

# Creating Terminology Server for FHIR using OMOP CDM

OMOP on FHIR Project  
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# What is Terminology Server?

a piece of software providing a range of terminology-related software services through an applications programming interface to its client applications.

- Wikipedia ([https://en.wikipedia.org/wiki/Terminology\\_server](https://en.wikipedia.org/wiki/Terminology_server))

# Why do we need a Terminology Server?

- We have needs for terminology.
  - Interoperability using same vocabulary
  - Easy of getting meanings from concept sets
  - And more...
- Terminology Server enables the programmatic access of terminology content
- With standard terminology services, the access technology can be reused
- Updates on terminology can be available to all client applications

ValueSet, ConceptMap, CodeSystem

**FHIR** RestAPI

*Terminology* **API**

**OMOP** CDM

*Concept/Vocabulary* **Database**

concept, concept\_relationship,  
concept\_ancestor, relationship, vocabulary

# Terminology in FHIR

- **CodeSystem**: a set of codes with meanings.
- **ValueSet**: a set of codes from those defined by one or more code systems to specify which codes can be used in a particular context.
- **ConceptMap**: a mapping from a set of concepts defined in a code system to one or more concepts defined in other code systems.
- **ExpansionProfile**: used to configure the behavior of the terminology server for expansions and validations.
- **NamingSystem**: a curated namespace that issues unique symbols within that namespace for the identification of concepts, people, devices, etc.

# Concept/Vocabulary in OMOP

- **concept**: a record that represents clinical information in all domains.
- **concept\_relationship**: defines direct relationships between concepts
- **concept\_ancestor**: hierarchical relationships between concepts
- **relationship**: defines type of a relationship.
- **concept\_class**: list of classifications used to differentiate concepts.
- **vocabulary**: list of vocabularies used in concept.

## ValueSet in FHIR

Concept and concept\_ancestor in OMOP CDM can be used to create ValueSet resources.

Example: Cholesterol | Bld-Ser-Plas (LOINC Codes for Cholesterol in Serum/Plasma)

- Find concepts with parent = LP43571-6 from concept\_ancestor.
- Add the concepts to expansion of valueset resource
- Example: <https://www.hl7.org/fhir/valueset-example-expansion.json.html>

## ConceptMap in FHIR

concept\_relationship in OMOP CDM can be used to create ConceptMap resources

Example: translate one concept in a domain to another concept in a different domain

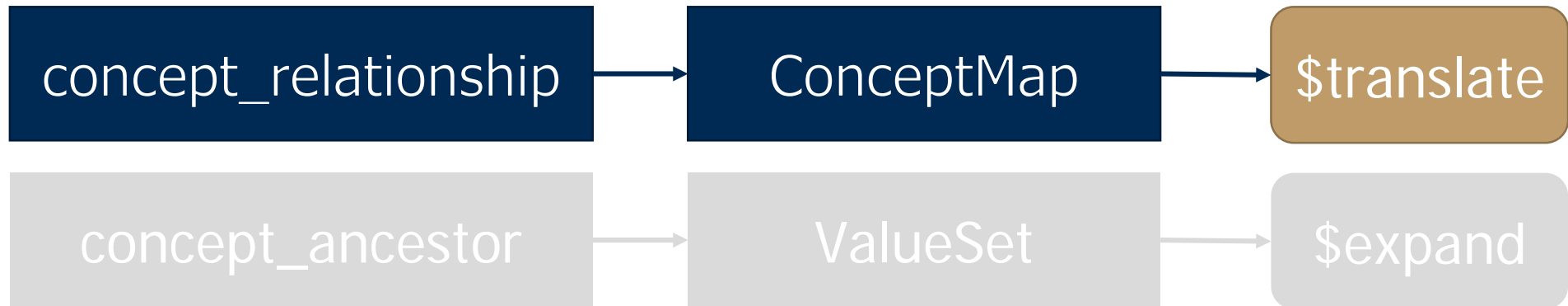
- Use relationship ID to define equivalence
- Return target coding system

FHIR equivalence:

relatedto | equivalent | equal | wider | subsumes | narrower | specializes | inexact |  
unmatched | disjoint

Need to find matching relationship from relationship table in OMOP.

# Current implementation of Terminology Server



- \$translate operation
- Local code system import for mapping to standard

- How it works:

[http://localhost:8080/fhir/ConceptMap/\\$translate?system=http://www.nlm.nih.gov/research/umls/rxnorm&code=311040&targetsystem=http://hl7.org/fhir/ndfirt](http://localhost:8080/fhir/ConceptMap/$translate?system=http://www.nlm.nih.gov/research/umls/rxnorm&code=311040&targetsystem=http://hl7.org/fhir/ndfirt)

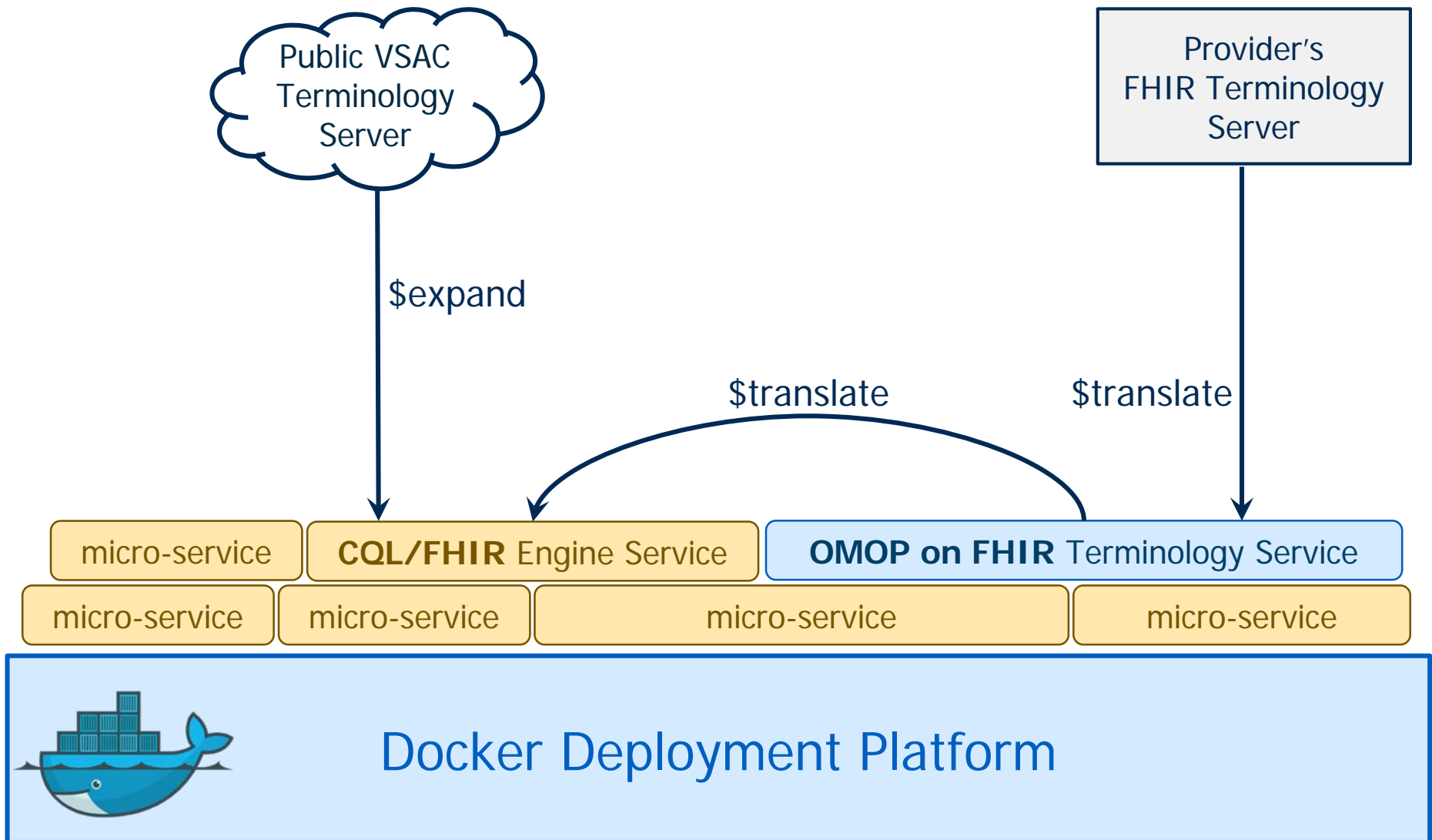
- Source:  
RxNorm | 311040 | Insulin, Aspart, Human 100 UNT/ML Injectable Solution
- Target:  
NDFRT | ??



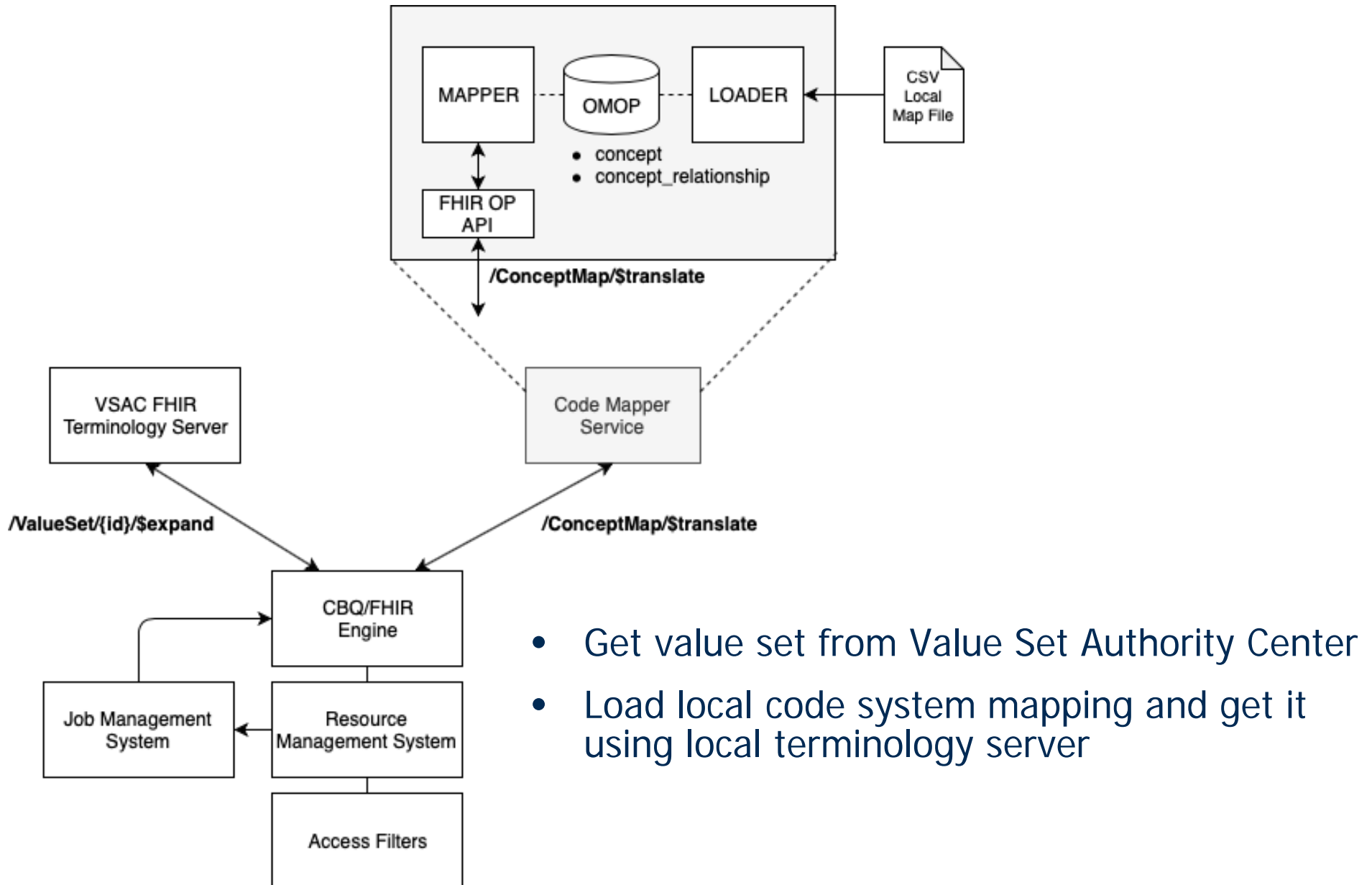
```
{
  "resourceType": "Parameters",
  "parameter": [
    {
      "name": "result",
      "valueBoolean": true
    },
    {
      "name": "match",
      "part": [
        {
          "name": "equivalence",
          "valueCode": "equivalent"
        },
        {
          "name": "concept",
          "valueCoding": {
            "system": "http://hl7.org/fhir/ndfrt",
            "code": "4014955",
            "display": "INSULIN,ASPART,HUMAN 100 UNT/ML INJ"
          }
        }
      ]
    }
  ]
}
```

# Local code system to standard code system mapping CSV file

```
# Coding map document.
# 1st line: defines mapping. Must be OMOP vocabularies (eg. RxNorm). Comma separated. 1st value is
source, 2nd is target.
# 2nd line: defines same mapping but with FHIR coding system name.
# If vocabulary does not exist, it will be created in the database.
# 3rd line: Definition of column. There are three required columns (if standard coding, description column is
optional).
# SOURCE_CODESYSTEM, SOURCE_CODE, SOURCE_DESC, TARGET_CODE
# Other columns will be ignored.
MUSCLabOb^MUSC Lab Observation,LOINC
urn:hssc:srhs:la01:observation:code,http://loinc.org
SOURCE_CODESYSTEM,SOURCE_CODE,SOURCE_DESC,TARGET_CODE,LOINC Code Description,
urn:hssc:srhs:la01:observation:code,789,"% ALBUMIN, SERUM",13980-8,Albumin/Protein.total in Serum or Plasma
by Electrophoresis,
urn:hssc:srhs:la01:observation:code,796,"% ALBUMIN, URINE",13992-3,Albumin/Protein.total in Urine by
Electrophoresis,
urn:hssc:srhs:la01:observation:code,790,"% ALPHA 1, SERUM",13978-2,Alpha 1 globulin/Protein.total in Serum or
Plasma by Electrophoresis,
```



# Bit more detail between the services



# Questions ??

<http://omoponfhir.org/> for more about OMOP on FHIR