

	severity	code	details
44	error	invalid	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))  Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
41	error	processing	Observation.category:genomicsCategory in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> - the discriminator [\$this] does not have fixed value, binding or existence assertions  Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
40	error	processing	Observation.category:genomicsCategory in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> - the discriminator [\$this] does not have fixed value, binding or existence assertions  Slicing cannot be evaluated: Could not match discriminator (\$this) for slice
39	error	processing	Observation.category:genomicsCategory in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> - the discriminator [\$this] does not have fixed value, binding or existence assertions  This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a> and slicing is CLOSED: Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
74	error	invalid	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice  DiagnosticReport.category:Genetics in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1</a> - the discriminator [\$this] does not have fixed value, binding or existence assertions
92	error	processing	Profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1</a> , Element matches more than one slice - organization, operator  This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> and slicing is CLOSED: Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept): Does not match slice 'valueQuantity' (discriminator: (\$this is Quantity))
93	error	invalid	Unable to find a profile match for urn:uuid:f0e67905-a55f-4bad-92dc-132ceb5dac1f among choices: <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>
95	error	structure	Unable to find a profile match for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac among choices: <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation</a>
99	error	structure	Slicing cannot be evaluated: Could not match discriminator (\$this) for slice  Observation.category:genomicsCategory in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a> - the discriminator [\$this] does not have fixed value, binding or existence assertions
72	error	processing	[Bundle.entry[6].resource/*Observation/null*/.category[1]]
71	error	processing	[Bundle.entry[6].resource/*Observation/null*/.category[0]]

	<b>severity</b>	<b>code</b>	<b>details</b>
10	error	processing	<p>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</p> <p>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</p> <p>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</p>
9	error	processing	[Bundle.entry[2].resource/*Observation/null*.category[0]]
97	error	structure	[Bundle.entry[8].resource.result[1]]
73	information	not-supported	[Bundle.entry[6].resource/*Observation/null*/]
49	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
50	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]
52	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[5].value.ofType(CodeableConcept)]
53	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[6].value.ofType(CodeableConcept)]
54	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[7].value.ofType(CodeableConcept)]
55	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[8].value.ofType(CodeableConcept)]
56	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[9].value.ofType(string)]
61	information	informational	[Bundle.entry[5].resource/*Observation/null*.component[14].value.ofType(CodeableConcept)]

	<b>severity</b>	<b>code</b>	<b>details</b>
		problem, but you should check that it's not intended to match a slice)	
62	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
77	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a> (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)]
57	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
58	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
59	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
68	information business-rule	Reference to draft CodeSystem <a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1">https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1</a> Binding for path Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept) in profile StructureDefinition[ <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> ] has no source, so can't be checked	[Bundle.entry[6].resource/*Observation/null*.component[0].value.ofType(CodeableConcept)] [Bundle.entry[5].resource/*Observation/null*.component[2].value.ofType(CodeableConcept)]
48	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 4.0.1">http://hl7.org/fhir/ValueSet/observation-category 4.0.1</a> ), 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[6].resource/*Observation/null*.category[1]]
66	information code-invalid	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
65	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
64	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
63	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
60	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (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)]
76	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a> (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)]
51	information informational		

severity	code	details
82	information code-invalid	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice) None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' ( <a href="http://hl7.org/fhir/ValueSet/report-codes 4.0.1">http://hl7.org/fhir/ValueSet/report-codes 4.0.1</a> ), and a coding is recommended to come from this value set (codes = <a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R210.2">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R210.2</a> )
17	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)
100	information structure	Details for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</a>
3	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 4.0.1">http://hl7.org/fhir/ValueSet/observation-category 4.0.1</a> ), 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> )
98	information structure	Details for urn:uuid:ee0ba942-c6ec-4921-937cd4c88e77abcd matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</a>
96	information structure	Details for urn:uuid:f0e67905-a55f-4bad-92dc-132ceb5dac1f matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1</a>
94	information not-supported	DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required (DiagnosticReport.category) due to lack of slicing validation (from <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1</a> )
11	information not-supported	Observation.category:genomicsCategory: Found 0 matches, but unable to check minimum required (Observation.category) due to lack of slicing validation (from <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> )
12	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)
13	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)
14	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)
15	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)
16	information informational	This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this

	<b>severity</b>	<b>code</b>	<b>details</b>
46	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 <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>Binding for path <code>Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)</code></p>
75	information code-invalid		<p><code>Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)</code> in profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a> has no source, so can't be checked</p> <p>None of the codings provided are in the value set 'Observation Category Codes' (<a href="http://hl7.org/fhir/ValueSet/observation-category 4.0.1">http://hl7.org/fhir/ValueSet/observation-category 4.0.1</a>), 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>)</p>
27	information code-invalid		<p><code>Bundle.entry[5].resource/*Observation/null*/.category[1]</code></p> <p>Entry '<code>urn:uuid:a3ea5c8d-7f22-4df9-8b3b-f0066d830ae8</code>' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p>
91	information informational		<p><code>Bundle.entry[10]</code></p> <p>Details for <code>urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac</code> matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Observation 4.0.1">http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</a></p>
86	information structure		<p>Details for <code>urn:uuid:ee0ba942-c6ec-4921-937c-d4c88e77abcd</code> matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Observation 4.0.1">http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</a></p>
85	information structure		<p>Details for <code>urn:uuid:f0e67905-a55f-4bad-92dc-132ceb5dac1f</code> matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Observation 4.0.1">http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</a></p>
84	information structure		<p>Details for <code>urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac</code> matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Observation 4.0.1">http://hl7.org/fhir/StructureDefinition/Observation 4.0.1</a></p>
42	information not-supported		<p><code>Bundle.entry[5].resource/*Observation/null*/</code></p> <p><code>Observation.category:genomicsCategory</code>: Found 0 matches, but unable to check minimum required (<code>Observation.category</code>) due to lack of slicing validation (from <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a>)</p>
43	information informational		<p>This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)</p>
18	information informational		<p>This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicStudyPanel 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)</p>
78	warning code-invalid		<p>Unknown Code 'R210.2' in the CodeSystem '<a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory</a>' 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>
83	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>
81	warning code-invalid		<p>Unknown Code 'R210.2' in the CodeSystem '<a href="https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory">https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory</a>' 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>
79	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</p>

severity	code	details
69 warning	code-invalid	<p>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>Unknown Code 'R210' in the CodeSystem '<a href="https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication">https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication</a>' 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>
1 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>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>
4 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>
5 warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
6 warning	not-found	A definition for CodeSystem ' <a href="http://www.ncbi.nlm.nih.gov/refseq">http://www.ncbi.nlm.nih.gov/refseq</a> ' could not be found, so the code cannot be validated
7 warning	not-found	A definition for CodeSystem ' <a href="http://www.genenames.org">http://www.genenames.org</a> ' could not be found, so the code cannot be validated
22 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>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>
28 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>
29 warning	not-found	A definition for CodeSystem ' <a href="http://www.ncbi.nlm.nih.gov/clinvar">http://www.ncbi.nlm.nih.gov/clinvar</a> ' could not be found, so the code cannot be validated
30 warning	not-found	A definition for CodeSystem ' <a href="http://www.genenames.org">http://www.genenames.org</a> ' could not be found, so the code cannot be validated
31 warning	not-found	A definition for CodeSystem ' <a href="http://www.ncbi.nlm.nih.gov/refseq">http://www.ncbi.nlm.nih.gov/refseq</a> ' could not be found, so the code cannot be validated
32 warning	not-found	A definition for CodeSystem ' <a href="http://varnomen.hgvs.org">http://varnomen.hgvs.org</a> ' could not be found, so the code cannot be validated
33 warning	not-found	A definition for CodeSystem ' <a href="http://varnomen.hgvs.org">http://varnomen.hgvs.org</a> ' could not be found, so the code cannot be validated
34 warning	not-found	A definition for CodeSystem ' <a href="http://www.ncbi.nlm.nih.gov/refseq">http://www.ncbi.nlm.nih.gov/refseq</a> ' could not be found, so the code cannot be validated
35 warning	not-found	A definition for CodeSystem ' <a href="http://varnomen.hgvs.org">http://varnomen.hgvs.org</a> ' could not be found, so the code cannot be validated
36 warning	not-found	A definition for CodeSystem ' <a href="http://www.ncbi.nlm.nih.gov/projects/SNP">http://www.ncbi.nlm.nih.gov/projects/SNP</a> ' could not be found, so the code cannot be validated
37 warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated

	<b>severity</b>	<b>code</b>	<b>details</b>
45	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]
67	warning	code-invalid	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> ) [Bundle.entry[6].resource/*Observation/null*/.subject.identifi