



SNOMED CT, FHIR, and more

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Phast-Services: manager of international projects

IHE International: co-chair of Pathology and Laboratory Medicine (**PaLM**) committee

Interop'Santé: chair of **HL7 France**

Where do I speak from?

Two tightly bound organizations



- Not-for-profit association
- Members: healthcare IT vendors
- Contributes to healthcare interoperability standards (HL7, IHE, SNOMED CT, LOINC ...)
- Current projects: HL7 IPS, Trillium II, Catalogs on FHIR, ...



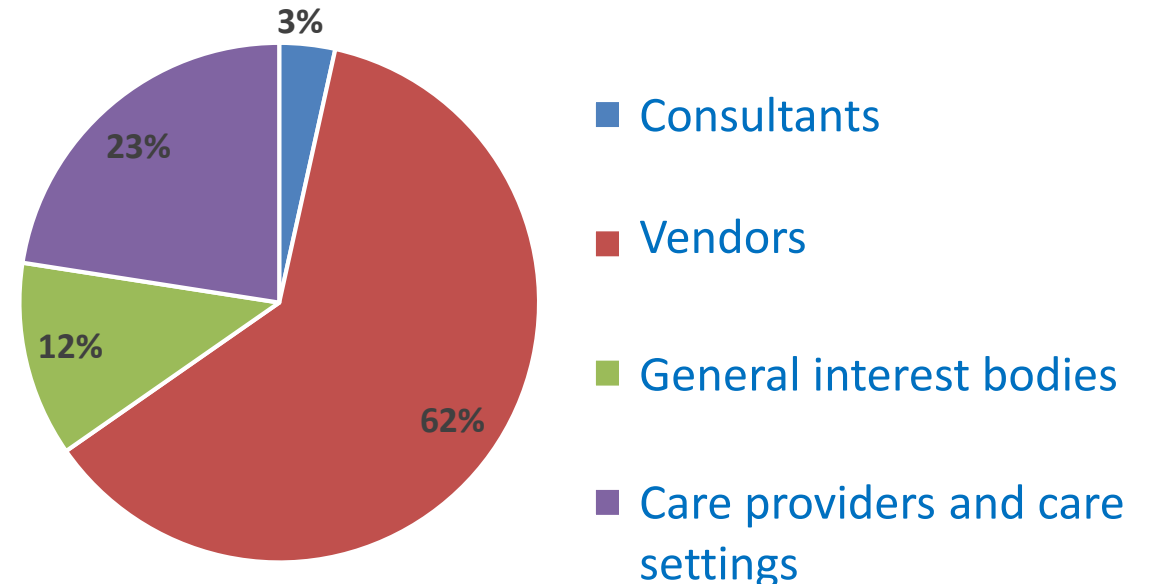
- Company: production structure
- Customers: hospitals & clinics
- Builds and markets services deploying the standards in secondary care settings.
- Current services: terminology services, metathesauri, expertise

Interop'Santé non-profit association

One goal: build standardized interoperability for health IT systems



160 members



The *Pathology and Laboratory Medicine (PaLM)* domain covers interoperability needs of clinical and pathology laboratory workflows, transfusion medicine workflows and biobanking.

This domain has 3 sponsors :



<http://www.cap.org>

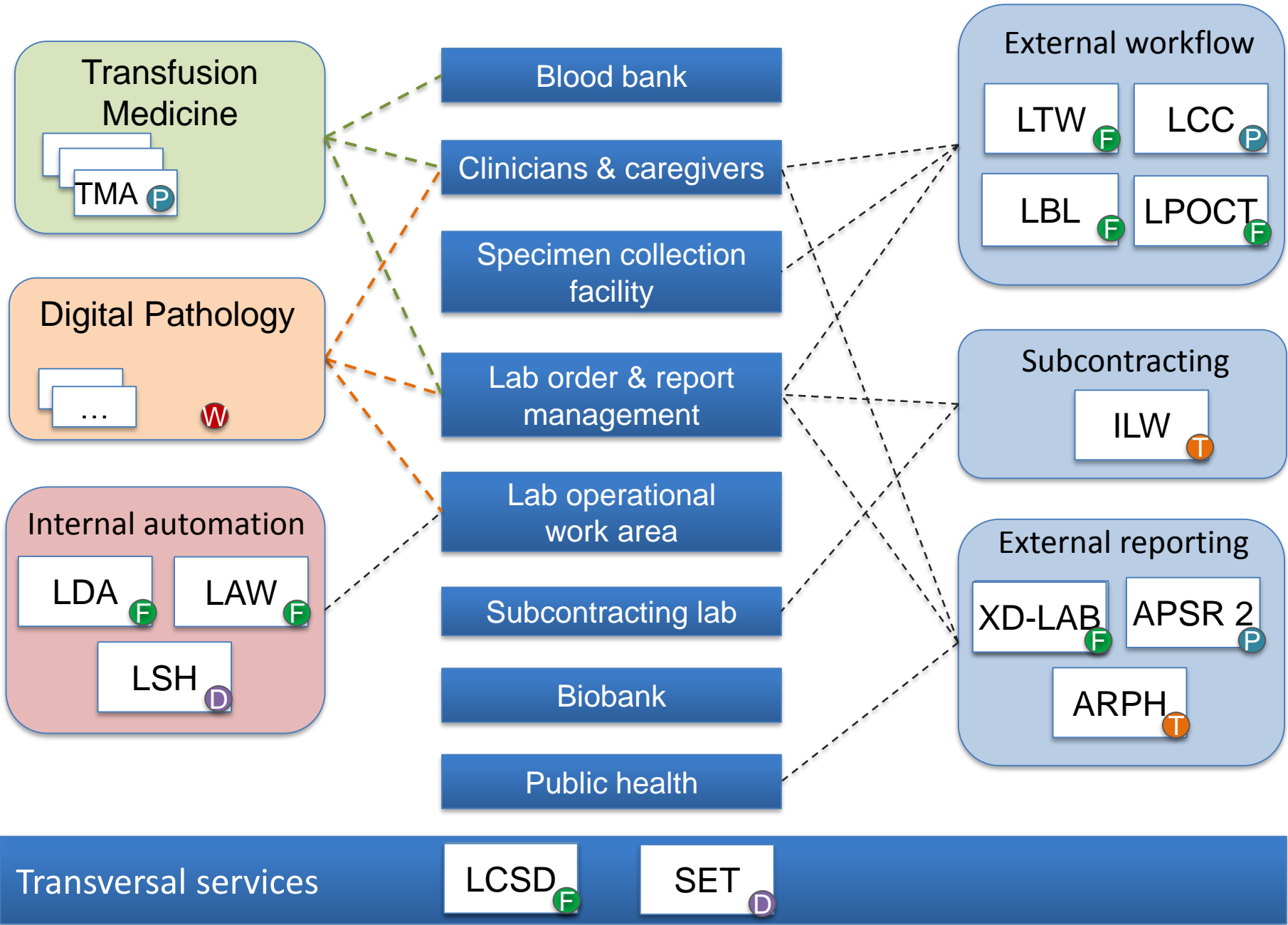
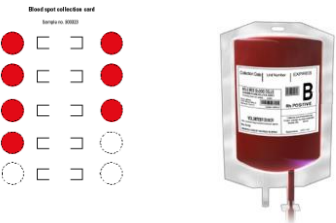
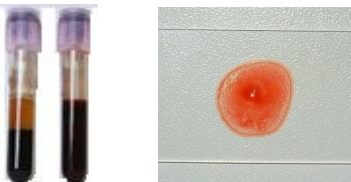
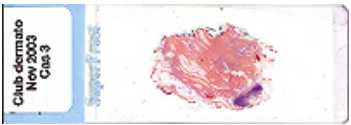


<https://www.jahis.jp/>



<http://phast.fr/>

IHE PaLM domain



- ^W White paper
- ^D Development
- ^P Public comment
- ^T Trial Implementation
- ^F Final Text



Agenda:

1. Meaningful data in health IT systems: why and how?
2. Adoption of SNOMED CT needs to be stepwise
3. Leveraging external FHIR terminology services
4. Leveraging FHIR catalogs



Agenda:

1. Meaningful data in health IT systems: why and how?
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3. Leveraging external FHIR terminology services
4. Leveraging FHIR catalogs

An example



Laboratory procedure requested:

- Microbiology study on nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication of the patient: none



Staphylococcus aureus identified.
vancomycin MIC ≥ 32 $\mu\text{g/ml}$ -> Resistant



Laboratory procedure requested:

- Microbiology study of nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication of the patient: none



The ordering physician and the clinical lab director have a common understanding of this statement because:

- a) they speak the same language (syntax),
- b) they use a common vocabulary (vernacular + medicine)
- c) both of them acquired full knowledge of these concepts by academic education.



This common understanding enables them to cooperate (interoperate) efficiently on this procedure.



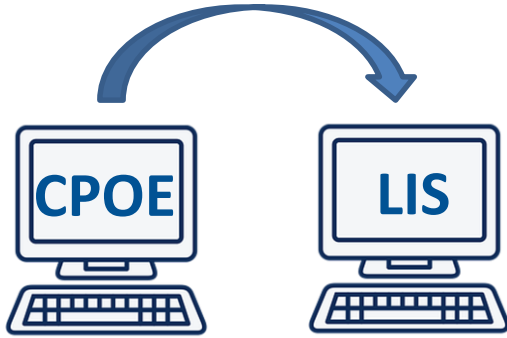
Laboratory procedure requested:

- Microbiology study of nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication: none

To cooperate even more efficiently, they need their IT systems to interoperate too, and to share this common understanding. This for more than one reason:

1. assist the user at data entry (creation of the procedure request),
2. integrate the request and acknowledge the event,
3. store each data element into the proper place,
4. display meaningfully the content of this request combined with relevant information available locally,
5. trigger actions to automate the process of the request,
6. later on, analyze the activity, or the population of patients based on the outcome of this activity,
7. ...





Laboratory procedure requested:

- Microbiology study of nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication: none

This common understanding between systems is only achieved if:

- a) The systems use a common language/syntax (**information models + concept binding**).
- b) The systems use a common vocabulary (**reference terminologies**).
- c) The systems have acquired some level of education (**parameterization**) about these concepts and models, to be able to operate on them (check consistency, trigger action, ...)
- d) The systems are able to retrieve knowledge about the concept received or to be keyed in (**terminology services**).



Common vocabulary:

What terminologies are needed and which concepts from each?



Laboratory procedure requested:

Microbiology study of nasal fluid specimen

Reason for ordering: preoperative diagnosis

Current medication: none

SNOMED CT: |Bacteriology general procedure|

SNOMED CT: |Laboratory procedure|

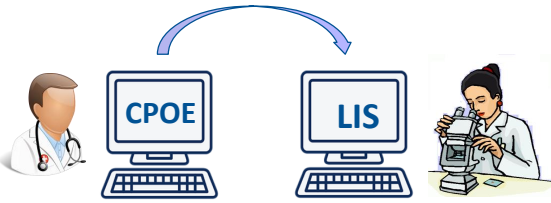
LOINC: |Reason for test or procedure|

LOINC: |Current medication|

SNOMED CT: |none used|

SNOMED CT: |Nasal fluid sample|

SNOMED CT: |Preoperative diagnosis|



Laboratory procedure requested:

- Microbiology study of nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication: none

Common language:

Using HL7 v2 language constrained by IHE LTW profile:

Syntax/language
SNOMED CT
LOINC

```
MSH|^~\&| ... |OML^O33^OML_O33| ... |EN|...  
PID| ... |EVERYMAN^ADAM^^JR^^^L|19800101|M|...  
PV1|...  
SPM|1|123^Card||168141000^Nasal fluid sample^SCT|...  
ORC|...  
TQ1|...  
OBR|1|123^Card||275721003^Bacteriology general procedure^SCT|...  
OBX|1|CE|67098-4^ Reason for test or procedure^LN||406520001^  
Preoperative diagnosis^SCT|...  
OBX|2|CE|18606-4^Current medication^LN||""|
```



Laboratory procedure requested:

- Microbiology study of nasal fluid specimen
- Reason for ordering: preoperative diagnosis
- Current medication: none

Another possible
common language:



Using HL7  FHIR[®] (as in the April R4 ballot):

ServiceRequest

category:

coding:

system: <http://snomed.info/sct>

code: 108252007

display: laboratory procedure

orderDetail:

coding:

system: <http://snomed.info/sct>

code: 275721003

display: bacteriology general procedure

reasonCode:

coding:

system: <http://snomed.info/sct>

code: 406520001

display: preoperative diagnosis

Patient

subject

specimen

supportingInfo

Specimen

type:

coding:

system: <http://snomed.info/sct>

code: 168141000

display: nasal fluid sample

Observation

code:

coding:

system: <http://loinc.org>

code: 18606-4

display: current medication

valueCodeableConcept:

coding:

system: <http://snomed.info/sct>

code: 262001002

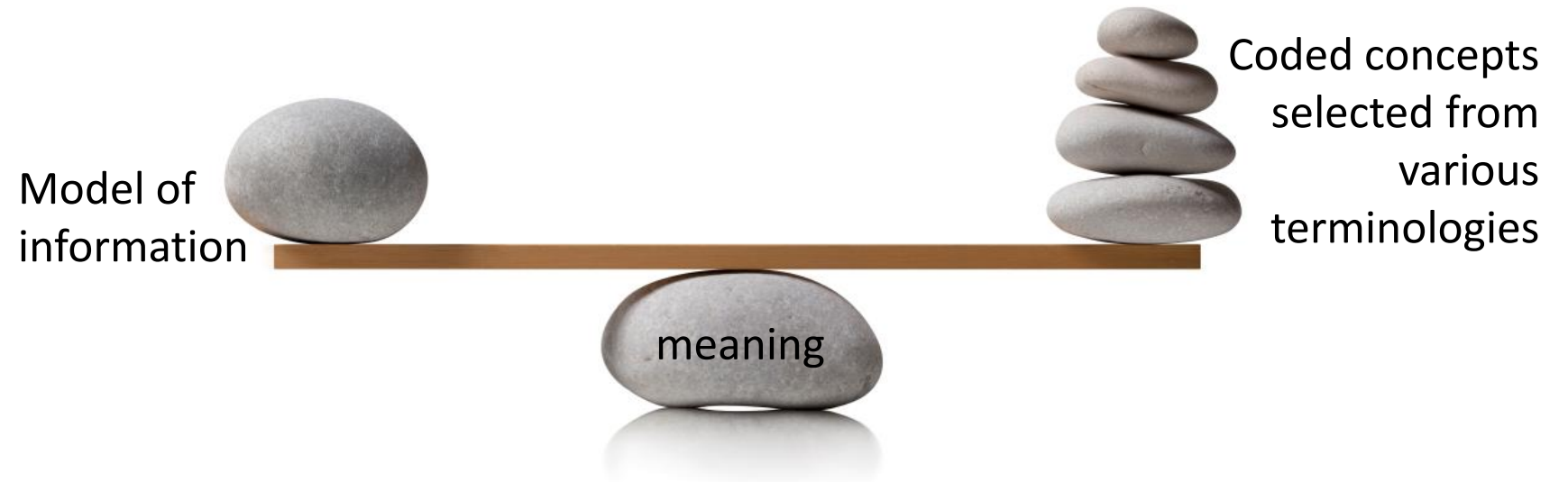
display: none used



The selection of concepts depends upon the chosen information model

- In the FHIR standard the concept of “reason for procedure” is built in the information structure:
`ServiceRequest.reasonCode: 406520001 |Preoperative diagnosis (qualifier value)|`
- In HL7 v2 a multipurpose OBX segment is used instead, therefore an additional concept is needed to specify which kind of observation is that:
`OBX-3: 67098-4^ Reason for test or procedure^LN`
`OBX-5: 406520001^ Preoperative diagnosis^SCT`

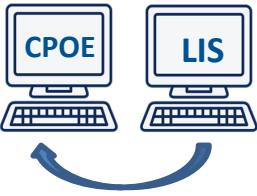
The information model carries a part of the meaning by itself.



Let's take a look at the results flow, now.



Staphylococcus aureus identified.
vancomycin MIC ≥ 32 $\mu\text{g/ml}$ -> Resistant



Common vocabulary



What terminologies are needed and which concepts from each?

SNOMED CT: |Staphylococcus aureus|

Bacteria identified: Staphylococcus aureus.

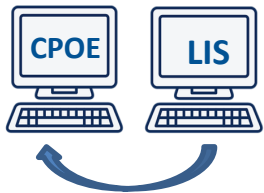
vancomycin MIC \geq 32 $\mu\text{g/ml}$ -> Resistant

HL7 table 0078: |Resistant|

UCUM: |ug/mL|

LOINC: |Vancomycin [Susceptibility] on isolate|

LOINC: |Bacteria identified in Nasopharynx by Culture|

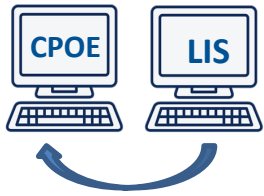


Staphylococcus aureus identified.
vancomycin MIC ≥ 32 $\mu\text{g/ml}$ -> Resistant



Using HL7 v2 language constrained by IHE LTW profile:

```
MSH|^~\&| ... |OUL^R22^OUL_R22| ... |EN|...
PID| ... |EVERYMAN^ADAM^^JR^^^L|19800101|M|...
:
SPM|1|123^Card||168141000^Nasal fluid sample^SCT|...
OBR|1|123^Card||275721003^Bacteriology general procedure^SCT|...
:
OBX|n|CE|67098-4^Bacteria identified in Nasopharynx by Culture^LN||3092008^
Staphylococcus aureus^SCT|... ← child order
:
OBR|2|123^Card||14788002^Antimicrobial susceptibility test^SCT|...
OBX|1|SN|19000-9^Vancomycin susceptibility on isolate^LN|1|>=^32|ug/mL||R|...
```



Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant

Other available language:
CDA R2 constrained by IHE
XD-LAB profile (focusing on
the entry)

entry (act)

procedure

Specimen collection, (specimen is a participant with typeCode="PRD" and a role of "SPEC")

Organizer (battery)

Culture

observation

Bacteria identified

Organizer (cluster)

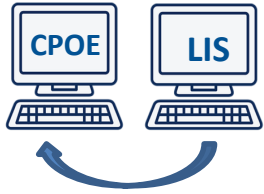
The antibiogram ...

specimen

... is performed on the isolate represented by the name of the bacteria identified

observation

Vancomycin MIC and its clinical interpretation



Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant

Using CDA R2 constrained
by IHE XD-LAB profile

entry (act)

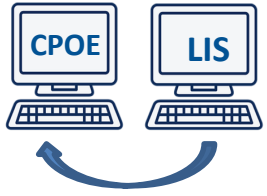
```
<entry typeCode="DRIV">  
  <templateId root="1.3.6.1.4.1.19376.1.3.1"/>  
  <act classCode="ACT" moodCode="EVN">  
    <code code="275721003" codeSystem="2.16.840.1.113883.6.96"  
      displayName="Bacteriology general procedure"/>  
  </act>  
</entry>
```

The entry

procedure

```
<entryRelationship typeCode="COMP">  
  <procedure classCode="PROC" moodCode="EVN">  
    <templateId root="1.3.6.1.4.1.19376.1.3.1.2"/>  
    ...  
    <participant typeCode="PRD">  
      <participantRole classCode="SPEC">  
        <playingEntity>  
          <code code="168141000" codeSystem="2.16.840.1.113883.6.96"  
            codeSystemName="SNOMED-CT" displayName="Nasal fluid sample"/>  
        </playingEntity>  
      </participantRole>  
    </participant>  
  </procedure>  
</entryRelationship>
```

The specimen collected



Staphylococcus aureus identified.
vancomycin MIC ≥ 32 $\mu\text{g/ml}$ -> Resistant

Using CDA R2 constrained
by IHE XD-LAB profile

entry (act)

procedure

...

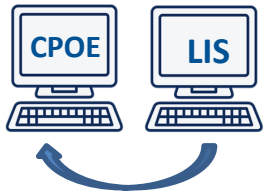
Organizer (battery)

...

observation

```
<observation classCode="OBS" moodCode="EVN">
  <templateId root="1.3.6.1.4.1.19376.1.3.1.6"/>
  <code code="67098-4" codeSystem="2.16.840.1.113883.6.1" codeSystemName="LOINC"
    displayName="Bacteria identified in Nasopharynx by Culture"/>
  <statusCode code="completed"/>
  <effectiveTime value="20150104131933+0100"/>
  <value xsi:type="CD" code="3092008" displayName="Staphylococcus aureus"
    codeSystem="2.16.840.1.113883.6.96" codeSystemName="SNOMED CT"/>
</observation>
```

The bacteria identified



Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant

Using CDA R2 constrained
by IHE XD-LAB profile

entry (act)

⋮

Organizer (cluster)

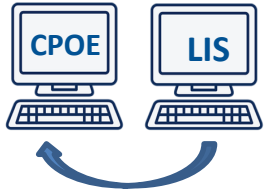
specimen

```
<entryRelationship typeCode="COMP">  
  <organizer classCode="CLUSTER" moodCode="EVN">  
    <templateId root="1.3.6.1.4.1.19376.1.3.1.5"/>  
    <statusCode code="completed"/>  
    <effectiveTime value="20150104155000+0100"/>
```

```
<specimen typeCode="SPC">  
  <specimenRole classCode="SPEC">  
    <id extension="55584739" root="1.3.6.1.4.1.19376.1.3.4"/>  
    <specimenPlayingEntity classCode="MIC">  
      <code code="3092008" displayName="Staphylococcus aureus"  
        codeSystem="2.16.840.1.113883.6.96" codeSystemName="SNOMED CT"/>
```

The antibiogram ...

... is performed on
the isolate
represented by the
name of the
bacteria identified



Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant

Using CDA R2 constrained
by IHE XD-LAB profile

entry (act)

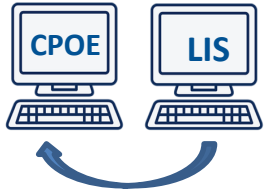
Organizer (cluster)

specimen

observation

```
<observation classCode="OBS" moodCode="EVN">
  <templateId root="1.3.6.1.4.1.19376.1.3.1.6"/>
  <code code="19000-9" codeSystem="2.16.840.1.113883.6.1" codeSystemName="LOINC"
    displayName="Vancomycin susceptibility on isolate"/>
  <statusCode code="completed"/>
  <effectiveTime value="20150104155000+0100"/>
  <value xsi:type="IVL_PQ">
    <low value="32" unit="ug/mL" inclusive="true"/>
  </value>
  <interpretationCode code="R" displayName="Resistant" codeSystem="2.16.840.1.113883.5.83"/>
</observation>
```

The vancomycin MIC



Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant

One more language

Using HL7  FHIR[®] (as in the April R4 ballot):

DiagnosticReport

status: final

category:

coding:

system: <http://hl7.org/fhir/v2/0074>

code: MCB

display: Mycobacteriology

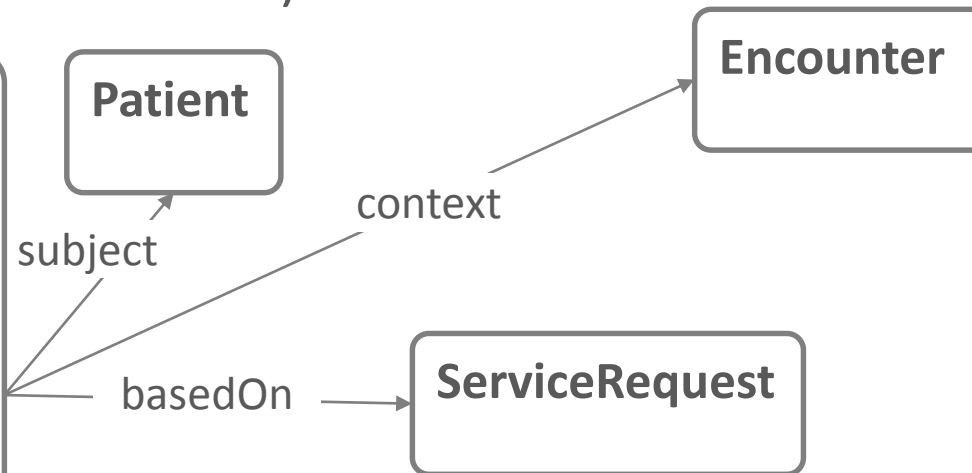
code:

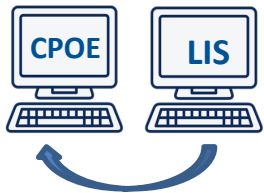
coding:

system: <http://loinc.org>

code: 67098-4

display: bacteria identified in nasopharynx





Staphylococcus aureus identified.
vancomycin MIC $\geq 32 \mu\text{g/ml}$ -> Resistant



Using HL7  FHIR[®]

DiagnosticReport

status: final

...

code:

coding:

system: <http://loinc.org>

code: 67098-4

display: bacteria

result

Specimen

type:

coding:

system: <http://snomed.info/sct>

code: 168141000

display: nasal fluid sample

specimen

Observation

code:

coding:

system: <http://loinc.org>

code: 67098-4

display: bacteria identified in
nasopharynx

valueCodeableConcept:

coding:

system: <http://snomed.info/sct>

code: 3092008

display: Staphylococcus aureus

hasMember

Observation

code:

coding:

system: <http://loinc.org>

code: 19000-9

display: Vancomycin susceptibility on
isolate

valueQuantity:

value: 32

comparator: >=

unit: $\mu\text{g/ml}$

system: <http://unitsofmeasure.org>

code: ug/mL

interpretation:

coding:

system: <http://hl7.org/fhir/v2/0078>

code: R

display: Resistant

Need to alert the clinician about this vancomycin-resistant Staph aureus carrier



1. Use a more specific concept for the observation “Bacteria identified”

➤ SNOMED CT: | Vancomycin resistant Staphylococcus aureus (organism) |

2. Provide an additional observation:

➤ Observation

▪ code: LOINC: | Pathologist interpretation of Unspecified specimen tests |

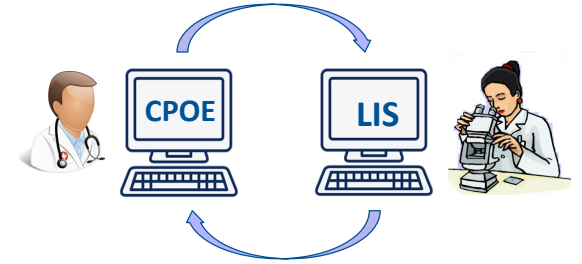
▪ value: SNOMED CT: | Multiple antimicrobial drug resistant bacteria (organism) |

3. In all cases, a conclusion is expected

➤ Example with FHIR: **DiagnosticReport.conclusion:**

“Nasal carriage of multi-drug resistant Staphylococcus aureus”

One more reason for adopting common reference terminologies



1. assist the user at data entry (creation of the procedure request),
2. integrate the request and acknowledge the event,
3. store each data element into the proper place,
4. display meaningfully the content of this request combined with relevant information available locally,
5. trigger actions to automate the process of the request,
6. later on, analyze the activity, or the population of patients based on the outcome of this activity,
7. Enhance collaborative surveillance of healthcare acquired infections



Agenda:

1. Meaningful data in health IT systems: why and how?
2. Adoption of SNOMED CT needs to be stepwise
3. Leveraging external FHIR terminology services
4. Leveraging FHIR catalogs

SNOMED CT provides not only concepts but also rich relationships

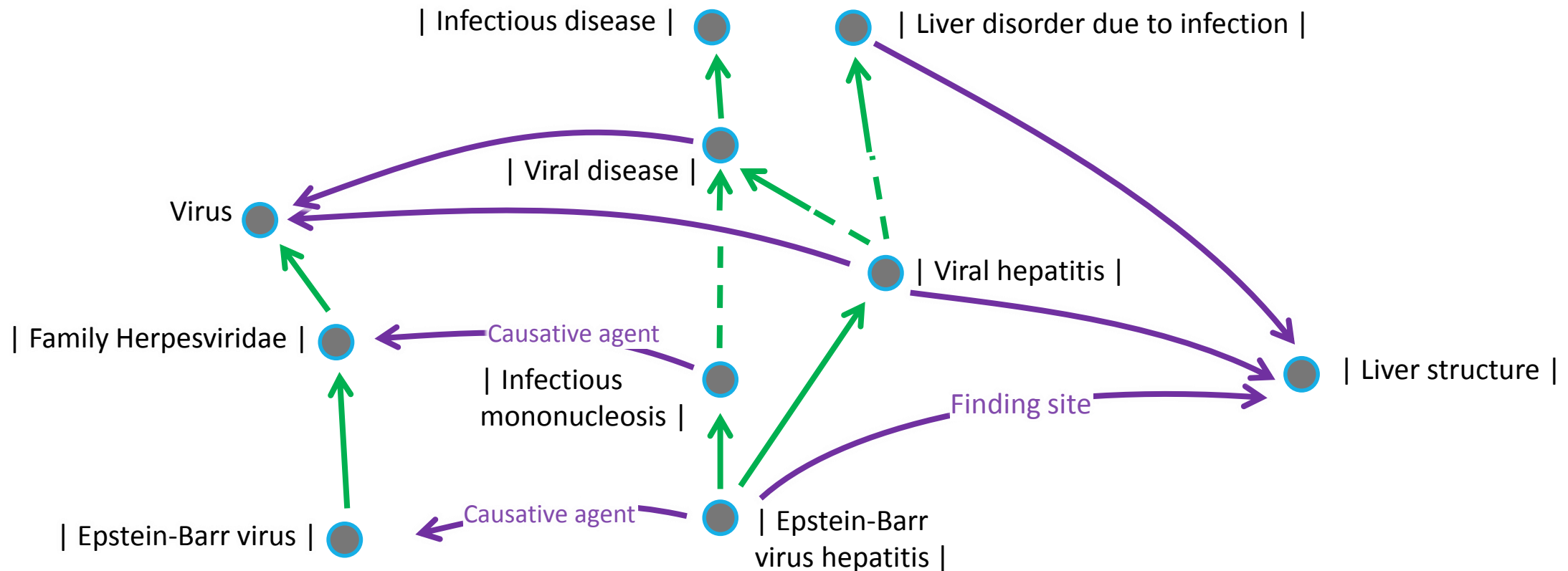


- Polyhierarchical (« *is a* » ↗)
- Multiple attributes (↶)

Microorganisms

Disorders

Anatomical structure



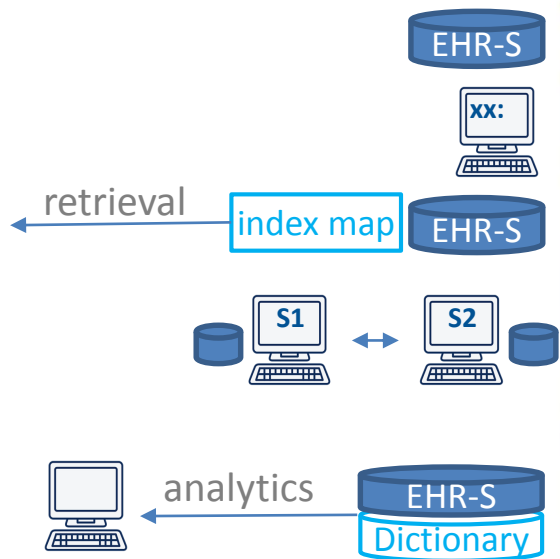
This enables various

Courtesy of
(Implementation course)



Implementation Approaches

SNOMED CT can be used as ...



What	Why
A code system	To store clinical information
An interface terminology	To capture and display clinical information
An indexing system	To retrieve clinical information
A common terminology	To communicate in a meaningful way
	To integrate heterogeneous data
A dictionary	To query, analyze and report
	To link health records to knowledge resources
Extensible foundation	To represent new types of clinical data

- Use of:
- concept model
 - SNOMED CT expressions

Need to start with the best combination of
(high priority need, low hanging fruit, low cost)



My guess for European countries: start using SNOMED CT as a common terminology to improve interoperability



➤ A small step ...

- IT Systems can rely on simple value sets bound to standardized interoperable information models (FHIR, CDA, HL7 v2, ...)

➤ ... which fulfills important needs,

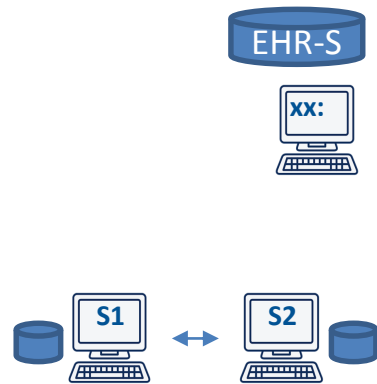
- National EHR/PHR systems
- Digitalized CPOE/result reporting in hospitals
- Cross-border healthcare (eHealth DSI, IPS, Trillium II, ...)

➤ ... and is a good priming.

- IT systems start acquiring SNOMED CT concepts through communication.
- Once sufficient data is SNOMED CT-encoded in the systems' databases, then other usages become possible (analytics, retrieval, link to knowledge resources ...)

Implementation Approaches

Start using SNOMED CT as:



What	Why
A code system	To store the clinical information exchanged
An interface terminology	To capture & display information exchanged
An indexing system	To retrieve clinical information
A common terminology	To communicate in a meaningful way
	To integrate heterogeneous data
A dictionary	To query, analyze and report
	To link health records to knowledge resources
Extensible foundation	To represent new types of clinical data



Agenda:

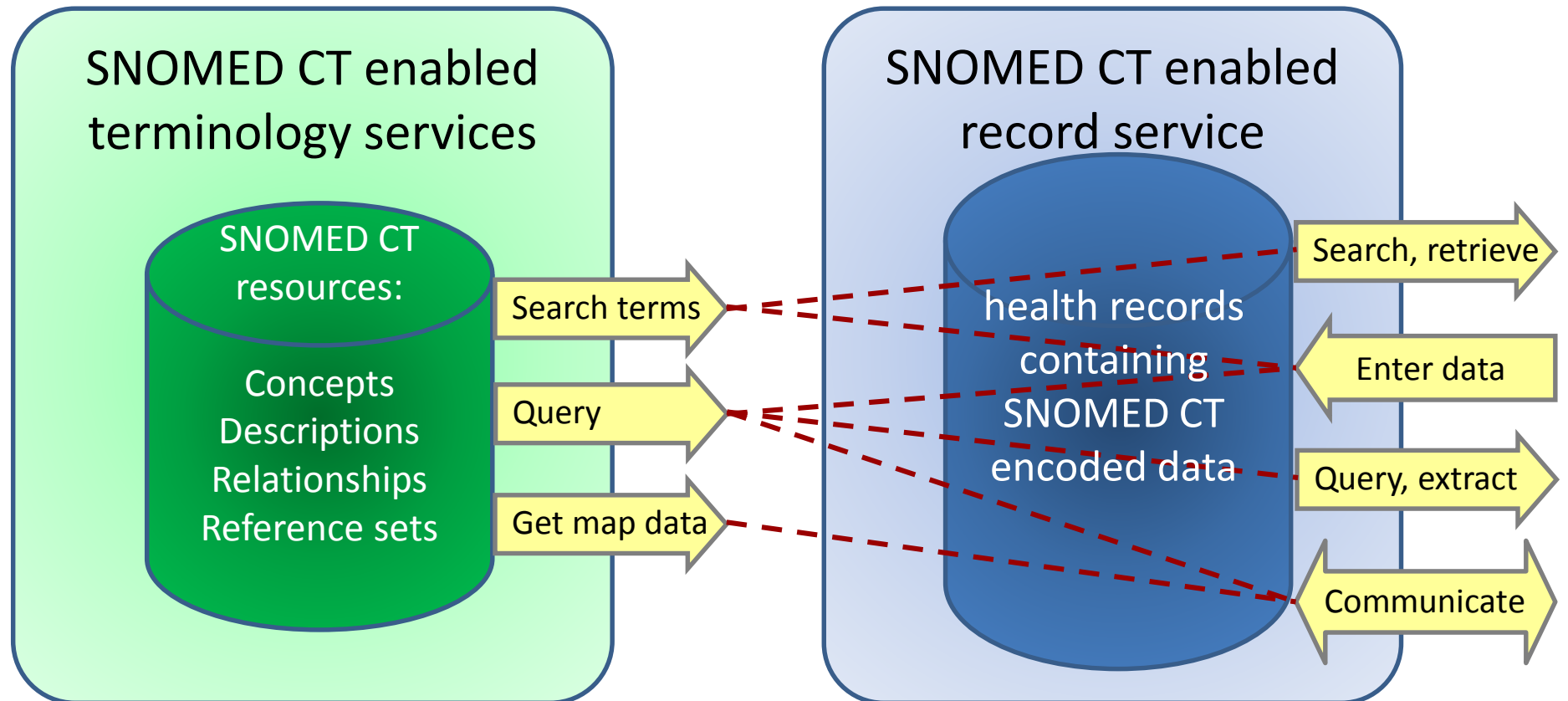
1. Meaningful data in health IT systems: why and how?
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Search terms: find a concept based on matching search term

Query:

- Testing membership in reference sets
- Testing concepts for subsumption
- Traversing, navigating hierarchies
- Testing defining relationships against given criteria
- Testing expressions for equivalence and subsumption

Get map data: Retrieve maps of SNOMED CT to or from other code systems.



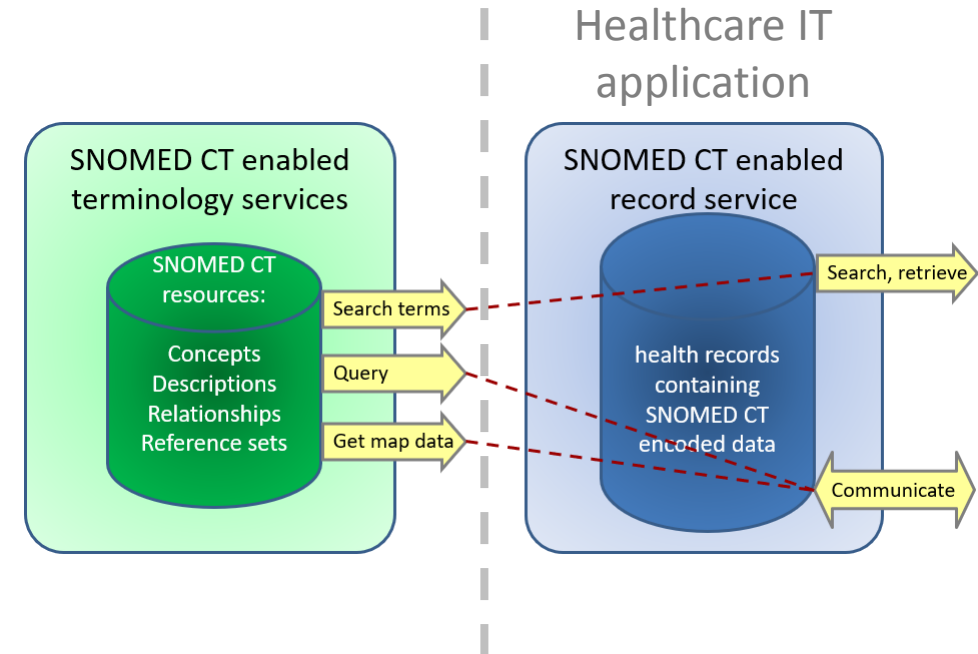
An easier first step for health IT solutions using SNOMED CT as a common terminology: Keep the terminology services external



To send meaningful data, the application must have been able to enter coded concepts, therefore **search terms**.

To receive and integrate meaningful data, the application must be able to:

- Check that a received concept belongs to a value set, therefore **test membership in reference sets (reduced Query service)**.
- Map back and forth SNOMED CT concepts to local codes, therefore **Get map data**.



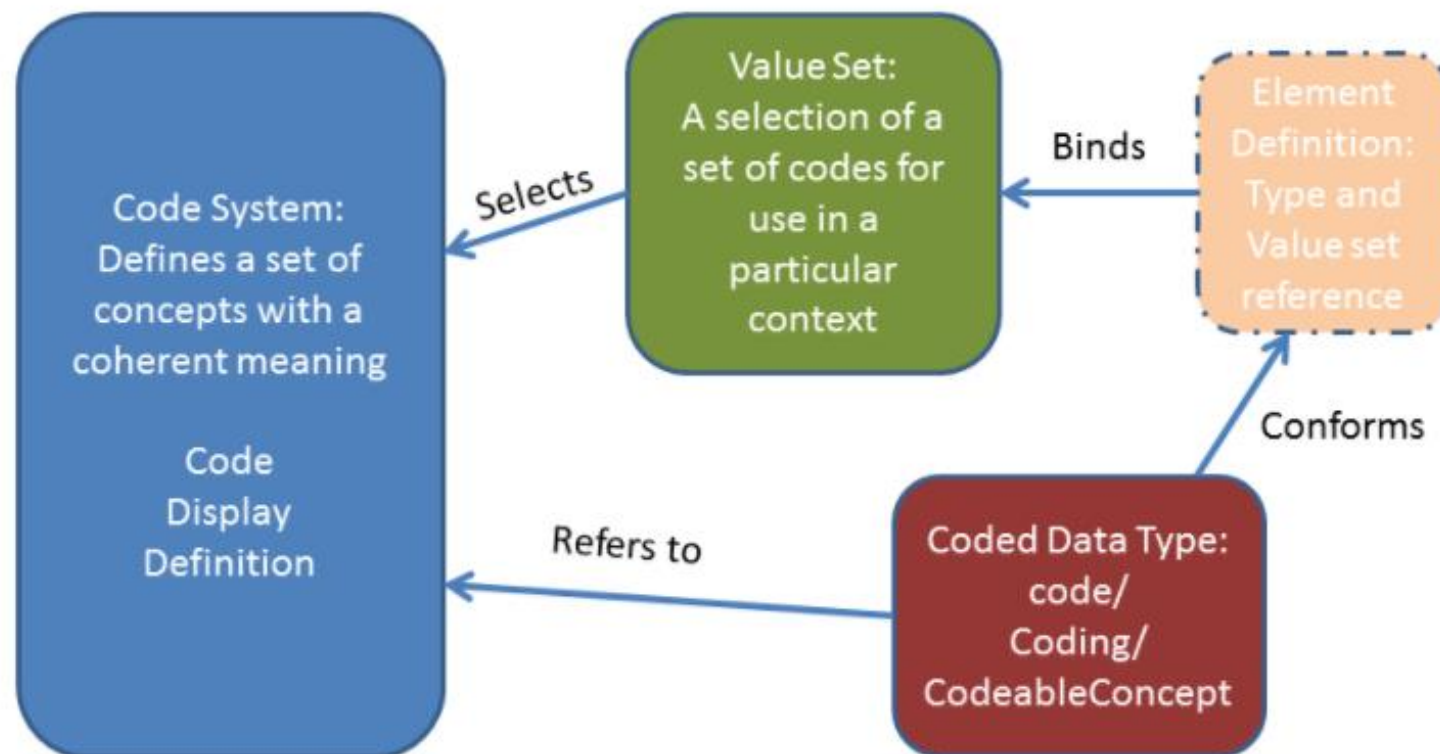
To use SNOMED CT for interoperability, the application needs limited terminology services



FHIR terminology services fit this first need



The terminology resources and their relationships in FHIR Terminology module



[4.0 Terminology Module](#)

Terminology services needed by a system, which uses SNOMED CT as a common terminology



Action	Minimal terminology services needed
Communicate in a meaningful way	<ul style="list-style-type: none">• Test membership in reference sets (to check that a received code is part of the value set)• Get map data (to convert back and forth with local code systems)
Capture & display the clinical information that is exchanged	<ul style="list-style-type: none">• Search terms• Expand value set (to obtain a pick list and browse through it to capture the proper concept)
Store the clinical information exchanged	none



FHIR resources and operations needed

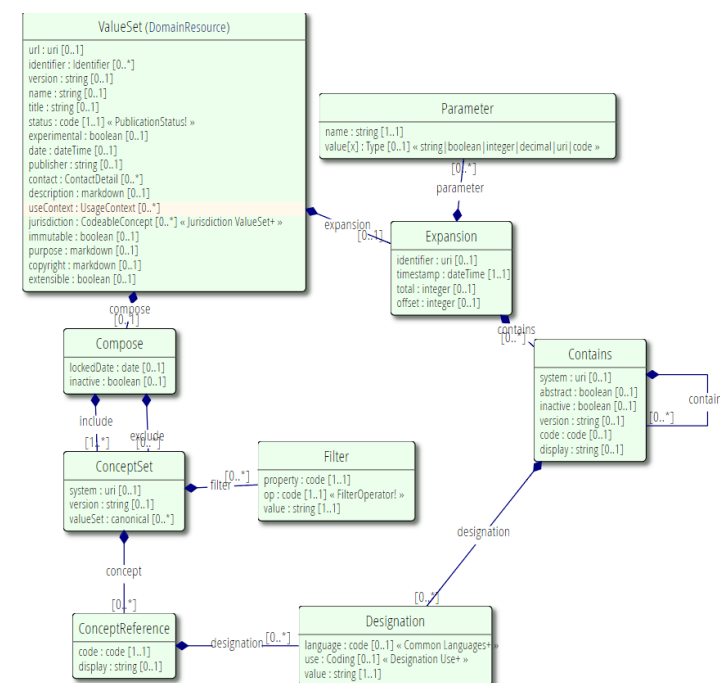


- Test membership in reference sets (verify that a code received belongs to the authorized value set)
 - Operation `$validate-code` on the `ValueSet` resource, which is candidate normative for R4.

Main parameters:

- the code received
- The value set to look into

See [4.8.1.2 Operation \\$validate-code on ValueSet](#)





FHIR resources and operations needed



- **Expand a value set** (to obtain a pick list and browse through it to capture the proper concept)
 - Operation **\$expand** on the **ValueSet** resource

Main parameters:

- the value set url
- a context
- a filter may be applied to restrict the subset of codes to be returned

See [4.8.1.1 Operation \\$expand on ValueSet](#)



FHIR resources and operations needed



- **Search term** (for user interface: find a concept corresponding to a textual description, to be captured for a field in a form)
 - Operation **\$expand** on the **ValueSet** resource, using filters

Main parameters:

- the value set url
- a filter on the display term, restricting to the concepts that contain this term.

Example:

GET [base]/ValueSet/23/\$expand?filter=urinalysis

See [4.8.1.1 Operation \\$expand on ValueSet](#)



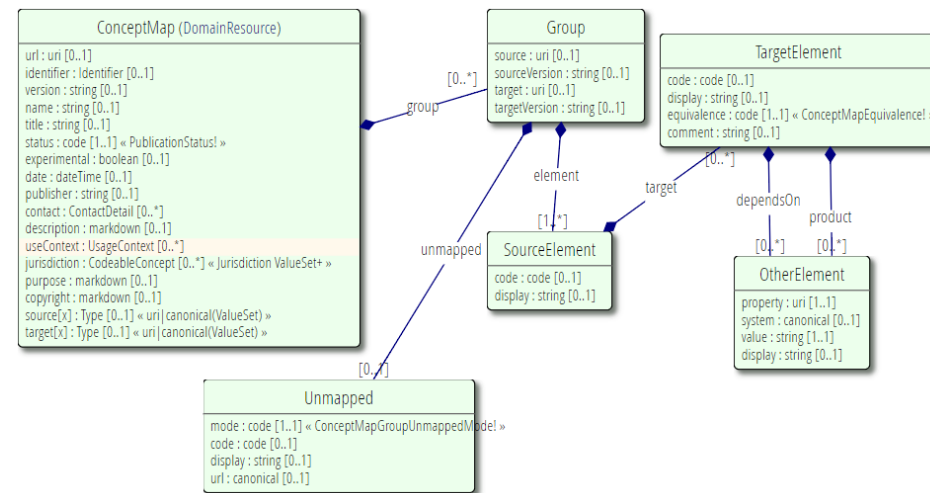
FHIR resources and operations needed



- **Get map data** (convert back and forth with local code systems)
 - Operation **\$translate** on the **ConceptMap** resource, candidate normative for R4.

The operation translates a code from one value set to another, based on the existing **ValueSet** and **ConceptMap** resources, and/or other additional knowledge available to the server.

- Works in both directions, and returns qualified matches for a concept.



See 4.9.1.1 [Operation \\$translate on ConceptMap](#)



Conclusion

- A full integration of the terminology SNOMED CT into a healthcare application, based on the RF2 release files represents a significant investment.
- For a first step aiming at using SNOMED CT as a common terminology, leveraging external FHIR terminology services is likely to minimize the adaptation effort.



Agenda:

1. Meaningful data in health IT systems: why and how?
2. Adoption of SNOMED CT needs to be stepwise
3. Leveraging external FHIR terminology services
4. Leveraging FHIR catalogs

Beyond terminology services



- Standardized vocabularies (SNOMED CT, LOINC, EDQM, UCUM, ICD-10, ...) are also part of master data sets,
 - like a reference **metathesaurus** (e.g.; the thesaurus of all medications that can be prescribed in a country)
 - or a **catalog** reflecting an organization (e.g.; the compendium of in vitro diagnostic services that can be placed to a reference laboratory).
- Other services are needed to share these master data sets.

Catalogs of orderable services/products with FHIR R4

OVERVIEW

SUMMARY OF OUTCOME FROM CONNECTATHON 17

Some examples of catalogs of orderable services/products

- ☐ Drug formulary
- ☐ Catalog of medical devices
- ☐ Laboratory compendium (similar content as eDOS HL7 v2 IG)
- ☐ List of tests supported by an IVD device
- ☐ Directory of health care services
- ☐ ...

Overview: Representing a catalog in FHIR R4

<http://build.fhir.org/catalog.html>
= Profile(Composition)

class = kind of catalog

section

entry Reference([EntryDefinition](#))

entry Reference([EntryDefinition](#))

...

entry Reference([EntryDefinition](#))

A catalog of orderable items is a collection of [EntryDefinition](#) resources.

Some of these [EntryDefinition](#) represent the orderable items themselves, while others represent resources supporting the orderable items.

EntryDefinition

type	(e.g. drug, service, device)
purpose	(orderable supporting)
referencedItem	Reference (Medication, Device, Organization, HealthcareService, ActivityDefinition, ObservationDefinition, SpecimenDefinition ...)
...	
Other metadata for this entry	
...	

Each orderable item (service/product) is represented by an EntryDefinition resource in the catalog, and is the referencedItem of that resource.

[0..*]
relatedEntry

RelatedEntry

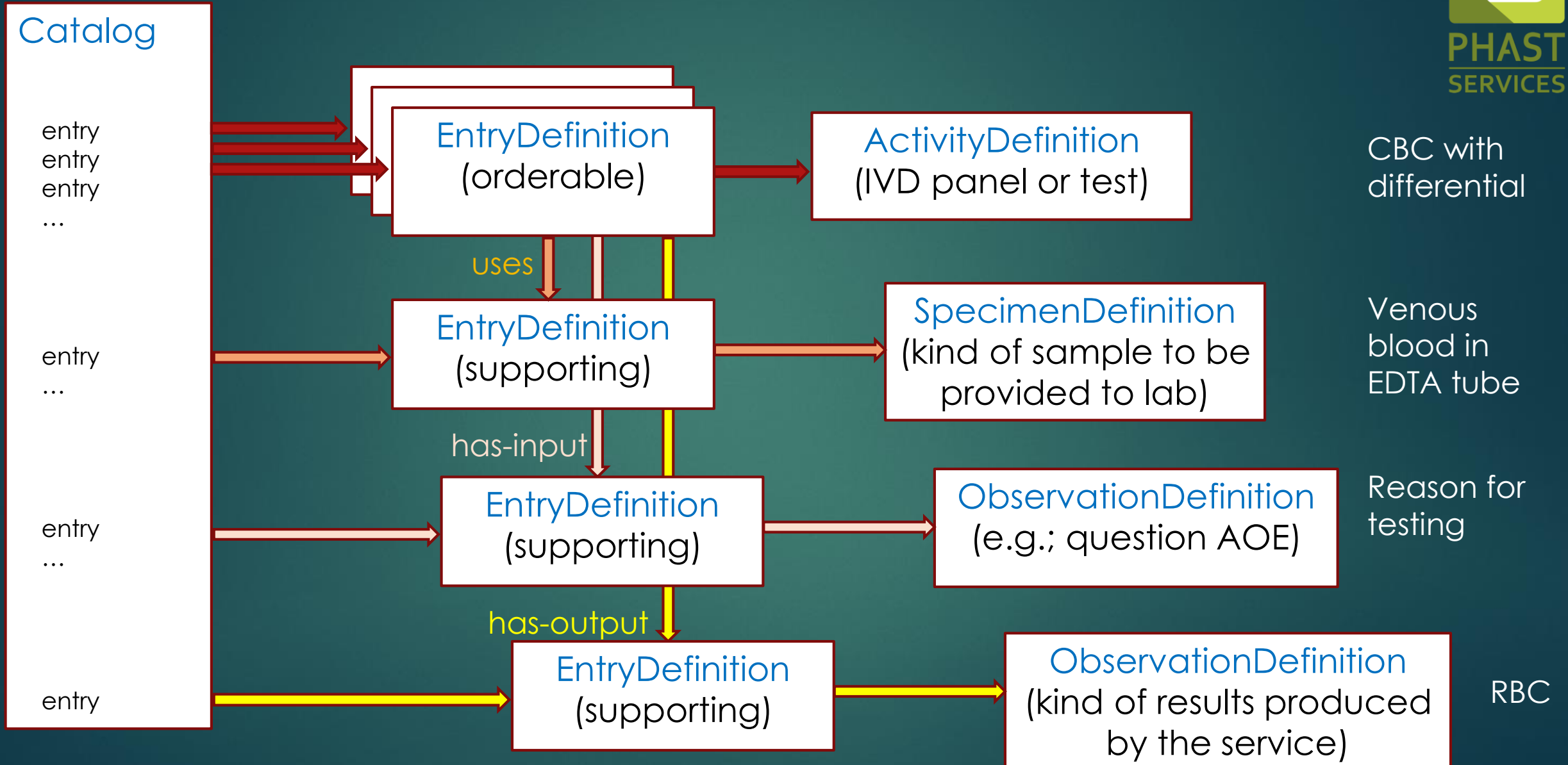
relationType	(e.g. has-input, uses, has-output, ...)
item	Reference (EntryDefinition)

Every resource contributing to the description of an orderable item is the item of a RelatedEntry referenced by the entries it contributes to.

Track “catalog” at connectathon 17


- ❑ Catalog: a clinical laboratory compendium
- ❑ 1 server (Phast) and 1 client (University of Utah)
- ❑ Scenario: Read access to the catalog
 - browse catalogs,
 - query the content of a catalog,
 - retrieve the detail of an orderable lab service (panel/test)
- ❑ Goal: verify the appropriateness of the new resources

Organization of the lab catalog



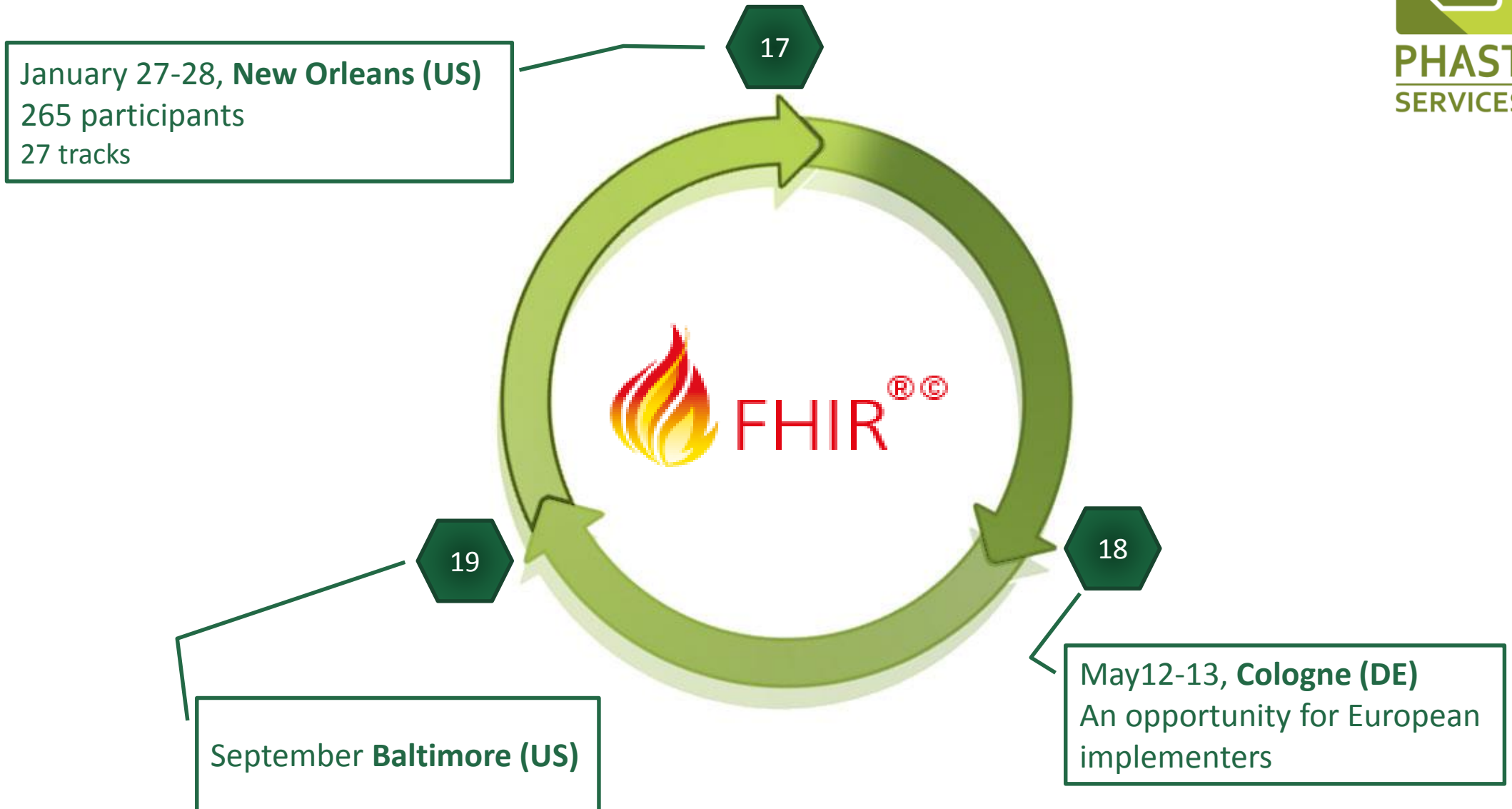
Outcome of connectathon 17



- ❑ Catalog Server  metathesaurus built by Phast-Services
- ❑ Catalog client developed by University of Utah
- ❑ Scenario: read access to the catalog; successful.
- ❑ Brings refinements to value sets, data types and search parameters for catalog resources.
- ❑ Catalog weekly calls (Wednesday 7pm, Europe central time).
See catalog project wiki page:

http://wiki.hl7.org/index.php?title=Order_Catalog_Interface

Three connectathons FHIR per year:



What's to be gained by attending a FHIR Connectathon?



- Join a community of FHIR users
 - Bring Questions and share your challenges
 - Help others by sharing your knowledge
- Develop and test your system and use the standard
- Demonstrate what's possible
- Refine the FHIR Specification



In summary:

- Health IT systems need international reference terminologies to support efficient coordination of care, locally, nationally, cross-border, cross-continent.
- SNOMED CT is the largest and inescapable one.
- Among the numerous purposes that SNOMED CT can serve, common terminology for meaningful information exchange seems an easy 1st step.
- Particularly for this 1st step, rather than a full integration by applications, it seems wise and affordable to leverage external FHIR terminology services.
- The new FHIR catalog services enable the sharing of large master data sets, making an extensive use of reference terminologies.



Thank you

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