

	severity	code	details
67	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)) 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)) 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)) Unable to find a profile match for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 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 Unable to find a profile match for urn:uuid:ee0ba942-c6ec-4921-937c-d4c88e77abcd among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 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) 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) 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) 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 This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
64	error	invalid	[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept)] [Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
92	error	invalid	[Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]
113	error	structure	[Bundle.entry[8].resource.result[2]]
109	error	processing	[Bundle.entry[8].resource.category[0]]
111	error	structure	[Bundle.entry[8].resource.result[1]]
59	information	informational	[Bundle.entry[5].resource/*Observation/null*/.component[1]]
66	information	informational	[Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept)]
53	information	informational	[Bundle.entry[5].resource/*Observation/null*/.component[9]]
54	information	informational	[Bundle.entry[5].resource/*Observation/null*/.component[1]]
93	information	code-invalid	[Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept)]
55	information	informational	[Bundle.entry[5].resource/*Observation/null*/.component[1]]

	severity	code	details
56	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
63	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
62	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
57	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
60	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
94	information informational	StructureDefinition/DiagnosticImplication 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
61	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
58	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
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	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
86	information code-invalid	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)	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)
85	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
84	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
83	information informational	StructureDefinition/Variant 0.2.1 (this may not be a	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

	severity	code	details
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)]
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(Range)]
51	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)]
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)]
49	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)]
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[2].value.ofType(CodeableConcept)]
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[1].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[8].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[7].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[6].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[5].value.ofType(CodeableConcept)]
13	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(CodeableConcept)]
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[3].value.ofType(CodeableConcept)]
12	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	[Bundle.entry[2].resource/*Observation/null*.component[3].value.ofType(CodeableConcept)]

	severity	code	details
10	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
9	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
108	information informational	Entry 'urn:uuid:a3ea5c8d-7f22-4df9-8b3b-f0066d830ae8' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
110	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)	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)
112	information structure	Details for urn:uuid:ee0ba942-c6ec-4921-937c-d4c88e77abcd matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	Details for urn:uuid:ee0ba942-c6ec-4921-937c-d4c88e77abcd matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
3	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)	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)
114	information structure	Details for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1	Details for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.1
11	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
50	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
21	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
23	information informational	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
48	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
47	information informational	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)	StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
46	information informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/

	severity	code	details
			StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
			This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
45	information informational		StructureDefinition/Variant 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
88	information business-rule		Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.2.1
			This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
22	information informational		StructureDefinition/GenomicStudyPanel 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
			This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/
91	information informational		StructureDefinition/DiagnosticImplication 0.2.1 (this may not be a problem, but you should check that it's not intended to match a slice)
102	information structure		Details for urn:uuid:ee0ba942-c6ec-4921-937cd4c88e77abcd matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1
103	information structure		Details for urn:uuid:5f2e9a9d-95ba-4179-b747-7ff59595b8ac matching against profile http://hl7.org/fhir/StructureDefinition/Observation 4.0.1
33	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)
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)
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#R210.2)
99	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
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)
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)
96	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
89	warning	code-invalid	Unknown Code 'R210' in the CodeSystem ' https://fhir.nwgenomics.nhs.uk/CodeSystem/

	severity	code	details
1	warning	code-invalid	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 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)
4	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)
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 ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
7	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
28	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)
34	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)
35	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
36	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
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
38	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
39	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
40	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
41	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
42	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
43	warning	not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated
65	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
87	warning	code-invalid	[Bundle.entry[1].resource/*Patient/null*/.identifier[0].type]
			[Bundle.entry[2].resource/*Observation/null*/.subject.identifier.type]
			[Bundle.entry[2].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
			[Bundle.entry[2].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]
			[Bundle.entry[2].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
			[Bundle.entry[4].resource/*DocumentReference/null*/.subject.identifier.type]
			[Bundle.entry[5].resource/*Observation/null*/.subject.identifier.type]
			[Bundle.entry[5].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[2].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[3].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[12].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept).coding]
			[Bundle.entry[5].resource/*Observation/null*/.component[1].value.ofType(CodeableConcept).coding]
			[Bundle.entry[6].resource/*Observation/null*/.subject.identifier.type]

severity	code	details
		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)