

16, Feb 2026

	severity	code	details
0	error	invalid	Object must have some content Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
59	error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
61	error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
64	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))
67	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)) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
89	error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
90	error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions 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))
92	error	invalid	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
110	error	processing	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator
111	error	invalid	Unable to find a profile match for urn:uuid:1e999bd1-eb54-404d-bdf6-efd30cea777 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
113	error	structure	Unable to find a profile match for urn:uuid:d89a409a-b66a-4061-97b6-fae122fc0be4 among choices:
115	error	structure	[Bundle.entry[8].resource.result[0]]
			[Bundle.entry[8].resource.result[1]]

severity	code	details
117 error	structure	<p>https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>Unable to find a profile match for <code>urn:uuid:0151824d-80f1-4fbf-b1ae-101ec87267e1</code> among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</p> <p>Slicing cannot be evaluated: Could not match discriminator (<code>\$this</code>) for slice</p>
60 error	processing	<p><code>Observation.category:genomicsCategory</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [<code>\$this</code>] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (<code>\$this</code>) for slice</p>
21 error	processing	<p><code>Observation.category:genomicsCategory</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [<code>\$this</code>] does not have fixed value, binding or existence assertions</p> <p>Slicing cannot be evaluated: Could not match discriminator (<code>\$this</code>) for slice</p>
20 error	processing	<p><code>Observation.category:genomicsCategory</code> in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [<code>\$this</code>] does not have fixed value, binding or existence assertions</p>
23 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>
81 information	informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p>
80 information	informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p>
79 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>
78 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>
77 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>
76 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>
75 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>
73 information	informational	

	severity	code	details
72	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[6].value.ofType(CodeableConcept)]
22	information not-supported	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1)	[Bundle.entry[5].resource/*Observation/null*.component[5].value.ofType(CodeableConcept)] [Bundle.entry[2].resource/*Observation/null*/]
70	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]
69	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
68	information code-invalid	Binding for path Bundle.entry[5].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	[Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
66	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]
74	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[7].value.ofType(CodeableConcept)]
82	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[15].value.ofType(CodeableConcept)]
84	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE)	[Bundle.entry[6].resource/*Observation/null*.category[1]]
63	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
118	information structure	Details for urn:uuid:0151824d-80f1-4fbfb1ae-101ec87267e1 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[8].resource.result[2]]
116	information structure	Details for urn:uuid:d89a409a-b66a-4061-97b6-fae122fc0be4 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[8].resource.result[1]]
114	information structure	Details for urn:uuid:1e999bd1-eb54-404d-bdf6-efd30ceea777 matching against profile https://	[Bundle.entry[8].resource.result[0]]

	severity	code	details
		fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	
112	information not-supported	DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required (DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1)	[Bundle.entry[8].resource]
4	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), 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]]
109	information informational	Entry 'urn:uuid:99c60d3a-0f0c-4e17-8574-9b7420a4bf7d' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	[Bundle.entry[10]]
104	information structure	Details for urn:uuid:0151824d-80f1-4fbf-b1ae-101ec87267e1 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[8].resource/*DiagnosticReport/null*.result[2]]
103	information structure	Details for urn:uuid:d89a409a-b66a-4061-97b6-fae122fc0be4 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[8].resource/*DiagnosticReport/null*.result[1]]
102	information structure	Details for urn:uuid:1e999bd1-eb54-404d-bdf6-efd30ceea777 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1	[Bundle.entry[8].resource/*DiagnosticReport/null*.result[0]]
100	information code-invalid	None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (http://hl7.org/fhir/ValueSet/report-codes 4.0.1), and a coding is recommended to come from this value set (codes = https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R185.1)	[Bundle.entry[8].resource/*DiagnosticReport/null*.code]
95	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)]
94	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)]
93	information code-invalid	Binding for path Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept) in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 has no source, so can't be checked	[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
91	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1)	[Bundle.entry[6].resource/*Observation/null*/]
86	information business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1	[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
83	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[16].value.ofType(CodeableConcept)]
62	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1)	[Bundle.entry[5].resource/*Observation/null*/]

	severity	code	details
71	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*Observation/null*.component[4].value.ofType(CodeableConcept)]
37	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[15].value.ofType(CodeableConcept)]
31	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].value.ofType(CodeableConcept)]
47	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[5].resource/*Observation/null*.category[1]]
30	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[8].value.ofType(CodeableConcept)]
36	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[14].value.ofType(CodeableConcept)]
29	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[7].value.ofType(CodeableConcept)]
32	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[10].value.ofType(CodeableConcept)]
28	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[6].value.ofType(CodeableConcept)]
38	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[16].value.ofType(CodeableConcept)]
26	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[4].value.ofType(CodeableConcept)]
34	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[12].value.ofType(CodeableConcept)]
25	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]
35	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[2].resource/*Observation/null*.component[13].value.ofType(CodeableConcept)]

	severity	code	details
24	information informational		StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
27	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)
33	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)
96	warning code-invalid		Unknown Code 'R185.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England'- GenomicTestDirectory' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
97	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) Unknown Code 'R185.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England'- GenomicTestDirectory' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
99	warning code-invalid		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 ' http://www.ncbi.nlm.nih.gov/refseq ' 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
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) 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)
2	warning code-invalid		None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH) None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)
101	warning code-invalid		A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
9	warning not-found		A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
10	warning not-found		A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
15	warning not-found		

	severity	code	details
		A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated	[Bundle.entry[2].resource/*Observation/null*/.component[10].value.ofType(CodeableConcept).coding]
13	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
65	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
12	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
57	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
56	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
55	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.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
53	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
52	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
51	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
50	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
49	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
48	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)
16	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
17	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
18	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
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)
85	warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)

	severity	code	details
87	warning	code-invalid	Unknown Code 'R185' in the CodeSystem ' https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication ' version '0.2.1' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment
11	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
14	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated