

16, Feb 2026

	severity	code	details
105	error	structure	Unable to find a profile match for urn:uuid:974de113-c730-454b-ae69-18f9bd2c0ec2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[2]]
99	error	structure	Unable to find a profile match for urn:uuid:4ad67914-efd6-40bb-9e0b-f1a9cd16f1cc among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[0]]
107	error	structure	Unable to find a profile match for urn:uuid:974de113-c730-454b-ae69-18f9bd2c0ec2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[2]]
98	error	structure	Slice 'DiagnosticReport.code.coding:GenomicTestDirectory': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport[0.2.1]). Note that other slices are allowed in addition to this required slice [Bundle.entry[13].resource.code]
109	error	structure	Unable to find a profile match for urn:uuid:dc330943-75ff-46d2-875d-e1349502bc44 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[3]]
111	error	structure	Unable to find a profile match for urn:uuid:dc330943-75ff-46d2-875d-e1349502bc44 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[3]]
113	error	structure	Unable to find a profile match for urn:uuid:a3e2d98f-606d-492f-a9e8-cb069fcf5739 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[4]]
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]]
115	error	structure	Unable to find a profile match for urn:uuid:a3e2d98f-606d-492f-a9e8-cb069fcf5739 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[4]]
103	error	structure	Unable to find a profile match for urn:uuid:75a85011-ec6b-4c31-9b84-eea486293f21 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[1]]
101	error	structure	Unable to find a profile match for urn:uuid:75a85011-ec6b-4c31-9b84-eea486293f21 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[1]]
94	error	processing	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 [Bundle.entry[13].resource.category[0]]
117	error	structure	Unable to find a profile match for urn:uuid:3ed3e5cc-0956-4079-99e5-94f9a1726bf5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[5]]
119	error	structure	Unable to find a profile match for urn:uuid:3ed3e5cc-0956-4079-99e5-94f9a1726bf5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[5]]
121	error	structure	Unable to find a profile match for urn:uuid:23f6ea6e-f9d8-4ee4-94b5-218fca1117f8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant [Bundle.entry[13].resource.result[6]]
123	error	structure	Unable to find a profile match for urn:uuid:23f6ea6e-f9d8-4ee4-94b5-218fca1117f8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[6]]

	severity	code	details
			fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
127	error	structure	Unable to find a profile match for urn:uuid:78cd25f0-7c24-4295-9dff-b800f0ed8a33 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[13].resource.result[7]]
125	error	structure	Unable to find a profile match for urn:uuid:78cd25f0-7c24-4295-9dff-b800f0ed8a33 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication [Bundle.entry[13].resource.result[7]]
100	information	structure	Details for urn:uuid:4ad67914-efd6-40bb-9e0b-f1a9cd16f1cc matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[13].resource.result[0]]
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) [Bundle.entry[11].resource/*Observation/null*/.category[1]]
58	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[9].resource/*Observation/null*/.category[1]]
97	information	informational	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) [Bundle.entry[13].resource.code.coding[0]]
41	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[7].resource/*Observation/null*/.category[1]]
96	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[13].resource]
93	information	informational	Entry 'urn:uuid:56b375b4-feb9-48d7-b727-fdcdbec6587c' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message) [Bundle.entry[15]]
53	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[8].resource/*Observation/null*/.category[1]]
102	information	structure	Details for urn:uuid:75a85011-ec6b-4c31-9b84-eea486293f21 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 [Bundle.entry[13].resource.result[1]]
70	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[10].resource/*Observation/null*/.category[1]]
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]]
126	information	structure	Details for urn:uuid:78cd25f0-7c24-4295-9dff-b800f0ed8a33 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 [Bundle.entry[13].resource.result[7]]
36	information	code-invalid	[Bundle.entry[6].resource/*Observation/null*/.category[1]]

severity	code	details
124	information structure	None of the codings provided are in the value set 'Observation Category Codes' ( <a href="http://hl7.org/fhir/ValueSet/observation-category">http://hl7.org/fhir/ValueSet/observation-category</a>  4.0.1), and a coding is recommended to come from this value set (codes = <a href="http://terminology.hl7.org/CodeSystem/v2-0074#GE">http://terminology.hl7.org/CodeSystem/v2-0074#GE</a> ) Details for urn:uuid:23f6ea6e-f9d8-4ee4-94b5-218fca1117f8 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[6]]
122	information structure	Details for urn:uuid:23f6ea6e-f9d8-4ee4-94b5-218fca1117f8 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant</a>  0.2.1 [Bundle.entry[13].resource.result[6]]
128	information structure	Details for urn:uuid:78cd25f0-7c24-4295-9dff-b800f0ed8a33 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[7]]
118	information structure	Details for urn:uuid:3ed3e5cc-0956-4079-99e5-94f9a1726bf5 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant</a>  0.2.1 [Bundle.entry[13].resource.result[5]]
116	information structure	Details for urn:uuid:a3e2d98f-606d-492f-a9e8-cb069fcf5739 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[4]]
120	information structure	Details for urn:uuid:3ed3e5cc-0956-4079-99e5-94f9a1726bf5 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[5]]
24	information code-invalid	None of the codings provided are in the value set 'Observation Category Codes' ( <a href="http://hl7.org/fhir/ValueSet/observation-category">http://hl7.org/fhir/ValueSet/observation-category</a>  4.0.1), and a coding is recommended to come from this value set (codes = <a href="http://terminology.hl7.org/CodeSystem/v2-0074#GE">http://terminology.hl7.org/CodeSystem/v2-0074#GE</a> ) [Bundle.entry[5].resource/*Observation/null*/.category[1]]
112	information structure	Details for urn:uuid:dc330943-75ff-46d2-875d-e1349502bc44 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[3]]
110	information structure	Details for urn:uuid:dc330943-75ff-46d2-875d-e1349502bc44 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant</a>  0.2.1 [Bundle.entry[13].resource.result[3]]
108	information structure	Details for urn:uuid:974de113-c730-454b-ae69-18f9bd2c0ec2 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[2]]
106	information structure	Details for urn:uuid:974de113-c730-454b-ae69-18f9bd2c0ec2 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication</a>  0.2.1 [Bundle.entry[13].resource.result[2]]
104	information structure	Details for urn:uuid:75a85011-ec6b-4c31-9b84-eea486293f21 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>  0.2.1 [Bundle.entry[13].resource.result[1]]
114	information structure	Details for urn:uuid:a3e2d98f-606d-492f-a9e8-cb069fcf5739 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication</a>  0.2.1 [Bundle.entry[13].resource.result[4]]
73	warning not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated [Bundle.entry[10].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding
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 [Bundle.entry[10].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding

	severity	code	details
75	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[10].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
76	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[10].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[10].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[10].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding]
79	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[10].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding]
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]
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) [Bundle.entry[11].resource/*Observation/null*/.subject.identifier]
84	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[11].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
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) [Bundle.entry[12].resource/*ServiceRequest/null*/.subject.identifier.type]
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) [Bundle.entry[13].resource/*DiagnosticReport/null*/.subject.identifier.type]
81	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[11].resource/*Observation/null*/]
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 [Bundle.entry[9].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding]
69	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[10].resource/*Observation/null*/]
33	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]
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 [Bundle.entry[5].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding]
30	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]
29	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]
28	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]
27	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated [Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
26	warning	not-found	

	severity	code	details
			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
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)
			[Bundle.entry[5].resource/*Observation/null*/.subject.iden
23	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
			[Bundle.entry[5].resource/*Observation/null*/]
35	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
			[Bundle.entry[6].resource/*Observation/null*/]
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)
			[Bundle.entry[4].resource/*DocumentReference/null*/.subject.identifier.type]
13	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[8].value.ofType(CodeableConcept).codin
12	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[7].value.ofType(CodeableConcept).codin
11	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[6].value.ofType(CodeableConcept).codin
10	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[5].value.ofType(CodeableConcept).codin
9	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[4].value.ofType(CodeableConcept).codin
8	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[3].value.ofType(CodeableConcept).codin
7	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[2].value.ofType(CodeableConcept).codin
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).codin
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.iden
3	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
			[Bundle.entry[2].resource/*Observation/null*/]
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)
			[Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]
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)
			[Bundle.entry[10].resource/*Observation/null*/.subject.iden
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 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[6].resource/*Observation/null*/.subject.iden

	severity	code	details
67	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[9].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding]
64	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
63	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[9].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]
61	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated [Bundle.entry[9].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[9].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
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) [Bundle.entry[9].resource/*Observation/null*/.subject.identifier]
57	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[9].resource/*Observation/null*/]
38	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
55	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated [Bundle.entry[8].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
52	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[8].resource/*Observation/null*/]
50	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*/.component[11].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[7].resource/*Observation/null*/.component[10].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[7].resource/*Observation/null*/.component[6].value.ofType(CodeableConcept).coding]
47	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
46	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
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 [Bundle.entry[7].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]
44	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated [Bundle.entry[7].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
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*/]

	severity	code	details
54	warning	code-invalid	<p>None of the codings provided are in the value set 'IdentifierType' (<a href="http://hl7.org/fhir/ValueSet/identifier-type 4.0.1">http://hl7.org/fhir/ValueSet/identifier-type 4.0.1</a>), 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 = <a href="http://terminology.hl7.org/CodeSystem/v2-0203#NH">http://terminology.hl7.org/CodeSystem/v2-0203#NH</a>)</p> <p>[Bundle.entry[8].resource/*Observation/null*/.subject.iden</p>