NHGRI Computational Genomics and Data Science Program Workshop: Overview and Recommendations

> Ajay Pillai on behalf of CGDS Group

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Goals

- Prioritize genomic research topics relevant to the NHGRI extramural Computational Genomics and Data Science (CGDS) program
- Identify current challenges that face the computational genomics and data science community
- Redefine the focus of the CGDS portfolio over the next 3-5 years

NHGRI CGDS Workshop

Meeting Organizers

- Drs. Mike Boehnke, Carol Bult, Trey Ideker, Aviv Regev, Lincoln Stein
- NHGRI CGDS Staff

Participants

- 39 invited extramural researchers (academic and industry)
- NHGRI, NCI, NIGMS, ADDS staff

NHGRI CGDS Workshop

Organization

- Sessions designed by organizing committee
- Speakers selected by small groups of participants

Session Topics

- Challenges in enabling new biology in basic science
- Challenges in enabling new clinical insights
- Data and computational resources
- Computations at scale
- Collaborating with non-NHGRI resources

NHGRI CGDS Workshop

Recommendations from sessions consolidated and refined by participants using dot-storming



- Statistical/computational tools enabling *interactive analysis and visualization of large data sets*
- 2. Methods and data that enhance *understanding of how genotype translates to phenotype*
- Tools, technologies, and policies to *ensure genomic data sharing*
- 4. Statistical/computational tools to *identify causal variants*

- 5. Multi-scale *phenotype-focused ontologies* and standards
- **6.** Efficient and scalable algorithms and methods for computeintensive applications
- 7. Vertically integrated data resources supporting horizontallyorganized knowledgebases
- 8. Methods enabling *scalable, intelligent, cost-effective FAIR* metadata*

FAIR*: Findable, Accessible, Interoperable, Reusability

- **9.** Cloud environment for NHGRI investigators to share data and tools
- 10. *Rigorous benchmarking* and development of 'gold standards'
- 11. Improvements needed to *integrate genomic medicine into Clinical Decision Systems*
- 12. *Integrating patients more fully into genomic medicine research* and clinical practice
- 13. Support informatics and computational needs for *single cell studies*

Moving from Recommendations to Potential Actions



Example Portfolio Analysis: Visualization (Interactive)

1 - Create query and aliases (avoid false negatives)

(e.g., "genome visualization"~10 OR "variant browser"~10 OR "population variation visualization"~10)



2 – Place constraints

(e.g., number of years, awards, nonawarded, HG-only, non-HG)

3 – Curate results (remove false positives)

Remove hits



Query	HG funded (before; after curation) last decade	Non-HG funded (before; after curation) last decade
"genome visualization"~10	42;11	131;42
"genome visualization"~10 AND interactive	3;3	27;14

- Query runs on abstract and specific aims.
- We can obtain detailed information on both awarded and non-awarded grants and potentially utilize the results to correctly reflect the needs.

Next Steps

- Publish report on website and advertise
- Expect to finish portfolio analysis by summer





Emphasis

The co-chairs: Drs. Mike Boehnke, Carol Bult, Trey Ideker, Aviv Regev, Lincoln Stein

Kevin Lee

NHGRI CGDS Staff: Lisa Brooks, Valentina Di Francesco, Dan Gilchrist, Mike Pazin, Erin Ramos, Heidi Sofia, Jen Troyer, Chris Wellington, Ken Wiley.

Eric Green, Carolyn Hutter, and Jeff Schloss,