

severity	code	details	expression
17 error	invalid	<p>Wrong Display Name 'PANEL : R125.1, R125.2 - Thoracic aortic aneurysm and dissection v1.2' for https://fhir.nwgenomics.nhs.uk/CodeSystem/IGEAP#R125_Cardiology.</p> <p>Valid display is 'PANEL: R125.1, R125.2 - Thoracic aortic aneurysm and dissection v1.2' (for the language(s) 'en')</p> <p>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#R125_Cardiology, https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R125.1)</p> <p>Entry 'urn:uuid:b3518c03-9e70-45bd-a44d-fc59b8903625' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p> <p>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)</p> <p>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)</p> <p>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)</p> <p>Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0</p>	[Bundle.entry[6].resource/*DiagnosticReport/null*/.code.coding[0].display]
18 information	code-invalid	<p>code = https://fhir.nwgenomics.nhs.uk/CodeSystem/IGEAP#R125_Cardiology, https://fhir.nhs.uk/CodeSystem/England-GenomicTestDirectory#R125.1)</p> <p>Entry 'urn:uuid:b3518c03-9e70-45bd-a44d-fc59b8903625' isn't reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p> <p>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)</p> <p>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)</p> <p>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)</p> <p>Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0</p>	[Bundle.entry[6].resource/*DiagnosticReport/null*/.code]
24 information	informational	<p>informational</p> <p>reachable by traversing forwards from the MessageHeader. Check that this is meant to be included (needed to process the message)</p> <p>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)</p> <p>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)</p> <p>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)</p> <p>Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0</p>	[Bundle.entry[8]]
26 information	informational	<p>informational</p> <p>StructureDefinition/DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p>	[Bundle.entry[6].resource.resultsInterpreter[0]]
27 information	informational	<p>informational</p> <p>StructureDefinition/DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>StructureDefinition/DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>StructureDefinition/DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p>	[Bundle.entry[6].resource.code.coding[0]]
25 information	informational	<p>informational</p> <p>StructureDefinition/DiagnosticReport 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)</p> <p>Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0</p>	[Bundle.entry[6].resource.basedOn[0]]
14 information	business-rule	<p>business-rule</p> <p>CodeSystem/GenomicClinicalIndication 0.1.0</p>	[Bundle.entry[5].resource/*ServiceRequest/null*/.reasonCode[0]]
19 warning	code-invalid	<p>code-invalid</p> <p>None of the codings provided are in the value set</p>	[Bundle.entry[6].resource/*DiagnosticReport/null*/.subject.identifier.type]

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15 warning	code-invalid	'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) Unknown Code 'R125' 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[5].resource/*ServiceRequest/null*/.reasonCode[0].coding[0].code]
13 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/*ServiceRequest/null*/.subject.identifier.type]
11 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[4].resource/*Specimen/null*/.subject.identifier.type]
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[3].resource/*DocumentReference/null*/.subject.identifier.type]
1 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[1].resource/*Patient/null*/.identifier[0].type]