# Adequate Pharmacogenomics FHIR CG IG

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For comments, questions on slides refer to:

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General questions: HL7 CG WG members, FHIR community

#### Proposed actions

- Decide on what 'High Risk Allele' scope really is. Do we need something separate for Potential for Adverse Event.
- Vote on whether or not the increased efficiency in querying constitutes a strong imperative to change the PGx profiles to components.
- Select/Approve a group to finish selecting LOINC codes and implement switch to components.
- Vote on whether the change is important enough to be a quick version update.
- PROVIDE additional documentation on use. E.g. this power point has some textual statements that clarify how specific elements are used. Place such statements in the structure definition – not data element description.
- NOTE: level of evidence is currently adequate for the PGx use case as it allows any evidence system, such as PharmGKB's classification of evidence for PGx to be used. If a group needs to constrain then that group can do so with an implementation guide based on the CG WG implementation guide.

#### Goal

- 1. NOT a knowledge resource
- 2. NOT CDS delivery
- 3. NOT a guideline
- 4. Accommodate a variety of granular reporting of Pharmacogenomics found in current lab reports.

How do the current profiles differ from the existing abstract profile?

- The Medication Implication profile is an abstract profile which the current profiles are derived from
- The profile Medication Efficacy has a specific '.code' and '.value'
- The profile Medication Transporter has a specific '.code' and '.value'
- The profile Medication Metabolism has a specific '.code' and '.value'
- The profile High Risk Allele has a specific '.code' and '.value
  - For now, suggesting using High Risk Allele for communicating Adverse Drug Event. This needs conclusive discussion
- THE PROFILES ONLY DIFFER BY USING A SPECIFIC Observation.code and value-set for Observation.value

### Current Diagram



#### Current Diagram – note on Medication Task



http://build.fhir.org/ig/HL7/genomics-reporting/task-medchg.html

### Current Diagram – Med Task

For recommendation of action such as chose alternative medication, decrease dose, increase dose, Task-med-chg is used. Take note of 'intent', 'coding', 'coding.text', 'description' and 'reason reference'

				Binding: (unbound) (example)	
intent	ΣΙ	11	code	unknown   proposal Lolan   order   original-or Binding: TaskIntent (regume !)	
				Fixed Value: proposal	For intent use (proposal /
priority	I	01	code	routine   urgent   asap   stat Binding: RequestPriority (required)	For intent use proposal.
) code	ΣΙ	11	CodeableConcept	Task Type Binding: TaskCode (example)	
🛄 id		01	string	Unique id for inter-element referencing	
🖈 extension	I	0*	Extension	Additional content defined by implementations Slice: Unordered, Open by value:url	
- 🛢 coding	ΣΙ	0*	(Slice Definition)	Code defined by a terminology system Slice: Unordered, Open by value:system	For coding.text you can put a statement like
coding:change	ΣΙ	01	Coding	E.g. Consider alternative medication, Decrease of Binding: LOINC Answer List LL4049-4 (prefer a	'Change to alternative medication' in addition to
·· 💷 id		01	string	Unique id for inter-element referencing	the code value.
- 🖈 extension	Ι	0*	Extension	Additional content defined by implementations Slice: Unordered, Open by value:url	
🛄 system	ΣΙ	11	uri	Identity of the terminology system Fixed Value: http://loinc.org	de environtione in a construit findel familieuro e e construit la
·· 💷 version	ΣΙ	01	string	Version of the system - if relevant	description is a useful field for human readable-
💷 code	ΣΙ	11	code	Symbol in syntax defined by the system	text
display	ΣΙ	01	string	Representation defined by the system	
userSelected	ΣΙ	01	boolean	If this coding was chosen directly by the user	
text	ΣΙ	01	string	Medication usage suggestion narrative	
description	ΣΙ	01	string	Human-readable explanation of task	<b>reasonReference</b> – this links to the Medication
A focus	<b>T</b> T	4.4	Deference/Mediention	What task is acting on	- Implication that has a Delated Artificat for
reasonCode	I		01 CodeableConcept	Why task is needed Binding: (unbeand) (example)	support, such as a CPIC guideline.
reasonReference	I		01 Reference(Medicatio	n Why task is needed	

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#### Current Diagram – note on Med assessed

class 4-pharmacogenomics-fig1-implications

Medication-assessed – this is where the medication that the implication is about is identified. Use a code from a system such as RxNorm, SNOMED CT or LOINC

Genomic Observation Common F Medication Implication	roperties	Medication Us	age Implications (Task) 82117-3
component:medication-assessed	ΣΙ 1*	BackboneElement	Medication Assessed
💶 id	01	string	Unique id for inter-element referencing
🔹 🖈 extension	I 0*	Extension	Additional content defined by implementations
🗝 🖈 modifierExtension	?! 0* ΣΙ	Extension	Extensions that cannot be ignored even if unrecognized
- 🗘 code	ΣΙ 11	CodeableConcept	Type of component observation (code / type) Binding: LOINCCodes (example) Required Pattern: At least the following
💶 id	01	string	Unique id for inter-element referencing
- 🌍 extension	0*	Extension	Additional content defined by implementations
- 🔒 coding	1*	Coding	Code defined by a terminology system Fixed Value: (complex)
id	01	string	Unique id for inter-element referencing
- 🎲 extension	0*	Extension	Additional content defined by implementations
- 🔒 system	11	uri	Identity of the terminology system Fixed Value: http://loinc.org
·· <b>··</b> version	01	string	Version of the system - if relevant
🔒 code	11	code	Symbol in syntax defined by the system Fixed Value: 51963-7
display	01	string	Representation defined by the system
userSelected	01	boolean	If this coding was chosen directly by the user
🛄 text	01	string	Plain text representation of the concept

# Current Diagram – note on Guidelines and Reference material



### Current Diagram – note evidence level in PGx



#### Current Diagram – note on Narrative



### Changes - diagram



## Change entails

- Medication Implication profile will have:
  - Observation.code bound to LOINC code **61357-0** <u>https://loinc.org/61357-0/</u> See the term description. Consider to request something similar or use it. Also <u>https://loinc.org/51965-2/</u>
  - Observation.value bound to a value set of the current profile LOINC codes. Binding as 'preferred' (extensible) to 'Medication Efficacy', 'Medication Transporter,' 'Medication Metabolism', 'High Risk Allele'
- Components are for providing more granular information
  - Each of 'Medication Efficacy', 'Medication Transporter,' 'Medication Metabolism', 'High Risk Allele' would be a component, with a corresponding value-set (e.g. the component for the current Medication Efficacy would use the value set from the current profile)
  - Add component for Potential adverse drug event OR use the High Risk component? I will use High Risk component (we can
    vote to change this) in this document
  - Suggest to add associated phenotype to the component for Potential Adverse Drug Event. So one *can* state an assoiated phenotype.

Think: The observation is 'what kind of variant implication does the genetic variation have' where the values are 'Medication efficacy', 'Medication transporter', 'Medication Metabolism', 'High Risk'- 'Potential Adverse Drug event.'

Second layer are the components themselves with

component.value providing a more granular statement. The component.code is used to indicate which type of implication the component.value is for (e.g. 'Medication efficacy' in component.code with component.value of 'Intermediate metabolizer')

# Clinical Genomics (CG) Report and bundling – binding it all together

- The CG report structure has the variants, linked to implications, the implications linked to variants
- Task-med-chgs linked to implications and are included in the report as links for <u>RecommendedAction</u>
  - The implications do not have a link to task-med-chgs: so rev\_include is currently necessary to find the medicine implication statements (guideline, as an example, links are in the med implication statements)

#### Implementation - Receiving

- Will need to capture the Medication implication profile 1 profile to implement
- Each component may need to be handled separately
  - Not an increase in complexity the profile based is just as complex

### Implementation – Querying Efficiency

- The choice of components to handle the complexity makes querying easier. Semantically saying that the components are elements of an observation of a medication implication.
- Example query give me all implications for a patient : 'Observation where observation.code is for Medication Implication and patient ID is X'
- With profiles the query is more complex and would have required each observation.code to be enumerated to find all the profiles. 'Observation where observation.code is either Med Efficacy, Med ...and patient ID is X'
- The component approach is less logically complex requires less programming for system providing the data and the system querying for the data, also requires less processing at run-time.
- Getting a specific type of medication implication is possible through the observation.value E.g. 'Observation where observation.code is for Medication Implication and observsation.value is Medication Efficacy' would provide all the statements on medication efficacy.
- Note: task statements (e.g. lower, raise dose), supporting guidelines and variants are connected in the same manner whether components or profiles are used.

#### Implementation – Sending example

- Variant information is found through links to the correct variant observations.
- Molecular Phenotype example with Predicted Adverse Drug Response
  - "text interpretation": "Individuals with intermediate metabolizer status (IM) have reduced metabolism of tricyclics to less active compounds when compared to extensive metabolizers; the resultant higher plasma concentrations will increase the probability of side effects (concernCode =!A). Consider reducing the dose by 40% and monitor nortriptyline 10-hydroxynortriptyline plasma concentrations (list of references: 8, 26, CPIC guideline)."
  - "text activity": "Intermediate metabolizer. Two alleles showing decreased activity"
  - Confidence = '4'

## Implementation – Sending (placing data)

Molecular Phenotype example with Predicted Adverse Drug Response

From example	Where placed in Clinical Genomics IG
"text interpretation": "Individuals with intermediate metabolizer status have reduced metabolism of tricyclics to less active compounds when compared to extensive metabolizers; the resultant higher plasma concentrations will increase	
the probability of side effects."	Narrative of Medication Implication profile instance
"Consider reducing the dose by 40% and monitor nortriptyline 10-	
hydroxynortriptyline plasma concentrations."	Narrative in task-med-chg
"text activity": "Intermediate metabolizer. Two alleles showing decreased activity"	Narrative in medication metabolism component
metabolizer status code - "IM"	intermediate metabolizer goes in value of 'Medication Metabolism' component.value
Confidence = '4'	evidenceLevel component with PharmGKB as value.code-system and value = Level 1B High
"increase the probability of side effects", concernCode =!A	adverse Drug Event – use 'High Risk Allele' component with 'High' as value
list of references: 8, 26, CPIC guideline	in relatedArtifact of Medication Implication instance
"Consider reducing the dose"	in task-med-chg with 'proposed' intent

#### Future

- New slice on component can be made
- Additional values can be used in Observation.value
  - Dangerous Blessing: flexibility to accommodate future changes but can be misused
- As always, the implementer needs to use appropriate discretion. Make use of the text narrative section to provide human readable summaries. Have a clinical expert review the statements. But beware, computers need more boxes than humans. The human mind contains a model that abstracts complexity in a way not achievable with current computation. The computer needs codable concepts and labeled data elements.

# The Nots – guidelines, knowledge resources, and Clinical Decision Support (CDS) delivery

- Guidelines, Knowledge Resources and Clinical Decision Support delivery should be targets for our WG in conjunction with other working groups (and a formal invitation given to genomic bodies) depending on topic.
- Knowledge resources can be found in locations such as NCBI's repository or PharmGKB documentation. For example, using the HL7 Infobutton standard as implemented by PharmGKB, <a href="https://api.pharmgkb.org/v1/infobutton?mainSearchCriteria.v.dn=Clopidogrel">https://api.pharmgkb.org/v1/infobutton?mainSearchCriteria.v.dn=Clopidogrel</a> delivers a multitude of knowledge on *clopidogrel* in the context of pharmacogenomics. (aside: do we really need to create a standard for PharmGKB beyond recommending the use of the HL7 Infobutton Standard? Also, why is knowledge needing to be given in the message? Is a link sufficient?)
- My suggestion is to work on these questions with the Clinical Decision Support working group. Get a document together with the elements we think are important together and then ask the CDS working group to devise a means to communicate.

# Related Topic Medication assessed - consideration

- Link to medication resource. We did not do this currently. Suggested as we need a slot for, essentially, a generic.
  - Using RxNorm as codeable concept, assumes systems will be able to traverse from RxNorm to the meds on their system. Or requires human to jump from Generic to brand etc...
  - And Drug Class is not communicated so I propose
    - ADD TO OUR MEDICATION assessed medicineClassification 0..\*

- 🛅 medicineClassification	0*	BackboneElement	Categorization of the medication within a formulary or classification system
- 🧊 type	11	CodeableConcept	The type of category for the medication (for example, therapeutic classification, therapeutic sub-classification)
<ul> <li>Classification</li> </ul>	0*	CodeableConcept	Specific category assigned to the medication

- Or use medication knowledge to provide the medication assessed using Reference.
- Ultimately, the CG WG does not have purview over Medication specific knowledge. Expect medication
  assessed to mature as other portions of FHIR mature. BIG CAVEAT *Drug Class could be considered beyond
  our scope.*
- <u>https://www.hl7.org/fhir/medication.html</u> one reason Medication resource is not used is to avoid confusion with actual medications the patient has. However, I think we could use the resource can be used.

## https://loinc.org/61357-0/

#### 61357-0 Medication pharmaceutical advice.brief DocumentActive

Component

Medication pharmaceutical advice.brief

Property

Find

Time

Pt

System

**^Patient** 

Scale

Doc

Method

**Additional Names** 

Short Name

#### Medication pharm advice.brief Doc

#### **Term Description**

This collection contains information about a) the prescription item this pharmaceutical advice is related to, b) concerns such as interactions, contraindications, and allergies, and c) changed or alternative medication information. This term was originally created for, though not limited in use to, the IHE Pharmaceutical Advice profile in the Community Medication Prescription and Dispense set. Source: Regenstrief LOINC