

16, Feb 2026

	severity	code	details
0	error	invalid	Object must have some content [Bundle.entry[2].resource.component[14].value[x].coding[0] 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 [Bundle.entry[5].resource/*Observation/null*/.category[0]] 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 [Bundle.entry[5].resource/*Observation/null*/.category[2]]
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)) [Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)] This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
67	error	invalid	[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)] 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 [Bundle.entry[6].resource/*Observation/null*/.category[0]] 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 [Bundle.entry[6].resource/*Observation/null*/.category[1]]
92	error	invalid	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 and slicing is CLOSED: Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity)) [Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)] Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
110	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 [Bundle.entry[8].resource.category[0]]
111	error	invalid	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator [Bundle.entry[8].resource.performer[0]]
113	error	structure	Unable to find a profile match for urn:uuid:1e999bd1-eb54-404d-bdf6-efd30ceea777 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[0]]
115	error	structure	Unable to find a profile match for urn:uuid:d89a409a-b66a-4061-97b6-fae122fc0be4 among choices: [Bundle.entry[8].resource.result[1]]

	severity	code	details
			https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
117	error	structure	Unable to find a profile match for urn:uuid:0151824d-80f1-4fbf-b1ae-101ec87267e1 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[2]]
60	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 [Bundle.entry[5].resource/*Observation/null*/.category[1]]
21	error	processing	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*/.category[1]]
20	error	processing	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*/.category[0]]
23	information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) [Bundle.entry[2].resource/*Observation/null*/.component[1].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[14].value.ofType(CodeableConcept)]
80	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)]
79	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)]
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[11].value.ofType(CodeableConcept)]
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[10].value.ofType(string)]
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[9].value.ofType(Range)]
75	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[8].value.ofType(string)]
73	information	informational	

severity	code	details
		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)]
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[5].value.ofType(CodeableConcept)]
22	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)
		[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-80f1-4fbf-b1ae-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://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
		[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 StructureDefinition[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	<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> <p>[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]</p>
37	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[15].value.ofType(CodeableConcept)]</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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[9].value.ofType(CodeableConcept)]</p>
47	information code-invalid	<p>None of the codings provided are in the value set 'Observation Category Codes' (http://hl7.org/fhir/ValueSet/observation-category 4.0.1), and a coding is recommended to come from this value set (codes = http://terminology.hl7.org/CodeSystem/v2-0074#GE)</p> <p>[Bundle.entry[5].resource/*Observation/null*/.category[1]]</p>
30	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept)]</p>
36	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[14].value.ofType(CodeableConcept)]</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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept)]</p>
32	information informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>[Bundle.entry[2].resource/*Observation/null*/.component[10].value.ofType(CodeableConcept)]</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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept)]</p>
38	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[16].value.ofType(CodeableConcept)]</p>
26	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept)]</p>
34	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept)]</p>
25	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept)]</p>
35	information informational	<p>This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/</p> <p>[Bundle.entry[2].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept)]</p>

	severity	code	details
			StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
24	information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice) [Bundle.entry[2].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
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) [Bundle.entry[2].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept)]
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) [Bundle.entry[2].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept)]
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 [Bundle.entry[7].resource/*ServiceRequest/null*/.code.coding]
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) [Bundle.entry[7].resource/*ServiceRequest/null*/.subject.id]
99	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 [Bundle.entry[8].resource/*DiagnosticReport/null*/.code.coding[0].code]
8	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
7	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[2].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]
6	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[2].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
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) [Bundle.entry[2].resource/*Observation/null*/.subject.identifier]
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) [Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]
101	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[8].resource/*DiagnosticReport/null*/.subject.identifier.type]
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 [Bundle.entry[2].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[2].resource/*Observation/null*/.component[5].value.ofType(CodeableConcept).coding]
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
13	warning	not-found	[Bundle.entry[2].resource/*Observation/null*/.component[10].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
65	warning	not-found	[Bundle.entry[2].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept).coding 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	[Bundle.entry[2].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding 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	[Bundle.entry[5].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
56	warning	not-found	[Bundle.entry[5].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding 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	[Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
54	warning	not-found	[Bundle.entry[5].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept).coding 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	[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
52	warning	not-found	[Bundle.entry[5].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
51	warning	not-found	[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding 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	[Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated
49	warning	not-found	[Bundle.entry[5].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding 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	[Bundle.entry[5].resource/*Observation/null*/.subject.identi 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	[Bundle.entry[2].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
17	warning	not-found	[Bundle.entry[2].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
18	warning	not-found	[Bundle.entry[2].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept).coding A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
42	warning	code-invalid	[Bundle.entry[4].resource/*DocumentReference/null*/.subject.identifier.type] 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	[Bundle.entry[6].resource/*Observation/null*/.subject.identi 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 [Bundle.entry[6].resource/*Observation/null*/.component[0].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[6].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[9].value.ofType(CodeableConcept).coding