

2016-03-30 - SLPG Meeting

Date & Time

Wednesday 30th March 2016, 20:00 UTC

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Attendees

- Chair: [Linda Bird](#)
- Project Group: [Michael Lawley](#) [Ed Cheetham](#) [Alejandro Lopez Osornio](#) [Guillermo Reynoso](#)

Goals

To introduce the SPLG Confluence Space.

To discuss recent feedback about the SNOMED CT Expression Constraint Language.

To progress the SNOMED CT Template Syntax.

Apologies

- [Daniel Karlsson](#)
- [Harold Solbrig](#)
- [Rob Hausam](#)

Observers

Agenda and Meeting Notes

Item	Description	Owner	Notes	Action
1	Welcome, introductions and apologies	Linda Bird	SLPG meetings will be recorded and recordings will be accessible to SLPG members	<input checked="" type="checkbox"/> Check attendance details and apologies
2	Agenda review	Linda Bird	Review proposed agenda for today's meeting <ul style="list-style-type: none">• Introducing the SLPG Confluence Space• Discuss recent feedback about the Expression Constraint Language<ul style="list-style-type: none">◦ Immediate children/parents◦ Comments• Review template syntax discussion<ul style="list-style-type: none">◦ Scope and purpose of syntax◦ Simplifications to Finding context example◦ Cardinality in templates◦ Populating a template from a data structure	<input checked="" type="checkbox"/> Review agenda
3	SPLG Confluence Space	Linda Bird	Make sure everyone has access to Confluence and the SPLG space	<input checked="" type="checkbox"/> Introduce SPLG space <input checked="" type="checkbox"/> Make sure everyone has access to SPLG space
4	SNOMED CT Expression Constraint Language	Linda Bird	Discuss recent feedback about the ECL 1. Immediate children/parents <ul style="list-style-type: none">• FHIR Community have use case for these operators (user interface display)• Are these just 'syntactic sugar' for<ul style="list-style-type: none">▪ < 404684003 Clinical finding : 116680003 is a = 404684003 Clinical finding ▪ > 404684003 Clinical finding : R 116680003 is a = 404684003 Clinical finding • What syntax would we use - for example:<ul style="list-style-type: none">▪ <! 404684003 Clinical finding ▪ >! 404684003 Clinical finding 	<input checked="" type="checkbox"/> Discuss ECL feedback <input checked="" type="checkbox"/> Linda Bird to draft updates to ECL specification
			2. Comments <ul style="list-style-type: none">• If we introduce comments into the ECL, should we also be introducing them into CG?• Should comments be part of the normative (brief) syntax included in interoperable sharing of ECs?• Or should comments be included in the non-normative (full) syntax?• If we introduce comments, what syntax should we use? For example:<ul style="list-style-type: none">◦ /* */	
			OUTCOMES 1. The group agreed to add new operators to the ECL specification to support the FHIR community's use case for immediate children/parents. The preferred syntax was "<!" and ">!". 2. The group decided that comments should be added to the the brief syntax of the ECL using the "/*... */" syntax.	

5	SNOMED CT Template Syntax	Linda Bird	<p>Review discussion on optionality and populating attribute groups:</p> <ol style="list-style-type: none"> Scope and purpose of syntax <ol style="list-style-type: none"> Extract/disentangle SNOMED CT (and SNOMED CT-relevant) content from a FHIR Condition resource (i) into a free-standing and 'recognisable' SNOMED CT expression, whilst (ii) 'leaving nothing behind' which may be of relevance to further processing Specify mappings from FHIR value sets (e.g. Condition.clinicalStatus) into SNOMED CT Transform the extracted expression into an 'optimally-processable' SNOMED CT expression (in particular grouping body site values with morphology) Specify constraints on what the extracted/disentangled SNOMED CT expression could or couldn't contain (by e.g. cardinality instructions). <p>NOTE - The rest of this discussion will be carried over until the next meeting.</p> <ol style="list-style-type: none"> (From a(ii) and b above) Simplify [finding context] refinement to either: <ul style="list-style-type: none"> 408729009 [finding context] = [[@findingContext]] 408729009 [finding context] = [[findingContextTable (\$clinicalStatus, \$verificationStatus)]] (From d above) How to specify cardinality in terminology binding when restricting valid values in an information model data element: <ul style="list-style-type: none"> 62014003 [Adverse reaction to drug (disorder); 246075003 [Causative agent] = [[[0..1]^ 111115 AMP reference set]] 62014003 [Adverse reaction to drug (disorder); !! [0..1] !! 246075003 [Causative agent] = [[^ 111115 AMP reference set]] (From c above) To indicate how the following data structure can be used to populate a template: <ul style="list-style-type: none"> Data Structure A <ul style="list-style-type: none"> Condition <ul style="list-style-type: none"> Code: CodeableConcept [0..1] MorphologyBS [0..*] <ul style="list-style-type: none"> BodySite: CodeableConcept [0..1] Morphology: CodeableConcept [0..1] Possible template syntax examples: <ul style="list-style-type: none"> [[\$code]]: { 363698007 [finding site] = [[\$BodySite]], 116676008 [associated morphology] = [[\$Morphology]] [[\$code]]: { 363698007 [finding site] = [[\$MorphologyBS.BodySite]], 116676008 [associated morphology] = [[\$MorphologyBS.Morphology]] [[\$code]]: !! For each M = \$MorphologyBS !! { 363698007 [finding site] = [[M.BodySite]], 116676008 [associated morphology] = [[M.Morphology]] Data Structure B <ul style="list-style-type: none"> Condition <ul style="list-style-type: none"> Code: CodeableConcept [0..1] BodySite: CodeableConcept [0..*] Morphology: CodeableConcept [0..1] Possible template syntax examples: <ul style="list-style-type: none"> To include the different finding sites <i>within</i> the same attribute group: <ul style="list-style-type: none"> [[\$code]]: { 363698007 [finding site] = [[\$BodySite]], 116676008 [associated morphology] = [[\$Morphology]] To include an attribute group for each <i>different</i> finding site (with the <i>same</i> associated morphology): <ul style="list-style-type: none"> [[\$code]]: !! For each BS = \$BodySite !! { 363698007 [finding site] = [[BS]], 116676008 [associated morphology] = [[\$Morphology]] Other examples discussed by email (double scope): <ul style="list-style-type: none"> [finding] : [[([0..*] [findingSite] = \$bodySite << 48566001 Bone structure of extremity (body structure) , [[[0..1] [assocMorph] = \$morphology < 72704001 Fracture (morphologic abnormality)]])]] 	<input checked="" type="checkbox"/> Review Template Syntax discussion
6	Confirm next meeting date/time	Linda Bird	Confirm date and time of next SLPG meeting - Wednesday 27th April	<input checked="" type="checkbox"/> Confirm date of next call

Meeting Files

File Modified

No files shared here yet.