

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
62	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
64	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
66	error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
68	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))
71	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))
94	error	processing	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
95	error	processing	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions
97	error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 and slicing is CLOSED: Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
98	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
116	error	processing	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator
117	error	invalid	[Bundle.entry[2].resource.component[14].value[x].coding[0]] [Bundle.entry[5].resource/*Observation/null*.category[0]] [Bundle.entry[5].resource/*Observation/null*.category[2]] [Bundle.entry[5].resource/*Observation/null*/] [Bundle.entry[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)] [Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)] [Bundle.entry[6].resource/*Observation/null*.category[0]] [Bundle.entry[6].resource/*Observation/null*.category[1]] [Bundle.entry[6].resource/*Observation/null*/] [Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] [Bundle.entry[8].resource.category[0]] [Bundle.entry[8].resource.performer[0]]

	severity	code	details
119	error	structure	Unable to find a profile match for urn:uuid:09c11cdc-52db-44c4-8be4-9f26749cd5e6 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[0]]
121	error	structure	Unable to find a profile match for urn:uuid:634e003d-d73f-4da5-b3d3-3a038e807cda among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[1]]
123	error	structure	Unable to find a profile match for urn:uuid:6d5a536e-d827-473e-b032-1a197e02f18a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[2]]
63	error	processing	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[5].resource/*Observation/null*.category[1]]
21	error	processing	Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[2].resource/*Observation/null*.category[0]]
24	error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1) Slicing cannot be evaluated: Could not match discriminator (\$this) for slice Observation.category:genomicsCategory in profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 - the discriminator [\$this] does not have fixed value, binding or existence assertions [Bundle.entry[2].resource/*Observation/null*/]
22	error	processing	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[2].resource/*Observation/null*.category[1]]
86	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)]
85	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[14].value.ofType(CodeableConcept)]
84	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[13].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[12].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[11].value.ofType(CodeableConcept)]
81	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[10].value.ofType(string)]
80	information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ [Bundle.entry[5].resource/*Observation/null*.component[9].value.ofType(Range)]

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79	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[5].resource/*Observation/null*.component[8].value.ofType(string)]
77	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)]
76	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[5].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[3].value.ofType(CodeableConcept)]
73	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)]
72	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)]
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[1].value.ofType(CodeableConcept)]
78	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)]
87	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)]
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[1].value.ofType(CodeableConcept)]
89	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]]
124	information structure	Details for urn:uuid:6d5a536e-d827-473e-b032-1a197e02f18a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[8].resource.result[2]]
122	information structure	Details for urn:uuid:634e003d-d73f-4da5-b3d3-3a038e807cda matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	[Bundle.entry[8].resource.result[1]]

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120	information structure	Details for urn:uuid:09c11cdc-52db-44c4-8be4-9f26749cd5e6 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
118	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)
5	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) Entry 'urn:uuid:9ff62c5d-2157-41f0-b44f-26c68789dd33' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)
115	information informational	Details for urn:uuid:6d5a536e-d827-473e-b032-1a197e02f18a matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1
110	information structure	Details for urn:uuid:634e003d-d73f-4da5-b3d3-3a038e807cda matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1
109	information structure	Details for urn:uuid:09c11cdc-52db-44c4-8be4-9f26749cd5e6 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1
108	information structure	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)
106	information code-invalid	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)
101	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)
100	information informational	Binding for path Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)
99	information code-invalid	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1
96	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)
91	information business-rule	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)
67	information informational	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required
65	information not-supported	[Bundle.entry[8].resource.result[0]]
		[Bundle.entry[8].resource]
		[Bundle.entry[2].resource/*Observation/null*.category[1]]
		[Bundle.entry[10]]
		[Bundle.entry[8].resource/*DiagnosticReport/null*.result[2]]
		[Bundle.entry[8].resource/*DiagnosticReport/null*.result[1]]
		[Bundle.entry[8].resource/*DiagnosticReport/null*.result[0]]
		[Bundle.entry[8].resource/*DiagnosticReport/null*.code]
		[Bundle.entry[6].resource/*Observation/null*.component[1].value.ofType(CodeableConcept)]
		[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
		[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
		[Bundle.entry[6].resource/*Observation/null*/]
		[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
		[Bundle.entry[5].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)]
		[Bundle.entry[5].resource/*Observation/null*/]

severity	code	details
75	information informational	(Observation.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (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)
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)
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)
40	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)
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)
23	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/GenomicStudyPanel 0.2.1) 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)
35	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)
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)
50	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) 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)
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)
39	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)
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

	severity	code	details
30	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>
27	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>
29	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>
28	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>
31	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>
9	warning	not-found	<p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</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> <p>Unknown Code 'R185.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England'</p>
102	warning	code-invalid	<p>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>
103	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> <p>Unknown Code 'R185.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England'</p>
105	warning	code-invalid	<p>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>
107	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>
4	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
10	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated</p>
8	warning	not-found	<p>A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated</p>
7	warning	not-found	

	severity	code	details
6	warning	code-invalid	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH) [Bundle.entry[2].resource/*Observation/null*.component[1].value.ofType(CodeableConcept).coding]
16	warning	not-found	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]
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 [Bundle.entry[2].resource/*Observation/null*.component[5].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[2].resource/*Observation/null*.component[6].value.ofType(CodeableConcept).coding]
17	warning	not-found	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[11].value.ofType(CodeableConcept).coding]
15	warning	not-found	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[9].value.ofType(CodeableConcept).coding]
14	warning	not-found	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[8].value.ofType(CodeableConcept).coding]
69	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' version 'null' could not be found, so the code cannot be validated. Valid versions: 3.0.0 or 3.1.0 [Bundle.entry[5].resource/*Observation/null*.component[1].value.ofType(CodeableConcept).coding]
60	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[5].resource/*Observation/null*.component[12].value.ofType(CodeableConcept).coding]
13	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[2].resource/*Observation/null*.component[7].value.ofType(CodeableConcept).coding]
59	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[5].resource/*Observation/null*.component[11].value.ofType(CodeableConcept).coding]
58	warning	not-found	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[7].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[5].resource/*Observation/null*.component[6].value.ofType(CodeableConcept).coding]
56	warning	not-found	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]
55	warning	not-found	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[3].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[5].resource/*Observation/null*.component[2].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]
51	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[5].resource/*Observation/null*.subject.identifi
49	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[5].resource/*Observation/null*/]
18	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[12].value.ofType(CodeableConcept).coding]
19	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
44	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	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
90	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)
92	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
53	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated