

severity	code	
31 error	value	The pattern [system http://terminology.hl7.org/CodeSystem/v2-0203 , code AN, and display 'Account number'] defined in fhir.nwgenomics.nhs.uk/StructureDefinition/HospitalProviderSpellIdentifier 0.1.0 not found. Issues: [ValidationMessage[level=ERROR,type=VALUE,location=Bundle.entry[7].resource.encounter.identifier.type.coding='VN' but is fixed to 'AN' in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/HospitalProviderSpellIdentifier 0.1.0]] ValidationMessage[level=ERROR,type=VALUE,location=Bundle.entry[7].resource.encounter.identifier.type.coding='VN' but is fixed to 'AN' in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/HospitalProviderSpellIdentifier 0.1.0]]
29 error	structure	The extension http://hl7.org/fhir/StructureDefinition/workflow-supportingInfo is not allowed to be used at this point [Bundle.entry.resource]; allowed for this version = e:DiagnosticReport, e:Observation, e:CommunicationRequest)
30 information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicClinicalIndication 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)
28 information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicTest 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)
27 information	informational	This element does not match any known slice defined in the profile https://fhir.nwgenomics.nhs.uk/StructureDefinition/GenomicTest 0.1.0 (this may not be a problem, but you should check that it's not intended to match a slice)
18 information	business-rule	Reference to draft CodeSystem https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication 0.1.0
21 information	code-invalid	None of the codings provided are in the value set 'LOINC Diagnostic Report Codes' (http://hl7.org/fhir/ValueSet/reports). Coding is recommended to come from this value set (codes = https://fhir.nwgenomics.nhs.uk/CodeSystem/IGEAP#R125.1)
22 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 from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = https://fhir.nwgenomics.nhs.uk/CodeSystem/v2-0203#NH)
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 from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = https://fhir.nwgenomics.nhs.uk/CodeSystem/v2-0203#NH)
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 from this value set unless it has no suitable code (note that the validator cannot judge what is suitable) (codes = https://fhir.nwgenomics.nhs.uk/CodeSystem/v2-0203#VN)
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19 warning	code-invalid	Unknown Code 'R125' in the CodeSystem ' https://fhir.nwgenomics.nhs.uk/CodeSystem/GenomicClinicalIndication ' - the code system is labeled as a fragment, so the code may be valid in some other fragment