

12, Feb 2026

	severity	code	details	expression
0	error	invalid	Object must have some content	[Bundle.entry[5].resource.presentedForm[0]]
5	error	structure	DocumentReference.content: minimum required = 1, but only found 0 (from http://hl7.org/ fhir/StructureDefinition/ DocumentReference 4.0.1) The extension http://hl7.org/ fhir/StructureDefinition/ workflow-supportingInfo is not allowed to be used at this point (this element is [Bundle.entry[5].resource]; allowed for this version = e:DiagnosticReport, e:Observation, e:CommunicationRequest) This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[2].resource/ *DocumentReference/null*/]
23	error	structure	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource]
22	information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource.resultsInterpreter[0]]
21	information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource.basedOn[0]]
20	information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (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' (http://hl7.org/fhir/ValueSet/ report-codes 4.0.1), and a coding is recommended to come from this value set (codes = https:// fhir.nwgenomics.nhs.uk/ CodeSystem/ IGEAP#CCP19v1_R112, https://fhir.nhs.uk/CodeSystem/ England- GenomicTestDirectory#R112.1) Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/ CodeSystem/ GenomicClinicalIndication 0.1.0	[Bundle.entry[5].resource.extension[0]]
16	information	code-invalid	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource/*DiagnosticReport/ null*/.code]
12	information	business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/ CodeSystem/ GenomicClinicalIndication 0.1.0 This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[4].resource/*ServiceRequest/ null*/.reasonCode[0]]
24	information	informational	This element does not match any known slice defined in the profile https:// fhir.nwgenomics.nhs.uk/ StructureDefinition/ DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)	[Bundle.entry[5].resource.code.coding[0]]

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1	informational	DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice) A definition could not be found for Canonical URL 'https://fhir.nwgenomics.nhs.uk/Endpoint/IGENE' but a definition can be found for 'http://fhir.nwgenomics.nhs.uk/Endpoint/IGENE' - change from https: to http: 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[0].resource/*MessageHeader/null*/.source.endpoint]
3	warning	code-invalid Attachments have data and/or url, or else SHOULD have either contentType and/or language 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[1].resource/*Patient/null*/.identifier[0].type]
18	warning	structure 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[5].resource/*DiagnosticReport/null*/.presentedForm[0]]
17	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[5].resource/*DiagnosticReport/null*/.subject.identifier.type]
8	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[3].resource/*Specimen/null*/.subject.identifier.type]
13	warning	code-invalid Unknown Code 'R112' in the CodeSystem 'https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication' version '0.1.0' - 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)	[Bundle.entry[4].resource/*ServiceRequest/null*/.reasonCode[0].coding[0].code]
6	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[2].resource/*DocumentReference/null*/.subject.identifier.type]

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11 warning	code-invalid	<p>None of the codings provided are in the value set 'IdentifierType' (http://hl7.org/fhir/ValueSet/identifier-type 4.0.1), and a coding should come from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = http://terminology.hl7.org/CodeSystem/v2-0203#NH)</p> <p>Unknown Code 'R112.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory' version '0.1.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>	[Bundle.entry[4].resource/*ServiceRequest/null*/.subject.identifier.type]
10 warning	code-invalid	<p>Unknown Code 'R112.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory' version '0.1.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>	[Bundle.entry[4].resource/*ServiceRequest/null*/.code.coding[0].code]
15 warning	code-invalid	<p>Unknown Code 'R112.1' in the CodeSystem 'https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory' version '0.1.0' - note that the code system is labeled as a fragment, so the code may be valid in some other fragment</p>	[Bundle.entry[5].resource/*DiagnosticReport/null*/.code.coding[1].code]