

Genomic Medicine Centers Meeting VII

Genomic Clinical Decision Support – Developing Solutions for Clinical and Research Implementations October 2-3, 2014



National Human Genome Research Institute

Introductions

- Introductions and Welcome:
 - Marc and Blackford
 - Around the room
- Logistics
 - Bathrooms
 - Breaks
 - Overview of Agenda





Meeting Objectives

- GM 7 will convene key thought leaders in genomic implementation and application of clinical decision support to:
 - Compare current state with ideal state of genomic clinical decision support to define gaps and strategies to close the gaps
 - Identify and engage US and international health IT initiatives that would support recommended strategies
 - Define a prioritized research agenda for GCDS





Potential Examples of GCDS

- 1. Medication dosing support
- 2. Order facilitators
- 3. Alerts and reminders

1. Relevant information display

- 2. Expert systems Workflow support
- 3. Clinical genomics example

- CDS automatically adjusts warfarin dosing as a result of known alleles in the VKORC1 and CYP2C9 genes
- An order for colonoscopy is recommended at a younger age as a result of known pathogenic mutations in genes associated with colon cancer
- During medication ordering, gene variants known to affect drug pharmacokinetics are checked and clinicians are alerted to potential gene- drug interactions
- Context aware infobuttons in the problem list leverage genome data to provide genetic risk information for a patient with breast cancer
- The EHR provides a 10-year cardiovascular disease risk score based on clinical, environmental, and genetic risk factors
- The EHR schedules a genetic counseling consultation during prenatal visit due to presence of an X-linked disease gene variant

Our Key GCDS Questions

- 1. Is clinical decision support an essential element in the successful implementation of genomic medicine?
 - Does genomic clinical decision support differ significantly from decision support used for other purposes? Ifyes, what are the key differences?
 - What is the ideal state of genomic clinical decision support?
 - How can the impact of genomic clinical decision support be defined and measured?
- 2. What are data issues that impact genomic CDS?
- 3. How do we manage knowledge for genomic clinical decision support?
- 4. What are implementation issues surrounding genomic CDS?
- 5. What are areas that should be prioritized for the research agenda for GCDS?





GM7 Survey

 Survey instrument based on the 14 key recommendations from Masys *et al,* and Welch *et al.*

- Survey response rate
 - 30 invited attendees
 - 25 responded
 - 83% response rate





Recall the 14 Elements of Masys and Welch

1 Maintain **separation** of primary molecular observations from the clinical interpretations of those data

2 Support **lossless** data compression from primary molecular observations to clinically manageable subsets

3 Maintain linkage of molecular observations to the laboratory **methods** used to generate them

4 Support compact representation of clinically actionable **subsets** for optimal performance

5 Simultaneously support human-viewable formats and machine-readable formats in order to facilitate implementation of decision support rules

6 Anticipate fundamental changes in the understanding of human molecular variation

7 Support both individual clinical care and **discovery** science

8 CDS knowledge must have the potential to incorporate **multiple** genes and clinical information

9 Keep **CDS knowledge** separate from variant classification

10 CDS knowledge must have the capacity to support **multiple EHR** platforms with various data representations with minimal modification

11 Support a large number of **gene variants** while simplifying the CDS knowledge to the extent possible

12 Leverage current and developing CDS and genomics **standards**

13 Support a **CDS knowledge base** deployed at and developed by multiple independent organizations

14 Access and transmit only the genomic information necessary for CDS

Welch, B. M., Eilbeck, K., Fiol, G. D., Meyer, L. J., & Kawamoto, K. (2014). Technical

desiderata for the integration of genomic data with clinical decision support. *Journal of Biomedical Informatics*. doi:10.1016/j.jbi.2014.05.014 Masys, D. R., et al. (2012). Technical desiderata for the integration of genomic data

into Electronic Health Records. Journal of Biomedical Informatics, 45(3), 419–422.

Mean Element Importance



- 1. Separation of clin interp
- 2. Lossless compression
- 3. Methods linkage
- 4. Actionable subsets
- 5. Human /Machine readable
- 6. Changes in understanding
- 7. Discovery science
- 3. CDS over multiple genes
- 9. CDS Knowledge separate
- 10. Multiple EHR
- 11. Support Gene variants
- 12. Standards: CDS and genomics
- 13. Deploy shared CDS KB
- 14. Access and transmit minimum



Mean Difference from Ideal Capability



- 1. Separation of clin interp
- 2. Lossless compression
- 3. Methods linkage
- 4. Actionable subsets
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Mean Importance vs Mean Difference from Ideal



Sum of Priorities Selections by Element



Sum of Priority Selections Across Respondants

- 1. Separation of clin interp
- 2. Lossless compression
- 3. Methods linkage
- 4. Actionable subsets
- 5. Human / Machine readable
- 6. Changes in understanding
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- 8. CDS over multiple genes
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Prioritization Insights from the Survey

From Import v Lo Diff from Ideal

- 1 Maintain separation of primary molecular observations from the clinical interpretations of those data
- 5 Simultaneously support human-viewable formats and machine-readable formats in order to facilitate implementation of decision support rules
 - 6 Anticipate fundamental changes in the understanding of human molecular variation
 - 10 CDS knowledge must have the capacity to support multiple EHR platforms with various data representations with minimal modification
 - 11 Support a large number of gene variants while simplifying the CDS knowledge to the extent possible
- 12 Leverage current and developing CDS and genomics standards
- 13 Support a CDS knowledge base deployed at and developed by multiple independent organizations

From Top 5 Rankings

- 1 Maintain separation of primary molecular observations from the clinical interpretations of those data
- 5 Simultaneously support human-viewable formats and machine-readable formats in order to facilitate implementation of decision support rules
- 8 CDS knowledge must have the potential to incorporate multiple genes and clinical information
- 12 Leverage current and developing CDS and genomics standards
- 13 Support a CDS knowledge base deployed at and developed by multiple independent organizations

KEY THEMES FROM GM7 SURVEY

- Consensus on Masys: 1, 5; Welch: 12, 13
 - Agreement around separation of data and knowledge stores
 - Create machine-, and human-readable knowledge artifacts
 - Leverage current and developing CDS and genomic standards
 - Deploy a shared knowledge-base at several institutions
- Other 6, 8, 10, 11





Panel 1: What are the data issues that impact genomic CDS?

- Moderators: Robert Freimuth, PhD and James Ostell, PhD
 - Relevant Desiderata Elements 1, 2, 9
- Discussion of Key Questions
 - I. What data types are essential for genomic CDS
 - a. Patient Level / Clinical Data?
 - b. Provider / Institutional Data?
 - c. Other?
 - II. How does the massive nature of genomic data influence development and implementation of genomic CDS?
 - III. Are there unique attributes of genomics data that present unique challenges to the development and implementation of genomic clinical decision support?
 - a. Persistent nature of germ-line variation
 - b. Rapidly changing knowledge around genomic variants
 - c. Somatic vs. germline variation





Panel 2: How do we manage knowledge for genomic clinical decision support?

- Moderators: Atul Butte, MD, PhD and Josh Peterson, MD, MPH
 - Relevant Desiderata Elements 4, 5, 6, 8, 11, 13
- Discussion of Key Questions
 - I. What are the necessary elements of knowledge management and representation to achieve ideal state? What standards exist or are needed to achieve ideal state?
 - II. What type of clinical decision support architecture (Wright and Sittig, 2008) is needed to achieve ideal state?
 - III. What governance issues arise in knowledge management?





Panel 3: What are implementation issues surrounding genomic CDS?

- Moderators: Kensaku Kawamoto, MD, PhD, MHS and Casey Overby, PhD
 - Relevant Desiderata Elements 3, 7, 10, 12
- Discussion and Key Questions
 - I. Are there common workflow issues to be considered when implementing GCDS? If so, what are they?
 - II. What are the intra- and inter- institutional data and knowledge exchange issues implementing GCDS at scale?
 - a. Storage
 - b. System Requirements
 - c. Standards
 - d. Security
 - III. What is the role for patient facing genomic clinical decision support?





Panel 4: Discussion – What are areas that should be prioritized for the research agenda for GCDS?

- I. What existing entities (e.g. CPIC informatics, AMIA Genomics working group, CDSC, Health e-Decisions, ONC-HealthIT, Professional Societies, Guideline Developers, etc.) are aligned with the recommended strategies and how can they be engaged?
- II. Avoidance of adverse reactions vs. improvement in health
- III. Quality improvement vs. cost containment



