

14, Feb 2026

	severity	code	details
0	error	invalid	Object must have some content [Bundle.entry[16].resource.value[x].coding[0]]
175	error	structure	Unable to find a profile match for urn:uuid:9b005ceb-e7bd-4bd2-a249-85201fef8815 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[6]]
207	error	structure	Unable to find a profile match for urn:uuid:4f8af9e7-4b76-4e3c-bccc-51d171a5afff among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[22]]
167	error	structure	Unable to find a profile match for urn:uuid:87646305-1486-4c9b-83b9-d9fdabf749bb among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[2]]
205	error	structure	Unable to find a profile match for urn:uuid:0beec652-fe5c-4cdc-ba59-e1d68ab1004d among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[21]]
203	error	structure	Unable to find a profile match for urn:uuid:2ab95d30-f7b8-42ab-ba07-832e5e7243f8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[20]]
169	error	structure	Unable to find a profile match for urn:uuid:1441e38c-0659-4356-bbae-3ad21c212802 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[3]]
201	error	structure	Unable to find a profile match for urn:uuid:1bc984b9-bba9-438e-8390-1d26d54a289a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[19]]
199	error	structure	Unable to find a profile match for urn:uuid:12677216-0d39-43c9-98a3-28146e96516c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[18]]
171	error	structure	Unable to find a profile match for urn:uuid:5f319603-e25a-4b48-90e3-c33dd1233523 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[4]]
197	error	structure	Unable to find a profile match for urn:uuid:e68338e0-3b55-453b-bf8e-5a1a62ac370a among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[17]]
195	error	structure	Unable to find a profile match for urn:uuid:cf131873-be9c-41f4-b70b-dbcb7f82d8ba among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[16]]
173	error	structure	Unable to find a profile match for urn:uuid:bbb8be3d-bb12-4abe-9fe3-28b92a3f5169 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[5]]
209	error	structure	Unable to find a profile match for urn:uuid:abf23bd2-7463-47a5-855a-e3d393f4287c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[23]]
193	error	structure	Unable to find a profile match for urn:uuid:4973f2c8-eb95-4507-bd5a-db8f008f1bea among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[15]]
191	error	structure	Unable to find a profile match for urn:uuid:0ef86e5c-fe06-4877-932f-6c29ff48633e among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[14]]
189	error	structure	Unable to find a profile match for urn:uuid:ed9e5f61-cb96-4bc5-9075-b1f9c1f74f66 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation [Bundle.entry[42].resource.result[13]]

severity	code	details
187 error	structure	fhir.nwgenomics.nhs.uk/StructureDefinition/Observation Unable to find a profile match for urn:uuid:13d48508-9836-4d35-a7b7-b8e693ed1a54 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
185 error	structure	Unable to find a profile match for urn:uuid:a675-40e2-abfd-4e53e35e63d2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
183 error	structure	Unable to find a profile match for urn:uuid:b5c57a4f-618e-4960-aa41-dbc32d8dec88 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
181 error	structure	Unable to find a profile match for urn:uuid:b5c6-4988-a27e-8e999d4b99a2 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
179 error	structure	Unable to find a profile match for urn:uuid:63fbea93-7607-4bb4-a72a-7fd57a73d0a5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
165 error	structure	Unable to find a profile match for urn:uuid:aec68303-151c-4281-821c-547c1693d826 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
177 error	structure	Unable to find a profile match for urn:uuid:256e20b0-5116-4bdd-94df-df2384174222 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
225 error	structure	Unable to find a profile match for urn:uuid:d12c-4585-8166-8a6cc9bf68c8 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
231 error	structure	Unable to find a profile match for urn:uuid:fc1a13dc-b6fe-4dd2-afda-22c549156048 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation-Variant
229 error	structure	Unable to find a profile match for urn:uuid:bb396bdd-0a5a-4f82-a5e6-8430202f29bc among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
227 error	structure	Unable to find a profile match for urn:uuid:c382-4875-987c-6f666bea4e50 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
157 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 0.2.0 - the discriminator [\$this] does not have fixed value, binding or existence assertions
211 error	structure	Unable to find a profile match for urn:uuid:5b6eb6a5-dad0-4be1-b056-2cbfc14ec406 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
158 error	invalid	Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0, Element matches more than one slice - organization, operator
223 error	structure	Unable to find a profile match for urn:uuid:050cf12b-71af-4b42-bf08-05311fd6d4af among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation
151 error	invalid	No definition could be found for URL value 'http://hl7.org/fhir/diagnosticreport-relevant-information-types'

severity	code	details
233 error	structure Unable to find a profile match for urn:uuid:fc1a13dc-b6fe-4dd2-afda-22c549156048 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[34]]
235 error	structure Unable to find a profile match for urn:uuid:afec69a1-659d-45d8-9f3e-5d4ff81a0528 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[35]]
160 error	structure DiagnosticReport.supportingInfo is not allowed to be used at this point (this element is [Bundle.entry.resource]; allowed for this version = e:DiagnosticReport)	[Bundle.entry[42].resource]
149 error	structure Invalid Resource target type. Found DocumentReference, but expected one of ([Observation, Procedure, DiagnosticReport])	[Bundle.entry[42].resource/*DiagnosticReport/null*.extension[0].extension[0].value.ofType(Reference)]
219 error	structure Unable to find a profile match for urn:uuid:1fdb32f4-67c6-45dd-9546-b570125680cd among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[28]]
237 error	structure Unable to find a profile match for urn:uuid:4a855a49-8900-498c-97bc-0dfdad1281d5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[36]]
217 error	structure Unable to find a profile match for urn:uuid:8b25e6d2-3ca1-40b1-a672-02575e4fb2b5 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[27]]
148 error	structure The type 'DocumentReference' is not a valid Target for this element (must be one of [CanonicalType[http://hl7.org/fhir/StructureDefinition/Procedure], CanonicalType[http://hl7.org/fhir/StructureDefinition/Observation], CanonicalType[http://hl7.org/fhir/StructureDefinition/DiagnosticReport]])	[Bundle.entry[42].resource/*DiagnosticReport/null*.extension[0].extension[0].value.ofType(Reference)]
162 error	structure Slice 'DiagnosticReport.code.coding:GenomicTestDirectory': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0). Note that other slices are allowed in addition to this required slice	[Bundle.entry[42].resource.code]
215 error	structure Unable to find a profile match for urn:uuid:ba71c65c-0ac2-45fe-89c4-3f7756abe39c among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[26]]
163 error	structure Unable to find a profile match for urn:uuid:2377ec82-69a1-41de-8103-a2daf8d851ea among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[0]]
213 error	structure Unable to find a profile match for urn:uuid:b550f3d4-91d4-4c4a-98a9-e69a99cea773 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[25]]
221 error	structure Unable to find a profile match for urn:uuid:ab4a6ded-b603-4890-9beb-28bfef29afe0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation	[Bundle.entry[42].resource.result[29]]
164 information	structure Details for urn:uuid:2377ec82-69a1-41de-8103-a2daf8d851ea matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[0]]
172 information	structure Details for urn:uuid:5f319603-e25a-4b48-90e3-c33dd1233523 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[4]]
159 information	not-supported DiagnosticReport.category:Genetics: Found 0 matches, but unable to check minimum required	[Bundle.entry[42].resource]

severity	code	details
170 information structure	(DiagnosticReport.category) due to lack of slicing validation (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0)	[Bundle.entry[42].resource.result[3]]
161 information informational	Details for urn:uuid:1441e38c-0659-4356-bbae-3ad21c212802 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[42].resource.code.coding[0]]
168 information structure	Details for urn:uuid:87646305-1486-4c9b-83b9-d9fdabf749bb matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[2]]
174 information structure	Details for urn:uuid:bbb8be3d-bb12-4abe-9fe3-28b92a3f5169 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[5]]
166 information structure	Details for urn:uuid:aec68303-151c-4281-821c-547c1693d826 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[1]]
178 information structure	Details for urn:uuid:256e20b0-5116-4bdd-94df-df2384174222 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[7]]
220 information structure	Details for urn:uuid:1fdb32f4-67c6-45dd-9546-b570125680cd matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[28]]
238 information structure	Details for urn:uuid:4a855a49-8900-498c-97bc-0dfdad1281d5 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[36]]
236 information structure	Details for urn:uuid:afec69a1-659d-45d8-9f3e-5d4ff81a0528 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[35]]
234 information structure	Details for urn:uuid:fc1a13dc-b6fe-4dd2-afda-22c549156048 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[34]]
176 information structure	Details for urn:uuid:9b005ceb-e7bd-4bd2-a249-85201fef8815 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[6]]
232 information structure	Details for urn:uuid:fc1a13dc-b6fe-4dd2-afda-22c549156048 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation-Variant 0.2.0	[Bundle.entry[42].resource.result[34]]
230 information structure	Details for urn:uuid:bb396bdd-0a5a-4f82-a5e6-8430202f29bc matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[33]]
228 information structure	Details for urn:uuid:9d215f0c-c382-4875-987c-6f66bea4e50 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[32]]
226 information structure	Details for urn:uuid:8c2d7c1e-d12c-4585-8166-8a6cc9bf6c8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0	[Bundle.entry[42].resource.result[31]]

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224	information structure	Details for urn:uuid:050cf12b-71af-4b42-bf08-05311fd6d4af matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[30]]
222	information structure	Details for urn:uuid:ab4a6ded-b603-4890-9beb-28fbef29afe0 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[29]]
218	information structure	Details for urn:uuid:8b25e6d2-3ca1-40b1-a672-02575e4fb2b5 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[27]]
216	information structure	Details for urn:uuid:ba71c65c-0ac2-45fe-89c4-3f7756abe39c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[26]]
214	information structure	Details for urn:uuid:b550f3d4-91d4-4c4a-98a9-e69a99cea773 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[25]]
210	information structure	Details for urn:uuid:abf23bd2-7463-47a5-855a-e3d393f4287c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[23]]
212	information structure	Details for urn:uuid:5b6eb6a5-dad0-4be1-b056-2cbfc14ec406 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[24]]
206	information structure	Details for urn:uuid:0beec652-fe5c-4cdc-ba59-e1d68ab1004d matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[21]]
180	information structure	Details for urn:uuid:63fbea93-7607-4bb4-a72a-7fd57a73d0a5 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[8]]
204	information structure	Details for urn:uuid:2ab95d30-f7b8-42ab-ba07-832e5e7243f8 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[20]]
202	information structure	Details for urn:uuid:1bc984b9-bba9-438e-8390-1d26d54a289a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[19]]
200	information structure	Details for urn:uuid:12677216-0d39-43e9-98a3-28146e96516c matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[18]]
182	information structure	Details for urn:uuid:758b3b6a-b5c6-4988-a27e-8e999d4b99a2 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[9]]
198	information structure	Details for urn:uuid:e68338e0-3b55-453b-bf8e-5a1a62ac370a matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[17]]
196	information structure	Details for urn:uuid:cf131873-be9c-41f4-b70b-dbbc7f82d8ba matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[16]]
194	information structure	Details for urn:uuid:4973f2c8-eb95-4507-bd5a-db8f008f1bea matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[15]]

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208	information structure	Details for urn:uuid:4f8af9e7-4b76-4e3c-bccc-51d171a5aff matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[22]]
192	information structure	Details for urn:uuid:0ef86e5c-fe06-4877-932f-6c29ff48633e matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[14]]
190	information structure	Details for urn:uuid:ed9e5f61-cb96-4bc5-9075-b1f9c1f74f66 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[13]]
188	information structure	Details for urn:uuid:13d48508-9836-4d35-a7b7-b8e693ed1a54 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[12]]
186	information structure	Details for urn:uuid:49c8a0ac-a675-40e2-abfd-4e53e35e63d2 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[11]]
184	information structure	Details for urn:uuid:b5c57a4f-618e-4960-aa41-dbc32d8dec88 matching against profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation 0.2.0 [Bundle.entry[42].resource.result[10]]
133	warning invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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[37].resource/*Observation/null*/]
134	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[37].resource/*Observation/null*/.subject.identifier]
131	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[36].resource/*Observation/null*/.subject.identifier]
130	warning invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[36].resource/*Observation/null*/]
124	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[34].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
128	warning not-found	A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated [Bundle.entry[35].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
127	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[35].resource/*Observation/null*/.subject.identifier]
126	warning invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[35].resource/*Observation/null*/]
136	warning invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[38].resource/*Observation/null*/]
152	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[42].resource/*DiagnosticReport/null*/.subject.identifier]
139	warning invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[39].resource/*Observation/null*/]
140	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 [Bundle.entry[39].resource/*Observation/null*/.subject.identifier]

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141 warning	not-found	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)
143 warning	invalid	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated Best Practice Recommendation: In general, all observations should have an effective[x] ()
144 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)
146 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)
150 warning	not-found	A definition for CodeSystem 'http://hl7.org/fhir/diagnosticreport-relevant-information-types' could not be found, so the code cannot be validated
137 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)
123 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)
120 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)
31 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
32 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)
33 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated
35 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
36 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)
37 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
39 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
40 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)

	severity	code	details
41	warning	not-found	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) A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
43	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
44	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)
45	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
47	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
48	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)
49	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
51	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
52	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)
53	warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
55	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
29	warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' 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)
27	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
2	warning	code-invalid	None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)
4	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
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)

	severity	code	details
7	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
8	warning	code-invalid	[Bundle.entry[3].resource/*Observation/null*/].subject.iden
9	warning	not-found	A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated
11	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
12	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) 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)
56	warning	code-invalid	[Bundle.entry[15].resource/*Observation/null*/].subject.iden
13	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/refseq ' could not be found, so the code cannot be validated
15	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
16	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)
17	warning	not-found	A definition for CodeSystem ' http://www.genenames.org ' could not be found, so the code cannot be validated
19	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
20	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)
21	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
23	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
24	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)
25	warning	not-found	A definition for CodeSystem ' http://www.ncbi.nlm.nih.gov/clinvar ' could not be found, so the code cannot be validated
122	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] ()
57	warning	not-found	A definition for CodeSystem ' http://varnomen.hgvs.org ' could not be found, so the code cannot be validated
59	warning	invalid	[Bundle.entry[16].resource/*Observation/null*/]

severity	code	details
96 warning	not-found	Best Practice Recommendation: In general, all observations should have an effective[x] () A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
98 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
99 warning	code-invalid	[Bundle.entry[27].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
100 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
102 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
103 warning	code-invalid	[Bundle.entry[28].resource/*Observation/null*/.subject.id]
105 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
106 warning	code-invalid	[Bundle.entry[29].resource/*Observation/null*/.subject.id]
107 warning	not-found	A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated
109 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
110 warning	code-invalid	[Bundle.entry[29].resource/*Observation/null*/.subject.id]
111 warning	not-found	A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated
113 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
114 warning	code-invalid	[Bundle.entry[30].resource/*Observation/null*/.subject.id]
116 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
117 warning	code-invalid	[Bundle.entry[31].resource/*Observation/null*/.subject.id]
119 warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () 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)
95 warning	code-invalid	[Bundle.entry[32].resource/*Observation/null*/.subject.id]

	severity	code	details
94	warning	invalid	<p>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>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
92	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>
60	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>
62	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
63	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>
68	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>
73	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
74	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>
76	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
77	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>
80	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>
82	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
83	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>
84	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>
86	warning	invalid	<p>Best Practice Recommendation: In general, all observations should have an effective[x] ()</p>
87	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</p>

	severity	code	details
		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)	
88	warning	not-found	A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated [Bundle.entry[24].resource/*Observation/null*/.value.ofType(CodeableConcept).coding[0].system]
90	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[25].resource/*Observation/null*/]
91	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[25].resource/*Observation/null*/.subject.id]
79	warning	invalid	Best Practice Recommendation: In general, all observations should have an effective[x] () [Bundle.entry[22].resource/*Observation/null*/]