

16, Feb 2026

severity	code	details
109 error	structure	Unable to find a profile match for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[2]]
107 error	structure	Unable to find a profile match for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[2]]
101 error	structure	Unable to find a profile match for urn:uuid:d4da6a50-ac55-47ba-801b-b2388045efc0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[0]]
111 error	structure	Unable to find a profile match for urn:uuid:321d7cb2-087d-41ce-9cfa-37f1dedb4d2c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[3]]
99 error	structure	Unable to find a profile match for urn:uuid:d4da6a50-ac55-47ba-801b-b2388045efc0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel [Bundle.entry[13].resource.result[0]]
113 error	structure	Unable to find a profile match for urn:uuid:321d7cb2-087d-41ce-9cfa-37f1dedb4d2c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 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 [Bundle.entry[13].resource.result[3]]
98 error	structure	Unable to find a profile match for urn:uuid:ce817834-a1a2-4e22-a081-1416ac8f17ea among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.code]
115 error	structure	Unable to find a profile match for urn:uuid:ce817834-a1a2-4e22-a081-1416ac8f17ea among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[4]]
117 error	structure	Unable to find a profile match for urn:uuid:e642034a-d28d-435a-8d29-8e26f0b211bf among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[4]]
105 error	structure	Unable to find a profile match for urn:uuid:cc344c76-11ac-40ac-9789-cf8de6ef0538 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[1]]
119 error	structure	Unable to find a profile match for urn:uuid:f1bb6c3d-8cee-478a-808f-4795e58aa68c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[5]]
129 error	structure	Unable to find a profile match for urn:uuid:f1bb6c3d-8cee-478a-808f-4795e58aa68c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[7]]
127 error	structure	Unable to find a profile match for urn:uuid:f1bb6c3d-8cee-478a-808f-4795e58aa68c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[7]]
95 error	invalid	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator [Bundle.entry[13].resource.performer[0]]
125 error	structure	Unable to find a profile match for urn:uuid:15b75274-ebba-4ebf-8cd1-840901a05cf among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[6]]
123 error	structure	Unable to find a profile match for urn:uuid:15b75274-ebba-4ebf-8cd1-840901a05cf among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[6]]

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121	error	structure	<p>Unable to find a profile match for urn:uuid:cc344c76-11ac-40ac-9789-cf8de6ef0538 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice</p>
94	error	processing	<p>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</p>
103	error	structure	<p>Unable to find a profile match for urn:uuid:e642034a-d28d-435a-8d29-8e26f0b211bf among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant</p>
97	information	informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p>
93	information	informational	<p>Entry 'urn:uuid:0f6c36c6-9f64-4e56-a7cf-c5de6069c7bc' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p>
53	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>
41	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>
100	information	structure	<p>Details for urn:uuid:d4da6a50-ac55-47ba-801b-b2388045efc0 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</p>
58	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>
102	information	structure	<p>Details for urn:uuid:d4da6a50-ac55-47ba-801b-b2388045efc0 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p>
96	information	not-supported	<p>DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required (DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1)</p>
104	information	structure	<p>Details for urn:uuid:e642034a-d28d-435a-8d29-8e26f0b211bf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</p>
70	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>
130	information	structure	<p>Details for urn:uuid:f1bb6c3d-8cee-478a-808f-4795e58aa68c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</p>
128	information	structure	<p>Details for urn:uuid:f1bb6c3d-8cee-478a-808f-4795e58aa68c matching against profile https://</p>

	severity	code	details
4	information code-invalid	fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 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) Details for urn:uuid:15b75274-ebba-4ebf-8cd1-840901a05cf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[2].resource/*Observation/null*.category[1]]
126	information structure	Details for urn:uuid:15b75274-ebba-4ebf-8cd1-840901a05cf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[6]]
124	information structure	Details for urn:uuid:15b75274-ebba-4ebf-8cd1-840901a05cf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1	[Bundle.entry[13].resource.result[6]]
36	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:cc344c76-11ac-40ac-9789-cf8de6ef0538 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[6].resource/*Observation/null*.category[1]]
122	information structure	Details for urn:uuid:cc344c76-11ac-40ac-9789-cf8de6ef0538 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1	[Bundle.entry[13].resource.result[5]]
120	information structure	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:ce817834-a1a2-4e22-a081-1416ac8f17ea matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1	[Bundle.entry[13].resource.result[5]]
82	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:ce817834-a1a2-4e22-a081-1416ac8f17ea matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1	[Bundle.entry[11].resource/*Observation/null*.category[1]]
116	information structure	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:321d7cb2-087d-41ce-9cfa-37f1dedb4d2c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[4]]
24	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE) Details for urn:uuid:321d7cb2-087d-41ce-9cfa-37f1dedb4d2c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1	[Bundle.entry[5].resource/*Observation/null*.category[1]]
114	information structure	Details for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[3]]
112	information structure	Details for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1	[Bundle.entry[13].resource.result[3]]
110	information structure	Details for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[2]]
108	information structure	Details for urn:uuid:ba81a6c7-006e-4dd4-a83c-0dd749ed1608 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1	[Bundle.entry[13].resource.result[2]]
106	information structure	Details for urn:uuid:e642034a-d28d-435a-8d29-8e26f0b21bf matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[1]]
118	information structure	Details for urn:uuid:ce817834-a1a2-4e22-a081-1416ac8f17ea matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[13].resource.result[4]]

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75	warning	not-found	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
76	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
77	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
78	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
79	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
74	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
83	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)
84	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
86	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)
88	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)
81	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
73	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
66	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
71	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)
33	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
32	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
31	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
30	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
29	warning	not-found	[Bundle.entry[5].resource/*Observation/null*.component[3].value.ofType(CodeableConcept).coding]

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

	severity	code	details
			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)
38	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
69	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
67	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
65	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
64	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
63	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
62	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
61	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
60	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
59	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)
57	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
55	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
54	warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)
52	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
50	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
49	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
48	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
47	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
46	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
45	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
44	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated

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43	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[7].resource/*Observation/null*.component[0].value.ofType(CodeableConcept).coding]
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) [Bundle.entry[7].resource/*Observation/null*.subject.identifier]
40	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[7].resource/*Observation/null*/]
72	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[10].resource/*Observation/null*.component[0].value.ofType(CodeableConcept).coding]