

# Introduction to LOINC and RELMA

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## Why LOINC?

“Within one laboratory, local jargon terms may be used which are usually well understood between colleagues, but would not be sufficiently widely known for communication with the outside world.”

U. Forsum et al., Pure Appl. Chem. 72:555-745, 2000 *Properties and Units in the Clinical Laboratory Sciences Part VII. Properties and Units in Clinical Microbiology*

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## Anatomy of a LOINC Term

5193-8:Hepatitis B virus surface Ab:ACnc:Pt:Ser:Qn:EIA

5193-8

Hepatitis B virus surface Ab

ACnc

Pt

Ser

Qn

EIA

LOINC Code

Component
Property Measured
Timing
System
Scale
Method

There are six major LOINC axes

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## Acknowledgements

- RELMA Development Team
  - John Hook, Mark Fisher, Ryan Phillips, Karen Ahmad, and more!
- Kathy Mercer
- Clem McDonald
- Dan Vreeman
- The Lab LOINC Committee
- Funding Support
  - NLM, Regenstrief Institute, NCI, CDC

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## Test comparisons

What you see in the order list

### Lab A

Test Name: Lyme Disease Serology  
Measures: *B. burgdorferi* Ab IgG  
Method: ELISA  
Scale: quantitative  
e.g.: Titer 1:40

LOINC Code = 5062-5

### Lab B

Test Name: Lyme Disease Antibody  
Measures: *B. burgdorferi* Ab IgM  
Method: Immune blot  
Scale: qualitative  
e.g.: Positive

LOINC Code = 6321-4

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## What is NOT part of a LOINC Name?

- The instrument used in testing
- Specific details about the specimen
- Priority (e.g. STAT)
- Where testing was done
- Who did the test
- Test interpretation
- Anything that is not an intrinsic part of the name of the result
- Other things that are carried in;
  - The OBX segment
  - An HL7 Version 3 Observation Object

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## Component/Analyte

The substance or entity that is measured, evaluated, or observed

- Sodium
- Glucose
- Brucella sp. organism
- Influenza A Virus antigen
- Cytomegalovirus Virus antibody
- Lipids.Total

5193-8:Hepatitis B virus surface Ab:ACnc:Pt:Ser:Qn:EIA

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## Property

*the most difficult LOINC axis*

The characteristic or attribute of the analyte that is measured, evaluated, or observed

### Major Categories

- **Mass:** Observations reported with mass (milligrams, grams, etc.) in the numerator of their units of measure
- **Substance:** Observations reported with moles or milliequivalents in the numerator of their units of measure
- **Catalytic activity:** Observations that report enzymatic activity
- **Arbitrary:** Results that report arbitrary units in the numerator of their units of measure
- **Number:** Counts

5193-8:Hepatitis B virus surface Ab:ACnc:Pt:Ser:Qn:EIA

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## Timing\*

The interval of time over which the observation or measurement was made

- Pt - at a point in time
- 12H - a twelve hour collection
- 24H - a twenty four hour collection

\*Public Health Lab timings are usually Pt

\*non-Pt timings are usually associated with Ratio Property

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## System

The system (context) or specimen type upon which the observation was made.

- Ser - Serum
- Bld - Whole blood (RBC)
- Ur- Urine
- BldA - Arterial blood
- Liver - Liver
- Flu - Body Fluid, unspecified
- Gast – Gastric fluid/contents
- Food – Food or feedstuff
- Tiss – Tissue
- XXX – To be specified in another part of the message

### Super System

Second subpart (^). When not included, “patient” is the default. Used to indicate blood product unit (BPU), a bone marrow donor, or a fetus.

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## Scale

- Qn - quantitative
  - Continuous numeric (real, integer, ratio)
  - Optional operator ( $>$ ,  $\geq$ ,  $\leq$ ,  $<$ )
- Ord - ordinal
  - a ranked set of possible values (1+, 2+, 3+)
- Nom - nominal
  - an unranked collection of possible values
  - a taxonomy (e.g. list of bacteria)
- Nar - narrative
  - free text narrative (e.g., visit note)

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## Method

- Methods only needed if interpretation affected
  - Different normal ranges
  - Test Sensitivity
- Listed only at the generic level
  - Agglutination
  - Immunoassay
  - Probe with target amplification
- Methods may also be sent in OBX.17

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## Mapping Terms to LOINC

### Things to Remember

- The thing ordered is not always the thing measured:
  - Blood Culture – live organism(s) identified
  - VDRL – *Treponema pallidum Ab*
  - Urinalysis – lots of different things
- The question (what am I measuring?) is not the answer (e.g. Pos)
  - You are mapping the question, not the answer!
- You must know the specifics of the component being tested for (what is this test actually measuring?)

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# RELMA®

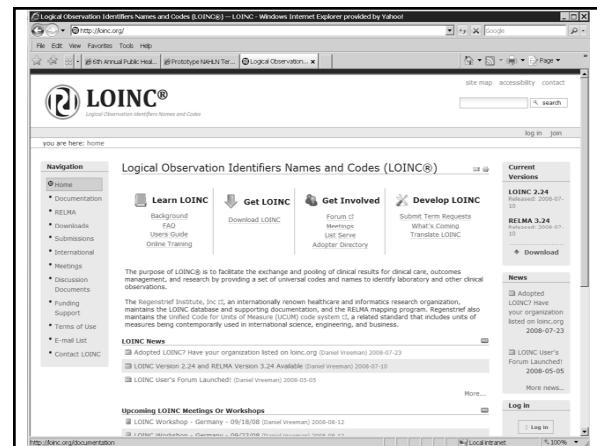
Regenstrief LOINC Mapping Assistant



## RELMA Functions

- Manual and automated mapping assistant
- Same free use as LOINC (see license)
- Comes with the LOINC files and indexes
- RELMA tools transform local words in local file
  - User creates file of local term/name and codes
- Assigns LOINC term to local test/battery code

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## RELMA Search logic

For a given test:

- User can select LOINC records containing:
  - A set of words connected by ANDs or ORs
  - Additional keyed in words
  - A particular category of test (e.g. microbiology)
  - Wild cards of ? and \*
  - Selected classes, systems, components,
  - Any intersection of the above

Note:

RELMA assumes *exact match* on word unless user adds terminal “\*” to indicate wild card.

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## RELMA Logic

Not → #<word A>

Or → <word A> | <word B>

And → <word B> <word A>

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## Importing Local Terms into RELMA



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## 4 Ways to Load Local Master Observation Files

- Direct entry into LMOF from within RELMA (painful)
- Create an Access table that mimics the LMOF structure (less painful but tedious)
  - Appendix A: RELMA Manual
- Create a delimited ASCII file from your local test catalog (good choice)
- Load directly from HL7 v2.x messages (best? choice)
  - Pulls data from OBR and OBX segments

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### Example Delimited File

	312370	MIC	309651	Nafcilin
665670	Basic Metabolic Panel	754689	Anion Gap	mmol/L
665670	Basic Metabolic Panel	755363	Creatinine SerP1 QN	mg/dL
665670	Basic Metabolic Panel	755364	Sodium SerP1 QN	mmol/L
665670	Basic Metabolic Panel	755365	Potassium SerP1 QN	mmol/L
665670	Basic Metabolic Panel	755366	Chloride SerP1 QN	mmol/L
665670	Basic Metabolic Panel	755367	Carbon Dioxide SerP1 QN	mmol/L
665670	Basic Metabolic Panel	755368	Glucose SerP1 QN	mg/dL
665670	Basic Metabolic Panel	755369	BUN SerP1 QN	mg/dL
665670	Basic Metabolic Panel	755370	Calcium Total SerP1 QN	mg/dL
667688	Factor VIII:R:Assay	755117	Factor VIII:R:Assay	%
667688	Factor X:Q:Assay	755119	Factor X Assay	%
667688	Factor XII:Q:Assay	755181	Factor XI Assay	%
684308	ABG and EtCO <sub>2</sub>	810892	Am-B	
684308	ABG and EtCO <sub>2</sub>	810893	Am-B	
685703	Cooxometry Panel Bld QN	755435	Carboxyhemoglobin Bld QN	%
685703	Cooxometry Panel Bld QN	755474	Methemoglobin Bld QN	%
685703	Cooxometry Panel Bld QN	755475	Oxyhemoglobin Bld QN	%
685733	CRP SerP1 QN	755479	C-Reactive Prot SerP1 QN	mg/dL
685796	Drug Sen SerP1 QL	755496	Tricyclic Antidepressants Sen SerP1 QL	
685796	Drug Sen SerP1 QL	755497	Salsalate SerP1 QN	mg/dL
685796	Drug Sen SerP1 QL	755498	Ethanol PI QN	mg/dL
685849	Hgb A1c HPLC Bld QN	755572	Hemoglobin A1c HPLC Bld QN	%
685849	Hgb A1c HPLC Bld QN	755573	Hemoglobin A1c, Postprandial	%
686283	PTH Intact Ser QN	755757	PTH Intact Ser QN	pg/mL
693154	B Cell Subsets	755895	B Cells IgG	%
693154	B Cell Subsets	755896	B Cells IgM	%
693154	B Cell Subsets	755897	B Cells IgD	%
693154	B Cell Subsets	755898	B Cells IgA	%
693154	B Cell Subsets	755906	Flow Cytometry Interpretation	
693154	B Cell Subsets	2773454	B cell Subset	
693154	B Cell Subsets	2724661	Flow Cytometry ASB Comment	

### Fields Segregated

**Import Local Terms from Delimited File**

Name of Local Term File: LOINC Workshop OBR  
Default Lab Section: You may create a new lab section by typing a name in the box.

Choose the delimiter that separates your fields:  Tab  Semicolon  Comma  Space

**Assign LMOF Attribute**

First Row Contains Column Headers Text Qualifier: "

To assign a column of data to one of the fields, click on the second header row of the grid, click on the arrow to the right of the dropdown list, and select the field into which you desire your data to be imported. Test Code and Test Description are required fields.

#1	#2	#3	#4	#5
Ignore Field	Ignore Field	Ignore Field	Ignore Field	Ignore Field
312370 MIC	309651 Nafcilin			
312370 MIC	708007 Moxifloxacin			
665670 Basic Metabolic Panel	754689 Anion Gap	mmol/L		
665670 Basic Metabolic Panel	755363 Creatinine SerP1 QN	mg/dL		
665670 Basic Metabolic Panel	755364 Sodium SerP1 QN	mmol/L		
665670 Basic Metabolic Panel	755365 Potassium SerP1 QN	mmol/L		

Case-sensitive Import

Import\_sample\_OBR.txt

### Open File in RELMA

**Import Local Terms from Delimited File**

Name of Local Term File: LOINC Workshop OBR  
Default Lab Section: Choose your working set. RELMA allows multiple sets in LMOF database

Choose the delimiter that separates your fields:  Tab  Semicolon  Comma  Space  Other:

First Row Contains Column Headers Text Qualifier: "

Select your delimiter To assign a column of data to one of the fields, click on the arrow to the right of the dropdown list, and select the field into which you desire your data to be imported. Test Code and Test Description are required fields.

#1	#2	#3	#4	#5
312370 MIC	309651 Nafcilin			
312370 MIC	708007 Moxifloxacin			
665670 Basic Metabolic Panel	754689 Anion Gap	mmol/L		
665670 Basic Metabolic Panel	755363 Creatinine SerP1 QN	mg/dL		
665670 Basic Metabolic Panel	755364 Sodium SerP1 QN	mmol/L		
665670 Basic Metabolic Panel	755365 Potassium SerP1 QN	mmol/L		

Case-sensitive Import

Import\_sample\_OBR.txt

### Ready to Import

**Import Local Terms from Delimited File**

Name of Local Term File: LOINC Workshop OBR  
Default Lab Section: You may create a new lab section by typing a name in the box.

Choose the delimiter that separates your fields:  Tab  Semicolon  Comma  Space  Other:

First Row Contains Column Headers Text Qualifier: "

Minimum assignments To assign a column of data to one of the fields, click on the second header row of the grid, click on the arrow to the right of the dropdown list, and select the field into which you desire your data to be imported. Test Code and Test Description are required fields.

#1	#2	#3	#4	#5
Battery Code	Battery Description	Test Code	Test Description	Units
312370 MIC	309651 Nafcilin			
312370 MIC	708007 Moxifloxacin			
665670 Basic Metabolic Panel	754689 Anion Gap	mmol/L		
665670 Basic Metabolic Panel	755363 Creatinine SerP1 QN	mg/dL		
665670 Basic Metabolic Panel	755364 Sodium SerP1 QN	mmol/L		
665670 Basic Metabolic Panel	755365 Potassium SerP1 QN	mmol/L		

Case-sensitive Import

Import\_sample\_OBR.txt

## Alternative (best) Way

- Use large set of HL7 messages
- Automatically make dataset of:
  - OBR ID
  - OBR description
  - OBX ID
  - OBX description
  - Sample of results with
    - Real values
    - Units
    - Abnormal flags
    - Normal ranges

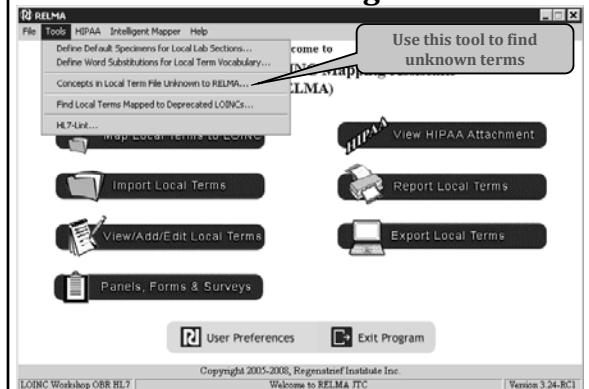
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## Preparing your Data for Mapping

- Improve mapping success by:
  - Expanding abbreviations
  - Standardizing colloquial terms
  - Ignoring “administrative” terms
  - Standardizing time references
- Can be done prior to importing
- Better to use tools built into RELMA

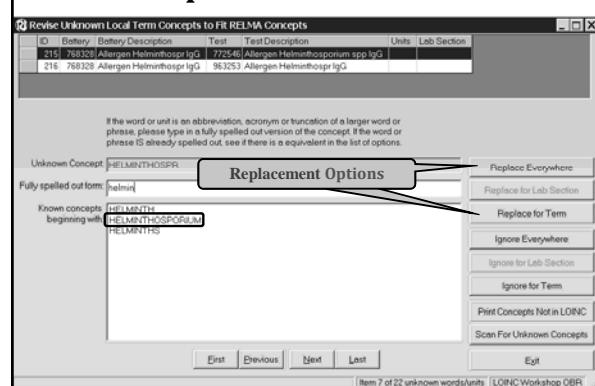
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## RELMA Cleaning Tools

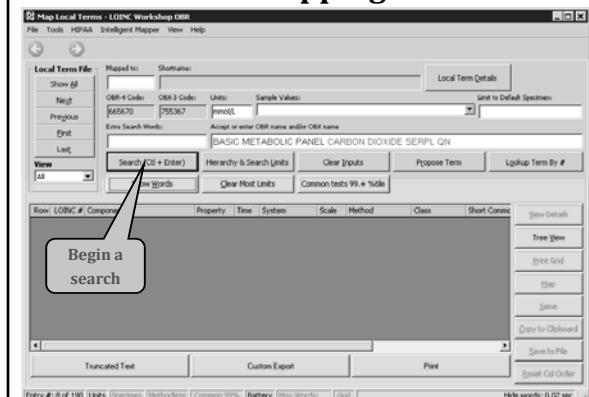


Use this tool to find unknown terms

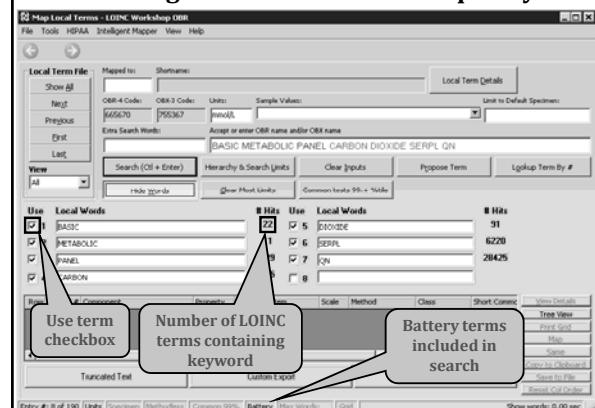
## Replace Local Terms



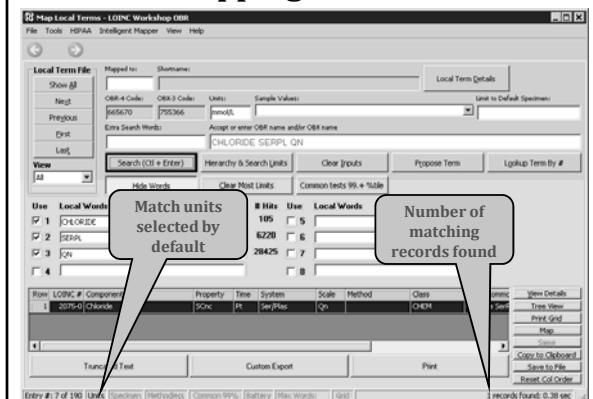
## Standard Mapping Screen



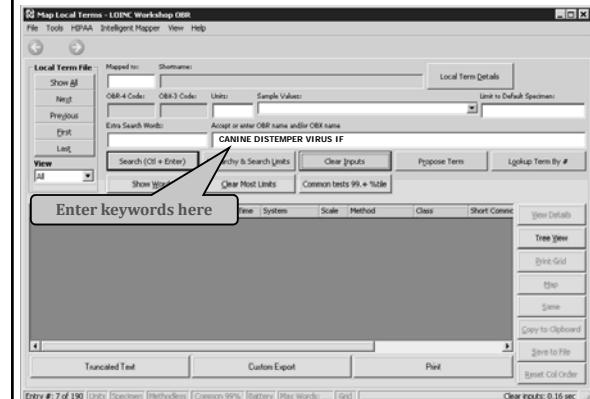
## Showing Search Terms and Frequency



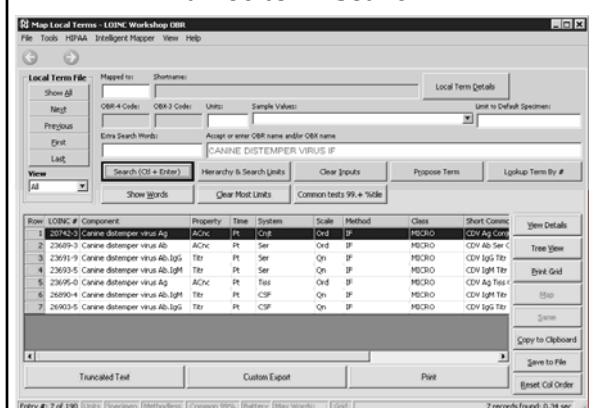
## Mapping Results



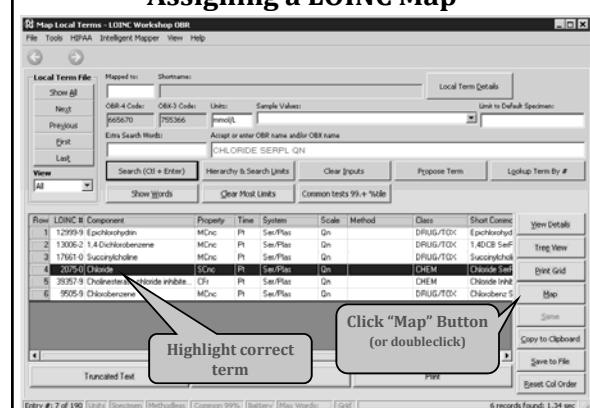
## Ad hoc term search



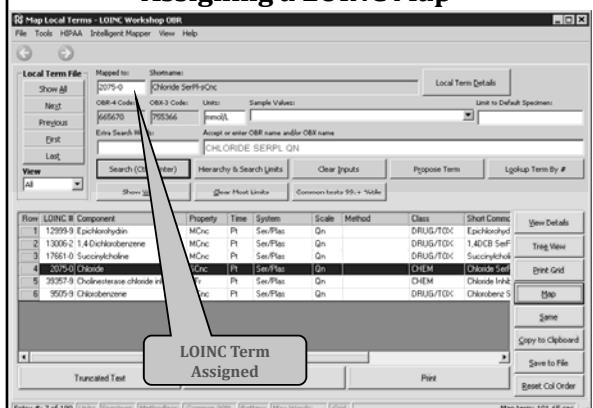
## Ad hoc term search



## Assigning a LOINC Map



## Assigning a LOINC Map



## Setting Search Limits

- Narrows search to specific subset of LOINC terms
- Reduces number of candidate terms
- Limits can be applied to all components
- Component attribute can be further restricted by number of words
- Tree structure allows for hierarchical constraints

## RELMA - Search Constraints

- Controls general search constraints including:
  - Limit to LOINC terms compatible with submitted units
  - Force match with any specimen contained in name
  - Methodless terms only (*now enhanced*)
  - Limit to components with N or fewer words in their name
  - Pop up search timing statistics after each search
- Use carefully or search may not be successful  
(Note parallel control switches at bottom of screen)

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## Terms Consistent with Specimen

The screenshot shows a software interface for managing local terms. In the search bar, 'ASPERGILLUS FLAVUS' is entered. The results grid shows three entries, all of which are 'CSF' (Composite Specimen Flavus). A callout box points to the message 'Only CSF Terms are returned'.

Row	LOINC #	Component	Property	Time	System	Scale	Method	Class	Short Comm
1	13119-0	Aspergillus flavus Ab	ACnc	Pr	CSF	Qn	Micro	MICRO	A flavus Ab
2	15415-2	Aspergillus flavus Ab	ACnc	Pr	CSF	Qd	Micro	MICRO	A flavus Ab C
3	21084-9	Aspergillus flavus Ab	ACnc	Pr	CSF	Qd	Immune diffusion	MICRO	A flavus Ab C

## Setting Search Limits

The screenshot shows the 'Search Constraints' tab of the software. Under 'Restrict Search', there is a checkbox labeled 'Methodless Terms Only'. A callout box points to this checkbox with the label 'Methodless Terms Restriction'. Another callout box points to the 'Override Methodless Terms Restriction' link below it.

## Limit to Methodless Terms

The screenshot shows the same software interface with the 'Methodless Terms Only' restriction applied. The results grid now only displays methodless terms, specifically 'Tir' (Titration) and 'BodyFld' (Body fluid). A callout box points to the message 'Only Methodless Terms Appear'.

Row	LOINC #	Component	Property	Time	System	Scale	Method	Class	Short Comm
1	13119-0	Aspergillus flavus Ab	ACnc	Pr	CSF	Qn	Micro	MICRO	A flavus Ab
2	15415-2	Aspergillus flavus Ab	ACnc	Pr	CSF	Qd	Micro	MICRO	A flavus Ab C
3	15725-1	Aspergillus flavus Ab IgG	ACnc	Pr	Ser	Qd	Micro	MICRO	A flavus Ab J
4	21084-9	Aspergillus flavus Ab	ACnc	Pr	Ser	Qd	Immune diffusion	MICRO	A flavus Ab J
5	23032-4	Aspergillus flavus Ab	ACnc	Pr	Ser	Qd	Immune diffusion	MICRO	A flavus Ab J
6	27309-4	Aspergillus flavus Ab	Tir	Pr	Ser	Qn	Immune diffusion	MICRO	A flavus Ab J
7	27307-1	Aspergillus flavus Ab	ACnc	Pr	BodyFld	Qd	Immune diffusion	MICRO	A flavus Ab J
8	31236-9	Aspergillus flavus Ab	ACnc	Pr	BodyFld	Qd	Immune diffusion	MICRO	A flavus Ab J
9	43093-0	Aspergillus flavus Ab	ACnc	Pr	Ser	Qn	Micro	MICRO	A flavus Ab J
10	43094-8	Aspergillus flavus Ab	ACnc	Pr	Ser	Qn	Micro	MICRO	A flavus Ab J

## Limit to Methodless Terms

The screenshot shows the same software interface with the 'Methodless Only' restriction applied. The results grid only displays methodless terms, specifically 'Tir' (Titration) and 'BodyFld' (Body fluid). A callout box points to the message 'Only Methodless Terms Appear'.

Row	LOINC #	Component	Property	Time	System	Scale	Method	Class	Short Comm
1	13119-0	Aspergillus flavus Ab	ACnc	Pr	CSF	Qn	Micro	MICRO	A flavus Ab
2	15415-2	Aspergillus flavus Ab	ACnc	Pr	CSF	Qd	Micro	MICRO	A flavus Ab C
3	15725-1	Aspergillus flavus Ab IgG	ACnc	Pr	Ser	Qd	Micro	MICRO	A flavus Ab J
4	21084-9	Aspergillus flavus Ab	ACnc	Pr	Ser	Qd	Immune diffusion	MICRO	A flavus Ab J
5	23032-4	Aspergillus flavus Ab	ACnc	Pr	Ser	Qd	Immune diffusion	MICRO	A flavus Ab J
6	40905-0	Aspergillus flavus Ab	Tir	Pr	Ser	Qn	Immune diffusion	MICRO	A flavus Ab J
7	47423-3	Aspergillus flavus B Ab	ACnc	Pr	Ser	Qd	Micro	MICRO	A flavus B Ab
8	47422-1	Aspergillus flavus H Ab	ACnc	Pr	Ser	Qd	Micro	MICRO	A flavus H Ab
9	6024-4	Aspergillus flavus Ab IgE	ACnc	Pr	Ser	Qd	ALLERGY	ALLERGY	A flavus IgE
10	34354-8	Aspergillus flavus Ab	ACnc	Pr	Ser	Qd	Micro	MICRO	A flavus Ab J

## More on Methodless Terms

- Some LOINC categories do not have methodless terms
- Checking methodless only will remove these from view on results grid
- Checking additional box allows these to be seen

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## Methodless Only Checked

**Only Methodless terms returned**

Row	LOINC # Component	Property	Time	System	Scale	Method	Class	Short Comm
1	22574-7 Viral hemorrhagic disease virus	Acn	Pt	Ser	Ord		MICRO	RHDV Ab S
2	22574-8 Viral hemorrhagic disease virus	Tb	Pt	Ser	Qn		MICRO	RHDV Ab T
3	21593-2 Viral hemorrhagic disease virus	Acn	Pt	Ser	Qn		MICRO	RHDV Ab S

Differ in one or more components

## Conditional Methodless

**More terms returned**

Row	LOINC # Component	Property	Time	System	Scale	Method	Class	Short Comm
1	22574-9 Viral hemorrhagic disease virus	Acn	Pt	Tes	Ord	Microscopy elect.	MICRO	RHDV Tiss
2	22574-8 Viral hemorrhagic disease virus	Acn	Pt	Ser	Ord		MICRO	RHDV Ab T
3	22574-7 Viral hemorrhagic disease virus	Acn	Pt	Ser	Qn		MICRO	RHDV Ab S
4	22574-2 Viral hemorrhagic disease virus	Tb	Pt	Ser	Qn		MICRO	RHDV Ab T4
5	22504-4 Viral hemorrhagic disease virus	Acn	Pt	Tes	Ord	EIA	MICRO	RHDV Ag Tu
6	22504-6 Viral hemorrhagic disease virus	Acn	Pt	Tes	Ord	IF	MICRO	RHDV Ag Tu
7	21592-0 Viral hemorrhagic disease virus	Acn	Pt	Tes	Ord	Immune stain	MICRO	RHDV Ag Tu

No methodless term; all shown

## Setting Search Limits

**Limit to Lab Tests Only (No Clinical LOINC Terms)**

Restrict Search

Common Test Restrictions

- Terms consistent with local units
- Terms consistent with specimen
- Terms consistent with local laterality
- Common Test Restrictions
- Lab Tests Only
- Pharma Tests Only
- Common Tests 99+ % Nucleic
- Include deprecated LOINCs

Methodless Terms Only

If no methodless version exists, then return the method specific terms

## Navigating through the Mapping Process



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## LOINC Mapping Tactics

- Try using method-less terms first
  - Specific methods (if needed) can be transmitted in:
    - OBX-17 (v2.x)
    - Observation.methodCode (v3.0)
- If specimen type is transmitted in the message (OBR or SPM), limit system to "XXX"
- Examine local units or real results to verify correct properties
  - Properties are rarely distinguishable in tests
- With every release - Update previous mappings to identify deprecated terms

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## Common Mapping Issues

- Locally Defined Test Name Ambiguity
  - Reuse of local test code
- "Analyte-free" Local Test Names
- Incongruent Value sets (Scale ambiguity)
- Result vs. Interpretation
- Available LOINC Terms too Specific
- Available LOINC Terms too General
- Panel vs. Discrete Test
  - Common in Microbiology

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## Search Hints and Tips

- Keywords with zero frequency are ignored
  - May need to rephrase – use synonym
- Some causes for no returned terms
  - Too many keywords in search – uncheck some
  - Limits applied that don't make sense
    - E.g. Method-less tests plus Method tree set to EIA
  - Did not find and revise words not in RELMA
    - Local units not in RELMA
- Units are GREAT discriminators
- You may have tests that need to be added to LOINC

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## Using LOINC and SNOMED Together



## Where do LOINC and SNOMED Fit?

### Remember:

- LOINC represents the question:
  - Is there any Botulism toxin in my specimen? (33708-9)
  - Organisms identified in specimen? (634-6)
- SNOMED represents the answer:
  - Negative (SCTID 260385009)
  - *E. coli* O157:H7 (SCTID 103429008)

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## Where do LOINC and SNOMED Fit?

- In an HL7 message, LOINC may be used:
  - In OBR-4 (Universal Service Identifier)
  - In OBX-3 (Observation Identifier)
  - In Version 3 - Observation.code
- SNOMED may be used:
  - In OBX-5 (where nominal values are needed)
  - Almost anywhere else in an HL7 message where coded values are needed

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