
| 278 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 | [Bundle.entry[11].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] |
| 279 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)] |
| 284 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#R59.3) | [Bundle.entry[13].resource/*DiagnosticReport/null*.code] |
| 286 information structure | Details for urn:uuid:1cd5764d-354d-4e78-9a25-b324dc8fd7f matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 | [Bundle.entry[13].resource/*DiagnosticReport/null*.result] |
| 287 information structure | | [Bundle.entry[13].resource/*DiagnosticReport/null*.result] |

| severity | code | details |
|----------|---------------------------|--|
| 288 | information structure | Details for urn:uuid:21308855-003f-42ee-ad92-3196f858a63e matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 Details for urn:uuid:567ffd5e-aef9-4527-90b8-f98f14b66868 matching against profile http://hl7.org/ [Bundle.entry[13].resource/*DiagnosticReport/null*/.result[.fhir/StructureDefinition/Observation 4.0.1] |
| 289 | information structure | Details for urn:uuid:8d15e519-01b2-43c7-beea-ab6f0253a05a matching against profile http://hl7.org/ [Bundle.entry[13].resource/*DiagnosticReport/null*/.result[.fhir/StructureDefinition/Observation 4.0.1] |
| 290 | information structure | Details for urn:uuid:9fcc7299-d358-4329-b676-84520d5b2a10 matching against profile http://hl7.org/ [Bundle.entry[13].resource/*DiagnosticReport/null*/.result[.fhir/StructureDefinition/Observation 4.0.1] |
| 291 | information structure | Details for urn:uuid:a271cce9-ed37-409e-99d4-a8932a4ff961 matching against profile http://hl7.org/ [Bundle.entry[13].resource/*DiagnosticReport/null*/.result[.fhir/StructureDefinition/Observation 4.0.1] |
| 292 | information structure | Details for urn:uuid:b6b0de97-b778-4bc8-8629-2ead5a0b326c matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 Entry 'urn:uuid:208a360c-7320-467d-bab5-63722296531f' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message) |
| 297 | information informational | [Bundle.entry[15]] |
| 299 | 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) Details for urn:uuid:1cd5764d-354d-4e78-9a25-b324dc8fd7f matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 |
| 301 | information structure | [Bundle.entry[13].resource.result[1]] |
| 303 | information structure | Details for urn:uuid:21308855-003f-42ee-ad92-3196f858a63e matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[2]] |
| 305 | information structure | Details for urn:uuid:567ffd5e-aef9-4527-90b8-f98f14b66868 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[3]] |
| 307 | information structure | Details for urn:uuid:8d15e519-01b2-43c7-beea-ab6f0253a05a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[4]] |
| 309 | information structure | Details for urn:uuid:9fcc7299-d358-4329-b676-84520d5b2a10 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[5]] |
| 311 | information structure | Details for urn:uuid:a271cce9-ed37-409e-99d4-a8932a4ff961 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[6]] |
| 313 | information structure | Details for urn:uuid:b6b0de97-b778-4bc8-8629-2ead5a0b326c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 264 | 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[11].value.ofType(CodeableConcept)] |
| 218 | 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[14].value.ofType(CodeableConcept)] |

| severity | code | details |
|-------------------------------|--|---|
| 263 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[10].value.ofType(string)] |
| 261 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[8].value.ofType(string)] |
| 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[9].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)] |
| 221 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[10].resource/*Observation/null*.category[1]] |
| 233 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*.category[2]] |
| 234 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[1]] |
| 235 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[1]] |
| 236 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[1]] |
| 237 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[1]] |
| 147 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[6].value.ofType(CodeableConcept)] |
| 239 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[1]] |
| 240 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[1]] |
| 241 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[1]] |
| 242 information informational | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ | [Bundle.entry[10].resource/*Observation/null*.component[1]] |

| 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) | |
| 243 | 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[|
| 244 | 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[|
| 245 | 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[|
| 246 | 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[|
| 247 | 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[|
| 248 | 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[|
| 249 | 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)] |
| 252 | 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[1].value.ofType(CodeableConcept)] |
| 254 | information code-invalid | Binding for path <code>Bundle.entry[10].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[10].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 255 | 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[2].value.ofType(CodeableConcept)] |
| 256 | 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[3].value.ofType(CodeableConcept)] |
| 257 | 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[4].value.ofType(CodeableConcept)] |
| 258 | 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[10].resource/*Observation/null*.component[5].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) |
| 259 information informational | | [Bundle.entry[10].resource/*Observation/null*.component[6].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) |
| 260 information informational | | [Bundle.entry[10].resource/*Observation/null*.component[7].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) |
| 262 information informational | | [Bundle.entry[10].resource/*Observation/null*.component[9].value.ofType(Range)] |
| | | 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) |
| 146 information informational | | [Bundle.entry[7].resource/*Observation/null*.component[5].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) |
| 157 information code-invalid | | [Bundle.entry[8].resource/*Observation/null*.category[1]] |
| | | 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/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 144 information informational | | [Bundle.entry[7].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/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 31 information informational | | [Bundle.entry[2].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/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 30 information informational | | [Bundle.entry[2].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/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 29 information informational | | [Bundle.entry[2].resource/*Observation/null*.component[1].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) |
| 80 information informational | | [Bundle.entry[5].resource/*Observation/null*.component[1].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) |
| 82 information code-invalid | | [Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| | | Binding for path <code>Bundle.entry[5].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 |
| 83 information informational | | [Bundle.entry[5].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) |
| 84 information informational | | [Bundle.entry[5].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 |

| | severity | code | details |
|----|---------------------------|---|--|
| 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> | [Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)] |
| 86 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept)] |
| 87 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept)] |
| 32 | 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> | [Bundle.entry[2].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)] |
| 88 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)] |
| 90 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[9].value.ofType(Range)] |
| 91 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[10].value.ofType(string)] |
| 92 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept)] |
| 93 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept)] |
| 94 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept)] |
| 95 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[14].value.ofType(CodeableConcept)] |
| 96 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[15].value.ofType(CodeableConcept)] |
| 97 | 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> | [Bundle.entry[5].resource/*Observation/null*/.component[16].value.ofType(CodeableConcept)] |

| severity | code | details |
|----------|---------------------------|--|
| 49 | information code-invalid | <p>None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE)</p> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 28 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 89 | information informational | <p>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>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 33 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 145 | information informational | <p>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>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 34 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 40 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 39 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 38 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 37 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 36 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 35 | information informational | <p>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> <p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 61 | information informational | <p>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>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> |
| 62 | information informational | <p>in the profile https://fhir.nwgenomics.nhs.uk/</p> |

| | severity | code | details |
|------------|---------------------------|-------------|--|
| | | | problem, but you should check that it's not intended to match a slice) |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 76 | information informational | | [Bundle.entry[5].resource/*Observation/null*.component[1] |
| | | | 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) |
| 77 | information informational | | [Bundle.entry[5].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] |
| | | | Details for urn:uuid:1cd5764d-354d-4e78-9a25-b324dc8fd7f matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 |
| 100 | information structure | | [Bundle.entry[6].resource/*Observation/null*.derivedFrom[|
| | | | 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) |
| 27 | information informational | | [Bundle.entry[2].resource/*Observation/null*.component[1] |
| | | | 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) |
| 98 | information code-invalid | | [Bundle.entry[6].resource/*Observation/null*.category[1]] |
| 140 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1].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) |
| | | | 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) |
| 137 | information informational | | [Bundle.entry[7].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/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 136 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 135 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 134 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 133 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 132 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) |
| 104 | information structure | | [Bundle.entry[6].resource/*Observation/null*.derivedFrom[|
| 131 | information informational | | [Bundle.entry[7].resource/*Observation/null*.component[1] |
| | | | Details for urn:uuid:1cd5764d-354d-4e78-9a25-b324dc8fd7f matching against profile http://hl7.eu/fhir/base/StructureDefinition/medicalTestResult-eu-core 2.0.0-ballot |
| | | | 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 |

| | severity | code | details |
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| | | 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[7].resource/*Observation/null*.category[2]] |
| 122 | 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) | |
| | | 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[3]] |
| 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) | |
| | | 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[4]] |
| 125 | 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) | |
| | | 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[5]] |
| 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) | |
| | | 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[6]] |
| 127 | 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) | |
| | | 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[7]] |
| 128 | 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) | |
| | | 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[8]] |
| 129 | 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) | |
| | | 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[9]] |
| 130 | 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) | |
| | | 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[0]] |
| 123 | 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) | |
| | | 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 | [Bundle.entry[7].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] |
| 142 | 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[2].resource/*Observation/null*.category[1]] |
| 4 | 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) | [Bundle.entry[2].resource/*Observation/null*.component[0]] |
| 15 | 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[1]] |
| 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 | [Bundle.entry[2].resource/*Observation/null*.component[1]] |

| | severity | code | details |
|-----|---------------------------|--|--|
| 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[9] |
| 23 | 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] |
| 109 | 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[1].value.ofType(CodeableConcept)] |
| 143 | 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[2].value.ofType(CodeableConcept)] |
| 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[7] |
| 21 | 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] |
| 20 | 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[5] |
| 19 | 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] |
| 18 | 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[3] |
| 107 | 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)] |
| 17 | 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[2] |
| 106 | information code-invalid | Binding for path <code>Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 has no source, so can't be checked | [Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] |
| 16 | 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 | [Bundle.entry[2].resource/*Observation/null*.component[1] |

| | severity | code | details |
|-----|-----------------|---------------|--|
| 26 | information | informational | <p>may not be a 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/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p> |
| 110 | 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> |
| 2 | warning | code-invalid | <p>None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)</p> |
| 271 | 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> |
| 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> |
| 285 | 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> |
| 273 | warning | not-found | <p>A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated</p> |
| 283 | warning | code-invalid | <p>Unknown Code 'R59.3' 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</p> |
| 13 | warning | not-found | <p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> |
| 12 | warning | not-found | <p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> |
| 5 | 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> |
| 281 | 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> |
| 11 | warning | not-found | <p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> |
| 10 | warning | not-found | <p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> |

| | severity | code | details |
|-----|-----------------|--------------|--|
| 9 | 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[4].value.ofType(CodeableConcept).coding] |
| 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] |
| 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] |
| 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] |
| | | | Unknown Code 'R59.3' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England' |
| 280 | warning | code-invalid | GenomicTestDirectory' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment [Bundle.entry[12].resource/*ServiceRequest/null*.code.code] |
| | | | 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] |
| 230 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/projects/SNP' could not be found, so the code cannot be validated [Bundle.entry[10].resource/*Observation/null*.component[11].value.ofType(CodeableConcept).coding] |
| 52 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated [Bundle.entry[5].resource/*Observation/null*.component[0].value.ofType(CodeableConcept).coding] |
| 115 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*.component[2].value.ofType(CodeableConcept).coding] |
| 116 | warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*.component[3].value.ofType(CodeableConcept).coding] |
| 117 | warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*.component[4].value.ofType(CodeableConcept).coding] |
| 118 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*.component[6].value.ofType(CodeableConcept).coding] |
| 119 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/projects/SNP' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*.component[10].value.ofType(CodeableConcept).coding] |
| 182 | warning | not-found | A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[12].value.ofType(CodeableConcept).coding] |
| 181 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/projects/SNP' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[11].value.ofType(CodeableConcept).coding] |
| 180 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[7].value.ofType(CodeableConcept).coding] |
| 179 | warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[4].value.ofType(CodeableConcept).coding] |
| 178 | warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[3].value.ofType(CodeableConcept).coding] |
| 177 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[2].value.ofType(CodeableConcept).coding] |
| 176 | warning | not-found | 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] |
| 175 | warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*.component[0].value.ofType(CodeableConcept).coding] |
| 174 | warning | not-found | |

| severity | code | details |
|-------------|--------------|--|
| | | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 173 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) |
| 120 warning | not-found | A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated |
| 160 warning | not-found | A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated |
| 158 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) |
| 139 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 |
| 114 warning | not-found | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 113 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated |
| 112 warning | not-found | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 111 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) |
| 53 warning | not-found | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 54 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated |
| 55 warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated |
| 56 warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated |
| 57 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated |
| 58 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 |
| 59 warning | not-found | A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated |
| 251 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 |
| 231 warning | not-found | A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated |
| 51 warning | not-found | [Bundle.entry[9].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system] |

| severity | code | details |
|--------------------|--------------|--|
| | | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 229 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated |
| 227 warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated |
| 226 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated |
| 225 warning | not-found | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 224 warning | not-found | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated |
| 223 warning | not-found | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated |
| 222 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) |
| 79 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 |
| 202 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 |
| 99 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) |
| 101 warning | not-found | A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated |
| 228 warning | not-found | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated |