Genomics informatics interoperability standards supporting precision medicine across health IT systems

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SNOMED International Genomics and Precision Medicine Clinical Reference Group April 2019 Business Meeting Genomics Session, 11th April 2019



"We believe that genomic data and the genomic aspects of precision medicine should be incorporated into the routine clinical workflow in as seamless and transparent a fashion as possible."

David McCallie Jr., Cerner Senior Vice President of Medical Informatics

genomeweb, Cerner Banking on Third-Party Apps for Genomic Integration, Starting with PGx, 28th Dec 2018



Introduction

Roles

- Associate Vice Chair for Standards, British Computer Society (BCS) Health and Care Group
- BCS representative, British Standards Institute (BSI) IST/35 Health Informatics Committee
- Member at large, Health Level Seven (HL7) UK Management Board
- · Member, IHE UK Steering Committee
- · Vendor representative, INTEROPen Board



Introduction

Agenda

- · Health IT vendor perspective
- · Standards landscape
- · Challenges and possible ways forward





Cerner overview



As of Q2 2018

735 ACUTE CLIENTS **66 1,600** AMBULATORY CLIENTS **471**

Updated 09/2018

FACILITIES

in 35+ COUNTRIES



Solution portfolio

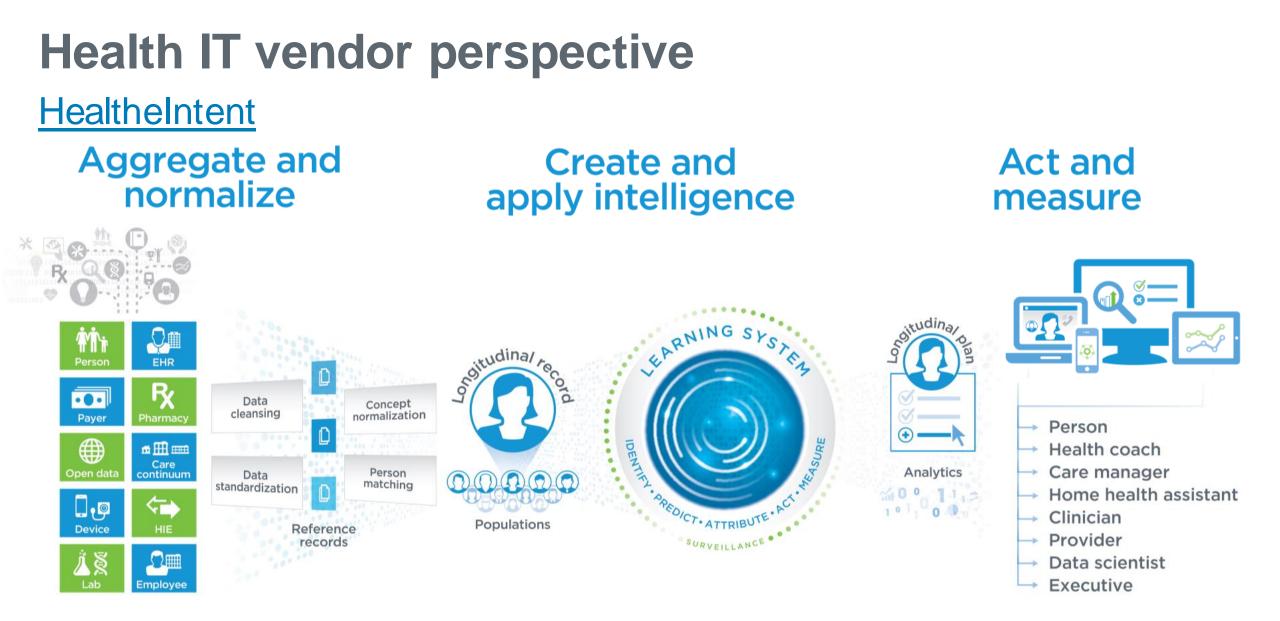


3 PLATFORMS • 1 ARCHITECTURE • 1 EXPERIENCE

Enable high performance organization • Lower total cost of ownership • Data liquidity



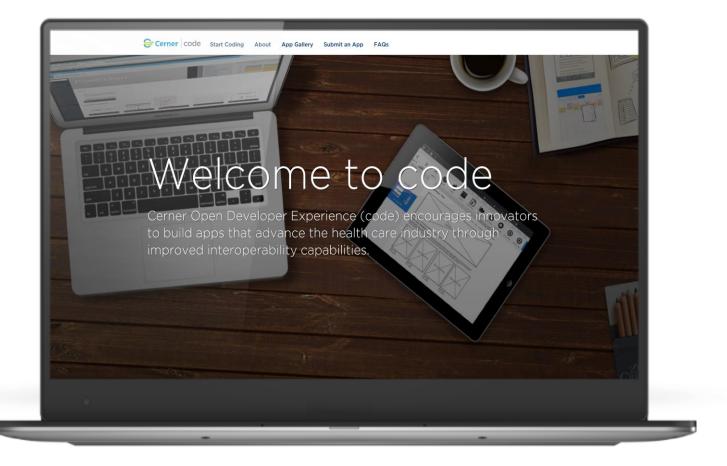






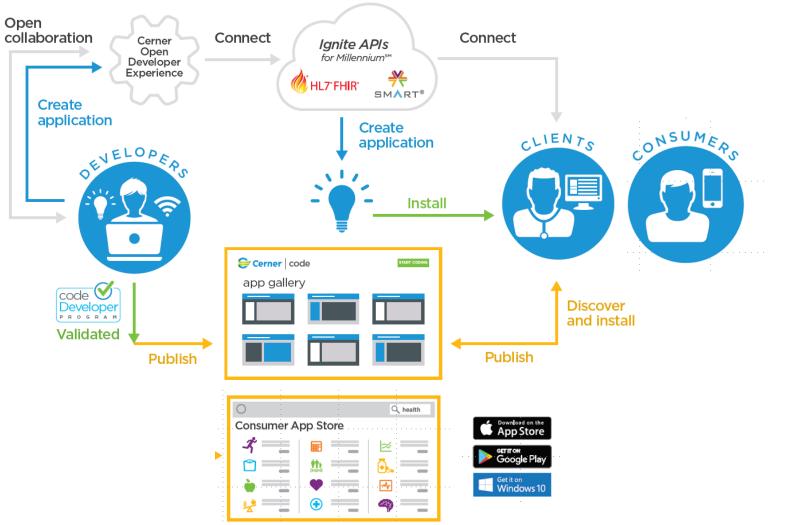
Open Platforms

- Cerner Open Developer Experience (code)
 - https://code.cerner.com/
- Open APIs Ignite
 - EHR
 - https://fhir.cerner.com/
 - Population Health
 - https://docs.healtheintent.com/





Open Platforms

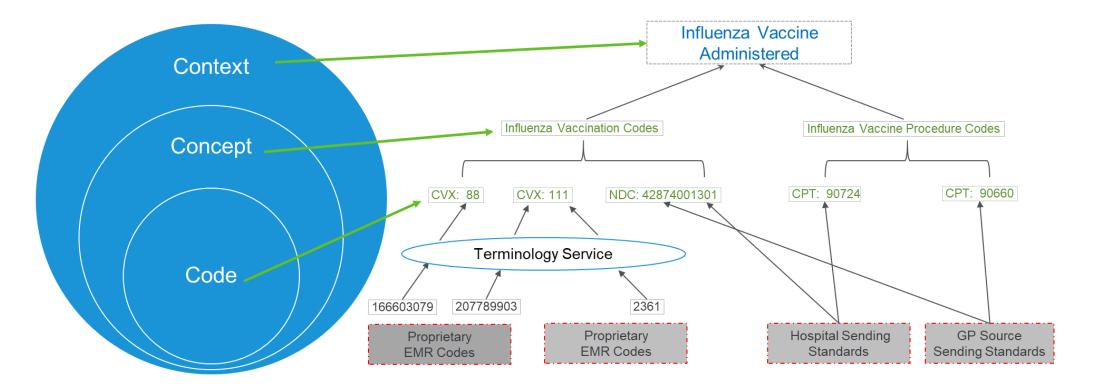




- Concept normalisation for population health and EHR FHIR API concept mapping
- Supports rules engine that drives registries
- <u>https://docs.healtheintent.com/api/v1/ontology/</u>



Discern Ontology



Standardisation and Normalisation Example: Influenza Vaccine

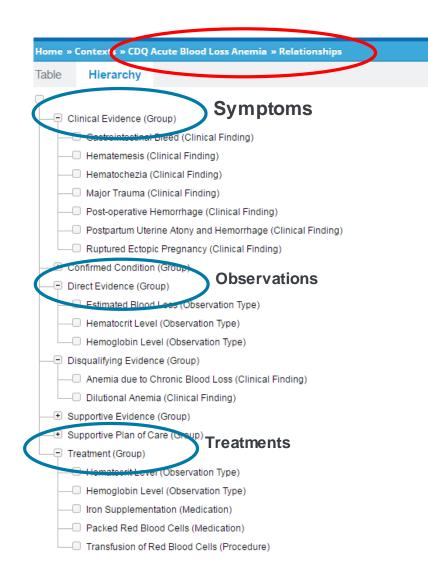


CDQ Acute Blood Loss Anemia Maybe(0)										
Home » Contex s » CDQ Acute Blood Loss Anemia » Concepts										
Context Con	cepts Templates Code Systems	Relationships Releases								
New Concept										
Туре	Name	♦ Aliases	Aaybes Upgra	adesJpdater	Updated (GMT)					
T										
Clinical Finding Group Medication	Acute Blood Loss Anemia	ACUTE_BLOOD_LOSS_ANEMIA	0 1	Peggy Payne	2016-10-17 22:22					
	Acute Postoperative Anemia	ACUTE_POSTOPERATIVE_ANEMIA	0 1	Peggy Payne	2016-10-17 22:26					
Observation Type	Anemia due to Chronic Blood Loss	ANEMIA_DUE_TO_CHRONIC_BLOOD_LOSS	0 1	Peggy Payne	2016-10-18 20:20					
Procedure	Dilutional Anemia	DILUTIONAL_ANEMIA	0 1	Peggy Payne	2016-10-17 22:43					
Clinical Finding	Dizziness	DIZZINESS	0 0	Peggy Payne	2016-10-17 22:48					
Clinical Finding	Fatigue	FATIGUE	0 0	Peggy Payne	2016-10-26 18:46					
Clinical Finding	Gastrointestinal Bleed	GASTROINTESTINAL_BLEED	0 1	Peggy Payne	2016-10-18 19:15					
Clinical Finding	Generalized Weakness	GENERALIZED_WEAKNESS	0 1	Peggy Payne	2016-10-18 19:52					
Clinical Finding	Hematemesis	HEMATEMESIS	0 0	Candy Barth	2015-04-07 13:55					
Clinical Finding	Hematochezia	HEMATOCHEZIA	0 0	Peggy Payne	2016-08-10 20:47					
Clinical Finding	Major Trauma	MAJOR TRAUMA	0 4	Peggy Payne	2016-10-27 17:49					



Home » Contexts » CDQ Acute	Blood Loss	Anemia » Concept » Hemoglobia	n Level [Observation Type	e] » Codes	
Concept Templates	Codes	Add a Code Children			
Save Cancel					
Code System	Code	Sources	CUI(s)	Ame(s)	Templates
			•		
CHV CPT (HCPCS Level I) Clinical Problem Statements ICPC-2 PLUS	14775-1	LNC	C0797949	Hgb BldA Oximetry-mCnc Hemoglobin:MCnc:Pt:BldA:Qn:Oximetry Hemoglobin [Mass/volume] in Arterial blood by Oxi Hemoglobin:Mass Concentration:Point in time:Bloc	
LOINC MEDCIN MedDRA Medical Entities Dictionary	20509-6	LNC	C0803320	Hgb Bld Calc-mCnc Hemoglobin:MCnc:Pt:Bld:Qn:Calculated Hemoglobin [Mass/volume] in Blood by calculation Hemoglobin:Mass Concentration:Point in time:Who	
NCI Read Codes v3 SNOMED CT	30313-1	LNC	C1114184	Hgb BldA-mCnc Hemoglobin:MCnc:Pt:BldA:Qn Hemoglobin [Mass/volume] in Arterial blood Hemoglobin:Mass Concentration:Point in time:Bloo	Hemoglobin Level
LOINC	30350-3	LNC	C1114213	Hgb BldV-mCnc Hemoglobin:MCnc:Pt:BldV:Qn Hemoglobin [Mass/volume] in Venous blood Hemoglobin:Mass Concentration:Point in time:Bloo	Hemoglobin Level
LOINC	20251_1	LNC	01114214	Hgb BldMV-mCnc Hemoglobin:MCnc:Pt:BldMV:Qn	Hemoglobin Level







Cerner Associates driving Genomics

- Kevin Power, Director of Genomics Solutions and Co-Chair on the HL7 Clinical Genomics Work Group
- Terah Collins, Senior Strategist for Precision Medicine and Genomics
- Marc Overhage, Chief Medical Informatics Officer
 & VP Intelligence Strategy



Genomic Precision Medicine related health IT system contexts

- Electronic Health Record (EHR)
- · Consumer
- Laboratory Information Management Systems (LIMS)
- · Population Health
- · Research



Genomic Precision Medicine client priorities

- · Pharmacogenomics
 - Oncology
 - Cardiovascular
 - Mental Health
 - Pain Management
- · Oncology
 - Somatic sequencing
- · Newborn screening
- · Rare and undiagnosed disease



- · Millennium
 - Model Experience -> Precision Medicine -> Pharmacogenomics
 - Decision Support Rules
 - High Risk Genotype
 - Drug-metabolizing enzymes
 - Transporters
 - Ignite API for Millennium
 - YouScript
 - https://code.cerner.com/en/apps/youscript
 - MyLegacy by Family Care Path
 - https://code.cerner.com/apps/mylegacy



Consumer

- · Millennium
 - · Ignite API for Millennium
 - Apple Health
 - https://code.cerner.com/apps/apple-health
 - MyLegacy by Family Care Path
 - https://code.cerner.com/apps/mylegacy
 - · HealtheLife, Patient Portal
- <u>CommonWell Members Enable Patient Access to Their Health Data</u>
- All of Us Research Program to Test Options for Returning Results

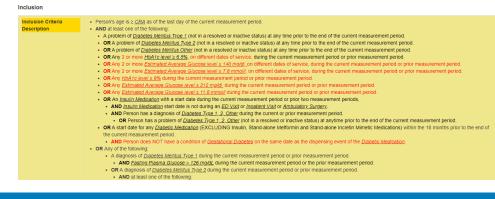


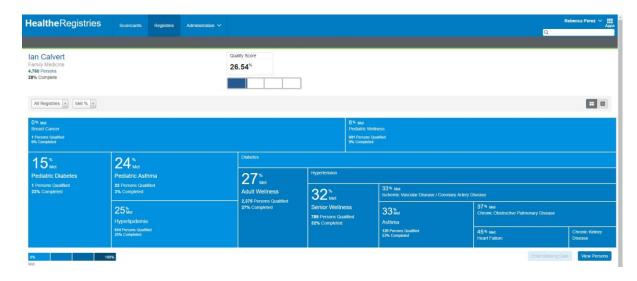
- · LabSequence
 - Workflow engine Polymerase Chain Reaction (PCR), Microarray, Multiplex Ligation-dependent Probe Amplification (MLPA), Next Generation Sequencing (NGS) etc
 - API available for customising and integrating with devices and software

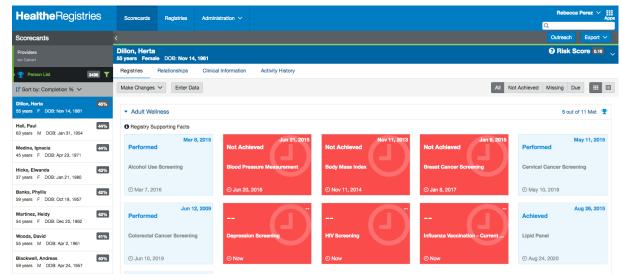


Population Health

- · HealtheIntent
 - · HealtheRegistries
 - Using Cerner
 'Synapse' Domain
 Specific Language and
 Rules Engine (using
 Drools and Clojure)



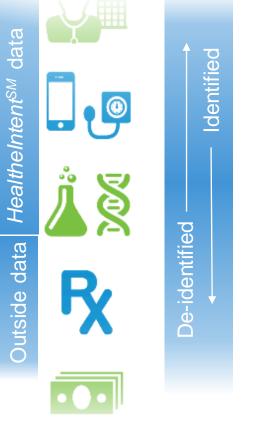






Research

- HealtheIntent
 - HealtheDataLab





Physical storage & processing

web services S3

Role-based access control



User tools

Client initiatives

- The Alliance for Genomics (TAG)
 - Pete Celano, Director of Consumer Health Initiatives, MedStar Health, MedStar Institute of Innovation
 - <u>https://www.cerner.com/blog/ep-100-future-of-precision-medicine</u>

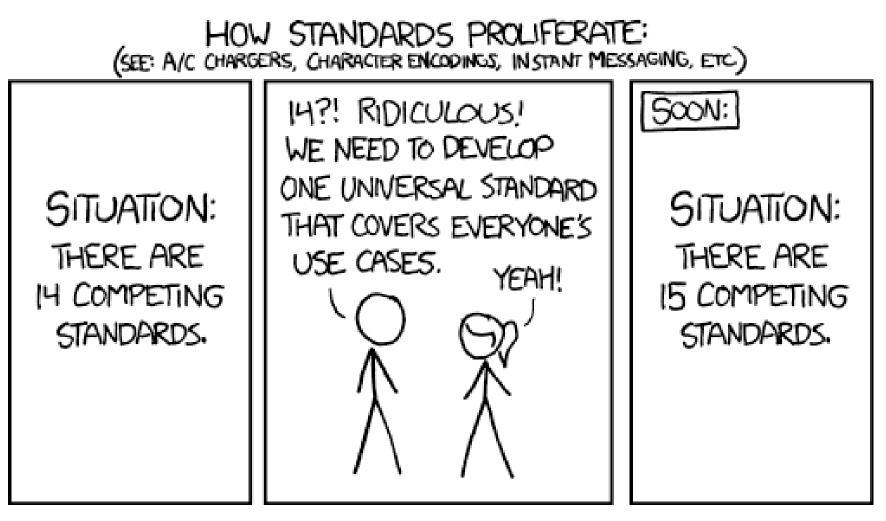


Client initiatives

- The Alliance for Genomics (TAG)
 - Presentations to date
 - Illumina
 - YouScript
 - Dignity Health and Catholic Health Initiatives, Precision Medicine Alliance LLC
 - Million Veteran Program
 - HudsonAlpha Institute for Biotechnology
 - American Heart Association, Institute for Precision Cardiovascular Medicine
 - <u>Helix</u>
 - <u>Myriad</u>
 - <u>AWS</u>
 - Cerner
 - Pharmacogenomics
 - HealtheDataLab











ISO/TC 215 Health Informatics Committee Proposal

- ISO/TC 215 N 2776 Proposal for Subcommittee on "Genomics informatics"
 - Proposed liaison organisations and survey of similar work
 - Other ISO Committees
 - ISO/TC 276 Biotechnology
 - ISO/IEC JTC 1/SC29 Coding of audio, picture, multimedia and hypermedia information
 - <u>Global Alliance for Genomics and Health (GA4GH)</u> and driver projects
 - HL7 International Clinical Genomics Working Group
 - <u>Clinical Data Interchange Standards Consortium (CDISC)</u>
 - <u>SNOMED International</u>
 - <u>Regenstrief Logical Observation Identifiers Names and Codes (LOINC)</u>
 - <u>National Cancer Institute Enterprise Vocabulary Services NCI Thesaurus (NCIt)</u>
 - <u>Massive Analysis and Quality Control (MAQC) Society</u>



ISO/TC 215 Health Informatics Committee Proposal

- ISO/TC215 SC1 Genomic Informatics proposed initial active programme of work
 - ISO/IS 25720:2009 Health Informatics Genomic sequence variation markup language (GSVML)
 - ISO/TS 20428:2017 Health Informatics Data elements and their metadata for describing structured clinical genomic sequence information in electronic health records
 - ISO/CD 21393 Health Informatics Omics Markup Language (OML) (in development)
 - ISO/NP TS 23357 Clinical genomics data sharing specification for next generation sequencing (in development)
 - ISO/AWI TR 21394 Health Informatics Whole genome sequence markup language (WGML) (in development)
 - ISO/AWI TR 22693 Health Informatics Structured clinical gene fusion report in electronic health records (in development)
 - ISO/PWI 22690 Health Informatics Reliability assessment criteria for high-throughput gene-expression data (in development)
 - ISO/DTS 22692 Health Informatics Quality control metrics for DNA sequencing (in development)



GA4GH Work Streams

- Key Technical Work Streams
 - Genomics Knowledge Standards
 - <u>Clinical and Phenotypic Data Capture</u>
- Other Work Streams
 - Foundational
 - Data Security
 - Regulatory & Ethics
 - Technical
 - Cloud
 - Data Use & Researcher Idenities (DURI)
 - Discovery
 - Large Scale Genomics



GA4GH Driver Projects

- · Key projects
 - Monarch Initiative
 - Human Phenotype Ontology (HPO)
 - <u>Clinical Genome Resource (ClinGen)</u>
 - Variant Interpretation for Cancer Consortium (VICC)



GA4GH Driver Projects

- · Other
 - All of Us Research Program
 - Australian Genomics
 - Autism Sharing Initiative
 - BRCA Challenge
 - <u>Canadian Distributed Infrastructure for Genomics (CanDIG)</u>
 - ELIXIR Beacon
 - <u>European Genome-phenome Archive (EGA)</u>, <u>European Variation Archive (EVA)</u>, and <u>European Nucleotide Archive (ENA)</u>
 - National Institute of Environmental Health Science (NIEH) EpiShare
 - European-Canadian Cancer Network (EUCANCan)
 - European Joint Programme on Rare Disease (EJP RD)
 - GEnome Medical alliance Japan (GEM Japan)



HL7 International Clinical Genomics Working Group

- <u>Confluence Work Group Home</u>
- <u>Wiki</u>
- <u>Existing Products</u>, v2, v3, CDA Implementation Guides Domain Analysis Models
- FHIR Implementation Guides
 - Genomics Implementation Guidance
 - <u>Genetic Reporting Implementation Guide</u> (Jan 2019 v0.3.0: STU 1 Ballot
 2)
 - New MolecularSequence resource plus DiagnosticReport and Observation resource profiles
 - General Genomic Reporting, Variant Reporting, Cytogenic Reporting, Pharacogenomic Reporting, Histocompatability Reporting



CDISC

 <u>Study Data Tabulation Model Implementation Guide</u> (SDTMIG)-Pharmacogenomics/Genetics (PGx) v1.0



SNOMED International

- Genomics and Precision Medicine Clinical Reference Group
- Seminars
 - Genomics Symposium (Ontologies for Clinical Value) -London, 12th, April 2018
 - "Making it real" Genomics session, Vancouver, 17th, October 2018
- Observable and Investigation Model Project Home
- SNOMED on FHIR



LOINC

- Standardised terms for genetic test ordering and reporting, and genetic findings (observations about the specimen's genetic characteristics)
- Links between LOINC and other genetic terminologies and coding systems
- https://loinc.org/genetics/ coming soon
- Deckard J, et al. J Am Med Inform Assoc 2015;22:621–627.
 Supporting interoperability of genetic data with LOINC



NCI

- National Cancer Institute Enterprise Vocabulary Services <u>NCI Thesaurus (NCIt)</u>
 - Provides reference terminology for many NCI and other systems. Covers vocabulary for clinical care, translational and basic research, and public information and administrative activities.
 - Features:
 - Stable, unique codes for biomedical concepts;
 - Preferred terms, synonyms, research codes, external source codes, and other information;
 - Over 100,000 textual definitions;
 - Links to NCI Metathesaurus and other information sources;
 - Over 400,000 cross-links between concepts, providing formal logic-based definition of many concepts;
 - Extensive content integrated from NCI and other partners, much available as separate NCIt subsets
 - Updated frequently by a team of subject matter experts.



Other related organisations and initiatives

- Online Mendelian Inheritance in Man (OMIM)
- Institut National de la Santé et de la Recherche Médicale (INSERM) Orphanet
- European Bioinformatics Initiative (EBI)
 - Other related <u>EBI initiatives</u> not referenced by ISO proposal, ChEMBL etc
- <u>University of Maryland School of Medicine Institute for Genome Science</u> <u>Disease Ontology (DO)</u>
- Alliance of Genome Resources Gene Ontology (GO)
- Basic Formal Ontology (BFO)
 - From which GO, HPO, DO, Information Artefact Ontology (IAO) among others are derived
- Open Biological and Biomedical Ontology (OBO) Foundry



Indirectly related organisations and initiatives

• HL7

- FHIR
 - Other pertinent <u>resources</u>
 - Phenotype
 - Patient, Condition (Diagnosis, Problem), Observation, Allergy etc
 - FamilyHistory
 - Interventions
 - MedicationRequest and related, Procedure, CarePlan, ServiceRequest etc
 - <u>Security</u>
 - Authenticatiion and Authorisation, Consent, Provenance, AuditEvent



Indirectly related organisations and initiatives

- HL7 continued
 - <u>Clinical Quality Language (CQL)</u>
 - FHIRPath
 - <u>Terminology Service</u>
 - Bulk data
 - Regional Implementation Guides, <u>Argonaut Data Query</u>, <u>CareConnect Core API</u>, <u>other</u>
- GraphQL (Using GraphQL with FHIR)
- Grading of Recommendations Assessment, Development and Evaluation (GRADE) working group
- MAGIC
- <u>Observational Health Sciences Data Science and Informatics (OHDSI) OMOP Data</u> <u>Model</u>



Indirectly related organisations and initiatives

- SMART Health IT
 - <u>SMART on FHIR</u> (in collaboration with HL7)
 - <u>CDS Hooks (in collaboration with HL7)</u>
 - Sync for Science (S4S)



Managing hypertension? <u>Launch JNC 8 Rx Pro</u>

Indirectly related organisations and initiatives

· CDS Hooks 1 EHR Med Order EHR triggers a CDS hook and **CDS** invokes a remote service **CDS Service** executes 2 Toprol XL **R**50 mg daily Services its own rules, leveraging FHIR data as needed information card \$200 per month 3 (patient pays \$30) EHR Returns CDS cards suggestion card (rendered and displayed by EHR) **FHIR Server** Try HCTZ as first-line Switch to HCTZ smart app link card



Indirectly related organisations and initiatives · ISO

- ISO/IEC JTC 1/SC 32 Data management and interchange
 - ISO/IEC 11179-1:2015 Information technology -- Metadata registries (MDR)
 - ISO/IEC 11179-1:2015 Part 1: Framework
 - ISO/IEC 11179-2:2005 Part 2: Classification
 - ISO/IECTR 11179-2 Inform Part 2: Classification (Under development)
 - ISO/IEC 11179-3:2013 Part 3: Registry metamodel and basic attributes
 - ISO/IEC 11179-3:2013/DAmd 1 (Under development)
 - ISO/IEC 11179-4:2004 Part 4: Formulation of data definitions
 - ISO/IEC 11179-5:2015 Part 5: Naming principles
 - ISO/IEC 11179-6:2015 Part 6: Registration
 - ISO/IEC DIS 11179-7 Part 7: Metamodel for data set registration (Under development)
 - Metadataworks
 - Genomics England
 - <u>Model Driven Data Management in Healthcare</u>' David Milward, Best Paper award at MODELSWARD 2019 International Conference on Model-Driven Engineering and Software Development
 - Metadata Management Language (MDML)



Indirectly related organisations and initiatives · ISO

- Identification of Medicinal Products (IDMP)
 - FHIR <u>MedicationKnowledge</u> resource
- ISO 13606 Health informatics -- Electronic health record communication
 - <u>openEHR</u>





Challenges

- Knowledge representation and reasoning conundrum
- Authentication and authorisation
- Ethics



Knowledge representation and reasoning conundrum

- Knowledge? Classifications/terminologies, ontologies, Domain Specific Languages (DSLs), information models...
- Relationship between classifications/terminologies and ontologies
- Relationship between different ontologies
- Relationship between information model representations of EHR, classifications/terminologies and ontologies
- Reasoning



Relationship between classifications/terminologies and ontologies

- Examples
 - ICD10 -> ICD11 <-> SNOMED CT alignment
 - LOINC <-> SNOMED CT



Relationship between different ontologies

- BFO as key component in alignment efforts
 - Alignment of SNOMED with BFO
 - Stakeholder Engagement for Consultant Terminologist Project <u>IHTSDO-836, entitled "Concept Model: Presence" David</u> <u>Sperzel, MD, MS, August-September 2016</u>
 - SNOMED CT and formal ontological principles p.17
 - <u>SNOMED CT standard ontology based on the ontology for</u> <u>general medical science EI-Sappagh et al. BMC Medical</u> <u>Informatics and Decision Making (2018) 18:76</u>



Relationship between information model representations of EHR, classifications/terminologies and ontologies

· ZOUD

- Standards based EHR information models
 - FHIR
 - · ISO 13606 / openEHR
 - INTEROPen Digital Health Rewired Panel Paper: FHIR and openEHR
- Articulating relationships
 - BFO and IAO as possible component in alignment efforts
 - FHIR and Ontology Barry Smith, Fourth Clinical and Translational Science Ontology Workshop 2015
 - <u>https://wolandscat.net/tag/ontologies/ https://wolandscat.net/tag/ontology/</u>
 - http://yosemiteproject.org/



Reasoning

- Rector AL, Rogers J, Taweel A. Models and inference methods for clinical systems: a principled approach. Stud Health Technol Inform. 2004;107(Pt 1):79–83
- · Workflow, CDS Hooks
- · DSLs, CQL
 - Beyond the Curly Braces: Exploring CQL on FHIR based CBK, Mobilizing Computable Biomedical Knowledge (MCBK) Marc Overhage with Motive Medical Intelligence, July 2018
- Rules engines/reasoners

Authentication and authorisation

- SMART on FHIR
- FHIR Security



- NIH National Human Genome Resource Institute
 - Ethical, Legal and Social Issues in Genomic Medicine
 - Genetic Discrimination
 - Intellectual Property and Genomics
 - Regulation of Genetic Tests
 - Privacy in Genomics
- Knowledge sharing and resource distribution



Pragmatism

The pragmatic maxim

Consider the practical effects of the objects of your conception. Then, your conception of those effects is the whole of your conception of the object.

Charles Sanders Peirce

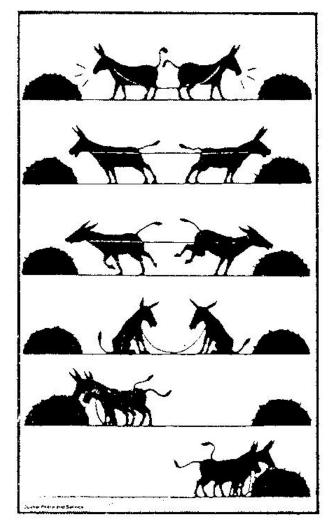


Continued collaboration...

- aligning classifications/terminologies with ontologies
- aligning different ontologies, with SNOMED, BFO and derivatives being a key components of alignment efforts
- aligning EHR information model representations of, classifications/terminologies and ontologies, with FHIR, SNOMED, BFO and derivatives being key components in alignment efforts
- progressing approaches to reasoning and clinical decision support with CDS Hooks and CQL being key components
- progressing approaches to Authentication and Authorisation with SMART on FHIR and FHIR Security being key components
- progressing ethics
- engaging industry and being pragmatic!



Cooperation is better than conflict!





Thank you.

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