

Bindings to FHIR Clinical Resources

This set of topics relates to the use of SNOMED CT to represent the values, meaning and/or maps to FHIR healthcare resources. See <http://build.fhir.org/resourcelist.html> for a list of the relevant resources.

Proposed collaborative work topics:

1. Principles and guidance for profiling/binding with SNOMED CT
2. Review existing base resource bindings to SNOMED CT
3. Sharing/collaboration on profiles, value set definitions, bindings and concept maps
4. Create SNOMED CT resource profiles that bind elements to SNOMED CT wherever that isn't already in the base resources - note this may require more than one profile per resource with different bindings/constraints for different modelling style

Perhaps a page per resource being worked on below this page would be useful, created as they begin to be worked on. Subpages currently include:

- [ConceptMaps for FHIR code systems to SNOMED CT](#)
- [Binding patterns](#)
- [Implementation Guidance Principles](#)
- [Resource bindings](#)
 - [AllergyIntolerance Binding](#)
 - [CarePlan](#)
 - [Condition Resource](#)
 - [DiagnosticReport](#)
 - [Diagnostic Service Section Codes](#)
 - [Immunization](#)
 - [Medication Resource](#)
 - [Mapping of FHIR Medication to SNOMED CT Concept](#)
 - [Observation binding](#)
 - [Procedure](#)
 - [ServiceRequest](#)
 - [Specimen binding](#)
 - [Vital Signs panel binding](#)
- [Molecular Pathology harmonising V2 and FHIR Datasets](#)
- [Developing and publishing ValueSets](#)
- [Free SNOMED CT set for FHIR](#)
 - [20190601 - Response on free set content addition by Jim Case](#)
 - [FHIR SNOMED CT Free Set - Consider for inclusion / rejection \(Red mappings\)](#)
 - [FHIR SNOMED CT Free Set - Proposal for Grahame Grieve \(Green mappings\)](#)
 - [FHIR SNOMED CT Free Set - Questions for HL7 \(Amber mappings\)](#)
- [COVID-19 Analysis](#)
 - [COVID 19 Diagnosis](#)
 - [COVID 19 General Comorbidities Absent](#)
 - [COVID 19 General Comorbidities Present](#)
 - [COVID-19 Symptoms Absent](#)
 - [COVID-19 Symptoms Present](#)
 - [Human coronavirus 229E+HKU1+OC43+NL63 RNA \[Presence\] in Nasopharynx by NAA with non-probe detection](#)
 - [Known Exposure](#)
 - [Reporting Priority](#)
 - [Travel History](#)
- [Flavours of Null and Degrees of Certainty](#)