

severity	code	details
62 error	structure	Unable to find a profile match for urn:uuid:96a207b4-ba2d-4792-bbf8-fca05ee0c4f7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
56 error	invalid	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
55 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
64 error	structure	Unable to find a profile match for urn:uuid:96a207b4-ba2d-4792-bbf8-fca05ee0c4f7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation Unable to find a profile match for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Patient
60 error	structure	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
59 error	structure	Unable to find a profile match for urn:uuid:01544798-1989-40c0-8c84-9739689d43e8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
68 error	structure	Unable to find a profile match for urn:uuid:542a39ce-8761-495c-8830-e807f973a4d7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication
70 error	structure	Unable to find a profile match for urn:uuid:542a39ce-8761-495c-8830-e807f973a4d7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
72 error	structure	Not a valid date format: '1982--0-2'
2 error	invalid	Unable to find a profile match for urn:uuid:01544798-1989-40c0-8c84-9739689d43e8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant
48 information	structure	Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1
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)
57 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)
41 information	structure	Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1
58 information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/

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54	information informational	<p>StructureDefinition/DiagnosticReport 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>Entry 'urn:uuid:0c3e8ee6-2b5d-4474-97ef-793502ee851e'</p> <p>isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p> <p>Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</p> <p>None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE)</p> <p>Details for urn:uuid:542a39ce-8761-495c-8830-e807f973a4d7 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p> <p>Details for urn:uuid:542a39ce-8761-495c-8830-e807f973a4d7 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</p> <p>None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE)</p> <p>Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</p> <p>Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</p> <p>Details for urn:uuid:01544798-1989-40c0-8c84-9739689d43e8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</p> <p>Details for urn:uuid:01544798-1989-40c0-8c84-9739689d43e8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p> <p>Details for urn:uuid:96a207b4-ba2d-4792-bbfb-fca05ee0c4f7 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p> <p>Details for urn:uuid:96a207b4-ba2d-4792-bbfb-fca05ee0c4f7 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</p> <p>Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</p> <p>Details for urn:uuid:4d07bf36-1da5-4eb2-99c7-32a9fe0789cd matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Patient 0.2.1</p> <p>None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier null).subject.identifier.type</p>
45	information structure	[Bundle.entry[7].resource/*ServiceRequest/null*.subject]
25	information code-invalid	[Bundle.entry[5].resource/*Observation/null*.category[1]]
73	information structure	[Bundle.entry[8].resource.result[2]]
71	information structure	[Bundle.entry[8].resource.result[2]]
5	information code-invalid	[Bundle.entry[2].resource/*Observation/null*.category[1]]
6	information structure	[Bundle.entry[2].resource/*Observation/null*.subject]
26	information structure	[Bundle.entry[5].resource/*Observation/null*.subject]
67	information structure	[Bundle.entry[8].resource.result[1]]
69	information structure	[Bundle.entry[8].resource.result[1]]
65	information structure	[Bundle.entry[8].resource.result[0]]
63	information structure	[Bundle.entry[8].resource.result[0]]
18	information structure	[Bundle.entry[4].resource/*DocumentReference/null*.subject]
61	information structure	[Bundle.entry[8].resource.subject]
49	warning code-invalid	[Bundle.entry[8].resource/*DiagnosticReport/null*.subject.identifier.type]

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		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)
46 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)
37 warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
43 warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
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)
4 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
7 warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)
8 warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
9 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
10 warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
11 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
12 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
13 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
19 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)
24 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
28 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
29 warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
30 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
31 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
32 warning	not-found	

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	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated	[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
33 warning	invalid	Coding has no system. A code with no system has no defined meaning, and it cannot be validated. A system should be provided
34 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
35 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
36 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
39 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
42 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)
27 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)