

16, Feb 2026

| severity       | code          | details  |
|----------------|---------------|--|
| 60 error       | structure     | Unable to find a profile match for urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Patient<br>Profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1, Element matches more than one slice - organization, operator  |
| 56 error       | invalid       | Unable to find a profile match for urn:uuid:cdd477e7-9fbc-4391-95c6-8e69a4121ca6 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation   |
| 62 error       | structure     | Slice<br>'DiagnosticReport.code.coding:GenomicTestDirectory': a matching slice is required, but not found (from https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1). Note that other slices are allowed in addition to this required slice  |
| 59 error       | structure     | Unable to find a profile match for urn:uuid:6f11285fc92-433d-b001-36150aa4afe7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant<br>Slicing cannot be evaluated: Could not match discriminator (\$this) for slice  |
| 64 error       | structure     | 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  |
| 55 error       | processing    | Unable to find a profile match for urn:uuid:6f11285fc92-433d-b001-36150aa4afe7 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation   |
| 66 error       | structure     | Unable to find a profile match for urn:uuid:61d28885caa2-4f41-b135-144b39c136d0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication  |
| 68 error       | structure     | Unable to find a profile match for urn:uuid:61d28885caa2-4f41-b135-144b39c136d0 among choices: https://fhir.nwgenomics.nhs.uk/StructureDefinition/Observation  |
| 70 error       | structure     | Not a valid date format: '1982--0-2'   |
| 2 error        | 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> ) |
| 40 information | code-invalid  | 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)<br>Details for urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1                 |
| 57 information | not-supported | Details for urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile http://hl7.org/fhir/StructureDefinition/Patient 4.0.1<br>Entry 'urn:uuid:3756906b-67e9-4522-acf6-85f168535e55' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)   |
| 26 information | structure     | 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  |
| 45 information | structure     | [Bundle.entry[1].resource/*Patient/null*/.birthDate]   |
| 54 information | informational | [Bundle.entry[6].resource/*Observation/null*/.category[1]]   |
| 25 information | code-invalid  | [Bundle.entry[7].resource/*ServiceRequest/null*/.subject]  |
|                |               | [Bundle.entry[10]]   |
|                |               | [Bundle.entry[5].resource/*Observation/null*/.category[1]]   |

| severity | code                      | details  |
|----------|---------------------------|--|
| 18       | information structure     | <p>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> <p>Details for<br/>urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Patient 4.0.1">http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</a></p>   |
| 41       | information structure     | <p>Details for<br/>urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Patient 4.0.1">http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</a></p>  |
| 48       | information structure     | <p>Details for<br/>urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Patient 4.0.1">http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</a></p>  |
| 61       | information structure     | <p>Details for<br/>urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Patient 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Patient 0.2.1</a></p> <p>Details for<br/>urn:uuid:cd477e7-9fb8-4391-95c6-8e69a4121ca6 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></p> |
| 63       | information structure     | <p>Details for urn:uuid:6f11285f-fc92-433d-b001-36150aa4afe7 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/Variant 0.2.1</a></p>  |
| 65       | information structure     | <p>Details for urn:uuid:6f11285f-fc92-433d-b001-36150aa4afe7 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></p>  |
| 67       | information structure     | <p>Details for urn:uuid:6f11285f-fc92-433d-b001-36150aa4afe7 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></p>  |
| 6        | information structure     | <p>Details for<br/>urn:uuid:da8b2d43-99e8-45dd-9e98-4f70b6038000 matching against profile <a href="http://hl7.org/fhir/StructureDefinition/Patient 4.0.1">http://hl7.org/fhir/StructureDefinition/Patient 4.0.1</a></p>  |
| 5        | information code-invalid  | <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>   |
| 69       | information structure     | <p>Details for urn:uuid:61d28885-caa2-4f41-b135-144b39c136d0 matching against profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticImplication 0.2.1</a></p>  |
| 71       | information structure     | <p>Details for urn:uuid:61d28885-caa2-4f41-b135-144b39c136d0 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></p>  |
| 58       | information informational | <p>This element does not match any known slice defined in the profile <a href="https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1">https://fhir.nwgenomics.nhs.uk/StructureDefinition/DiagnosticReport 0.2.1</a> (this may not be a problem, but you should check that it's not intended to match a slice)</p>  |
| 46       | 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>                                     |
| 49       | 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>                                     |
| 36       | warning not-found         |  |

| severity   | code         | details  |
|------------|--------------|--|
|            |              | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/projects/SNP' could not be found, so the code cannot be validated   |
| 42 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> ) |
| 1 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> ) |
| 4 warning  | invalid      | Best Practice Recommendation: In general, all observations should have an effective[x] ()  |
| 7 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> ) |
| 8 warning  | not-found    | A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated   |
| 9 warning  | not-found    | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated   |
| 10 warning | not-found    | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated   |
| 11 warning | not-found    | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated   |
| 12 warning | not-found    | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated   |
| 13 warning | not-found    | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated   |
| 19 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> ) |
| 24 warning | invalid      | Best Practice Recommendation: In general, all observations should have an effective[x] ()  |
| 27 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> ) |
| 28 warning | not-found    | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/clinvar' could not be found, so the code cannot be validated  |
| 29 warning | not-found    | A definition for CodeSystem 'http://www.genenames.org' could not be found, so the code cannot be validated   |
| 30 warning | not-found    | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated   |
| 31 warning | not-found    | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated   |

| <b>severity</b>   | <b>code</b> | <b>details</b>   |
|-------------------|-------------|--|
| <b>32</b> warning | not-found   | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated<br>[Bundle.entry[5].resource/*Observation/null*/.component[4].value.ofType(CodeableConcept).coding]           |
| <b>33</b> warning | invalid     | Coding has no system. A code with no system has no defined meaning, and it cannot be validated. A system should be provided<br>[Bundle.entry[5].resource/*Observation/null*/.component[6].code.coding[0]]                |
| <b>34</b> warning | not-found   | A definition for CodeSystem 'http://www.ncbi.nlm.nih.gov/refseq' could not be found, so the code cannot be validated<br>[Bundle.entry[5].resource/*Observation/null*/.component[7].value.ofType(CodeableConcept).coding] |
| <b>35</b> warning | not-found   | A definition for CodeSystem 'http://varnomen.hgvs.org' could not be found, so the code cannot be validated<br>[Bundle.entry[5].resource/*Observation/null*/.component[8].value.ofType(CodeableConcept).coding]           |
| <b>37</b> warning | not-found   | A definition for CodeSystem 'Chrom-Loc' could not be found, so the code cannot be validated<br>[Bundle.entry[5].resource/*Observation/null*/.component[13].value.ofType(CodeableConcept).coding]                         |
| <b>39</b> warning | invalid     | Best Practice Recommendation: In general, all observations should have an effective[x] ()<br>[Bundle.entry[6].resource/*Observation/null*/]  |
| <b>43</b> warning | not-found   | A definition for CodeSystem 'MedGen-Dis' could not be found, so the code cannot be validated<br>[Bundle.entry[6].resource/*Observation/null*/.component[0].value.ofType(CodeableConcept).coding]                         |