

16, Feb 2026

severity	code	details
98 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[8].resource.category[0]]
46 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1) [Bundle.entry[5].resource/*Observation/null*/]
44 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[2]]
43 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]]
42 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[0]]
77 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 [Bundle.entry[6].resource/*Observation/null*/.category[1]]
48 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)]
51 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)) [Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)]
11 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]]
101 error	structure	Unable to find a profile match for urn:uuid:e12cc617-eac0-43e5-b68e-54c6f1a7e5c5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[0]]
76 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] [Bundle.entry[6].resource/*Observation/null*/.category[0]]

severity	code	details
		does not have fixed value, binding or existence assertions
105 error	structure	Unable to find a profile match for urn:uuid:86eb4381-dea9-4542-b600-c530ad4eb314 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[2]]
79 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1) [Bundle.entry[6].resource/*Observation/null*/]
13 error	structure	Observation.effective[x]: minimum required = 1, but only found 0 (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1) [Bundle.entry[2].resource/*Observation/null*/]
80 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)]
103 error	structure	Unable to find a profile match for urn:uuid:1dd8f255-e922-4af7-9353-697d55da9497 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[8].resource.result[1]]
10 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]]
99 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]]
56 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)]
55 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)]
67 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)]
57 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)]
53 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)]
52 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)]
83 information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/ [Bundle.entry[6].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]

severity	code	details
		StructureDefinition/DiagnosticImplication 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
58	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)]
73	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)]
60	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(string)]
68	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[17].value.ofType(CodeableConcept)]
81	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)]
61	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(Range)]
62	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(string)]
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[12].value.ofType(CodeableConcept)]
71	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]]
82	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)]
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[18].value.ofType(CodeableConcept)]
64	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)]
65	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)]

severity	code	details
		problem, but you should check that it's not intended to match a slice)
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[15].value.ofType(CodeableConcept)]
59	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(CodeableConcept)]
54	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)]
78	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*/]
20	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)]
14	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)]
15	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)]
16	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)]
17	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(string)]
18	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)]
19	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)]
50	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)]
97	information informational	Entry 'urn:uuid:d90a5a99-e6ab-44f7-bf1a-5e2eece92554' 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]]
102	information structure	[Bundle.entry[8].resource.result[0]]

	severity	code	details
			Details for urn:uuid:e12cc617-eac0-43e5-b68e-54c6f1a7c5c5 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
30	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]]
92	information	structure	Details for urn:uuid:86eb4381-dea9-4542-b600-c530ad4eb314 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[8].resource/*DiagnosticReport/null*/.result[2]]
91	information	structure	Details for urn:uuid:1dd8f255-e922-4af7-9353-697d55da9497 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[8].resource/*DiagnosticReport/null*/.result[1]]
90	information	structure	Details for urn:uuid:e12cc617-eac0-43e5-b68e-54c6f1a7c5c5 matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1 [Bundle.entry[8].resource/*DiagnosticReport/null*/.result[0]]
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]]
88	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#R210.2) [Bundle.entry[8].resource/*DiagnosticReport/null*/.code]
104	information	structure	Details for urn:uuid:1dd8f255-e922-4af7-9353-697d55da9497 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[8].resource.result[1]]
106	information	structure	Details for urn:uuid:86eb4381-dea9-4542-b600-c530ad4eb314 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1 [Bundle.entry[8].resource.result[2]]
45	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*/]
47	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)]
12	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*/]
100	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]
84	warning	code-invalid	Unknown Code 'R210.2' 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]
89	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 [Bundle.entry[8].resource/*DiagnosticReport/null*/.subject.identifier.type]

	severity	code	details
85	warning	code-invalid	<p>validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)</p> <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>[Bundle.entry[7].resource/*ServiceRequest/null*/.subject.id</p>
87	warning	code-invalid	<p>Unknown Code 'R210.2' 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</p> <p>[Bundle.entry[8].resource/*DiagnosticReport/null*/.code.coding[0].code]</p>
74	warning	code-invalid	<p>Unknown Code 'R210' 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</p> <p>[Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding</p>
1	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>[Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]</p>
3	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p> <p>[Bundle.entry[2].resource/*Observation/null*/]</p>
5	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>[Bundle.entry[2].resource/*Observation/null*/.subject.identi</p>
6	warning	not-found	<p>A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated</p> <p>[Bundle.entry[2].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding</p>
7	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> <p>[Bundle.entry[2].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding</p>
8	warning	not-found	<p>A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated</p> <p>[Bundle.entry[2].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding</p>
24	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>[Bundle.entry[4].resource/*DocumentReference/null*/.subject.identifier.type]</p>
29	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p> <p>[Bundle.entry[5].resource/*Observation/null*/]</p>
31	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>[Bundle.entry[5].resource/*Observation/null*/.subject.identi</p>
32	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> <p>[Bundle.entry[5].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding</p>
34	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> <p>[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding</p>
35	warning	not-found	<p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p> <p>[Bundle.entry[5].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding</p>
36	warning	not-found	<p>A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated</p> <p>[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding</p>

	severity	code	details
37	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
38	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[8].value.ofType(CodeableConcept).coding
39	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[12].value.ofType(CodeableConcept).coding
40	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[13].value.ofType(CodeableConcept).coding
49	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
70	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[6].resource/*Observation/null*/]
72	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.identi
33	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