



ALL IN ONE DAY PRECISION MEDICINE

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THE STEPS TOWARDS PRECISION HEALTH ANALYTICS

1

Patient visit

PRIMARY ANALYSIS



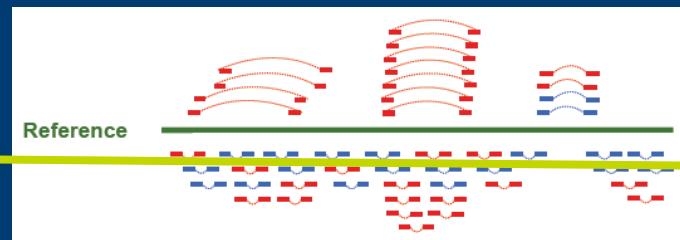
Individual

1-4 DAYS

2

Genes causing disease and key pathways identified

SECONDARY ANALYSIS, DNA/RNA PIPELINE + MORE



Multiple individuals

WEEKS

HC
HC
HC
HC
HC
HC
HC
HC
HC
HC
HC

*Multiple sample
compute starts here*

Joint
genotyping

Variant
store

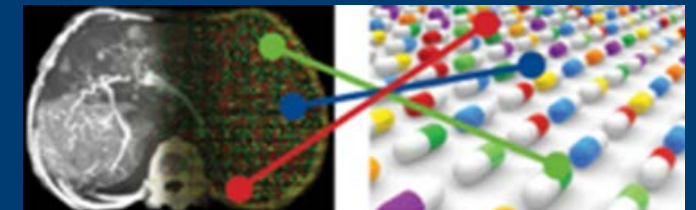
Pop/dis
study

3

Gene-targeted drugs identified, treatment begins in earnest

PRECISION MEDICINE

Predicted actionable variants,
data-driven association



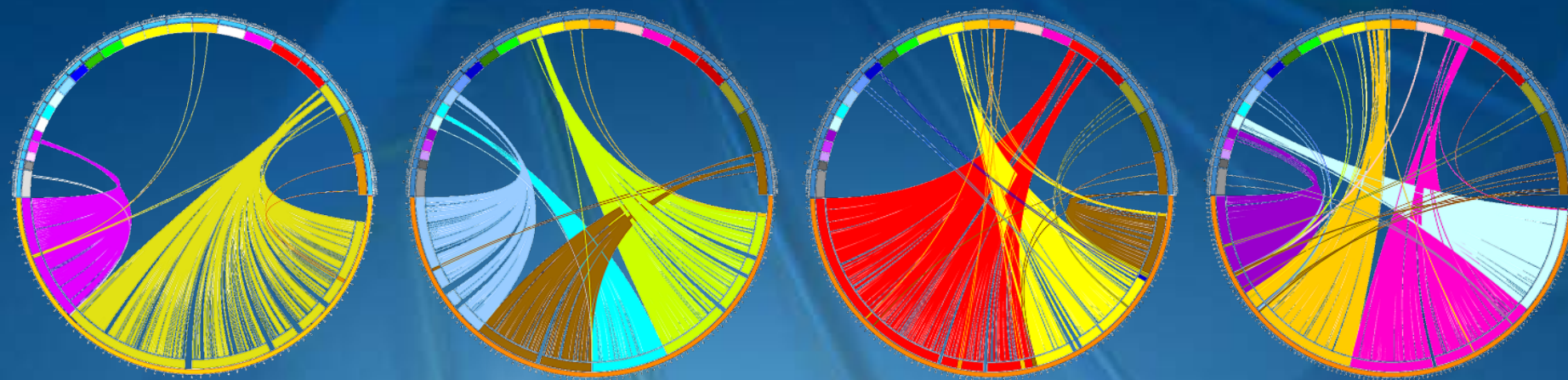
Clinically actionable variants

Clinical trial groups →
Data curation →

Knowledge
database

MONTHS

We Must Go Faster! And not via Phones, Faxes, & Fed-exing.

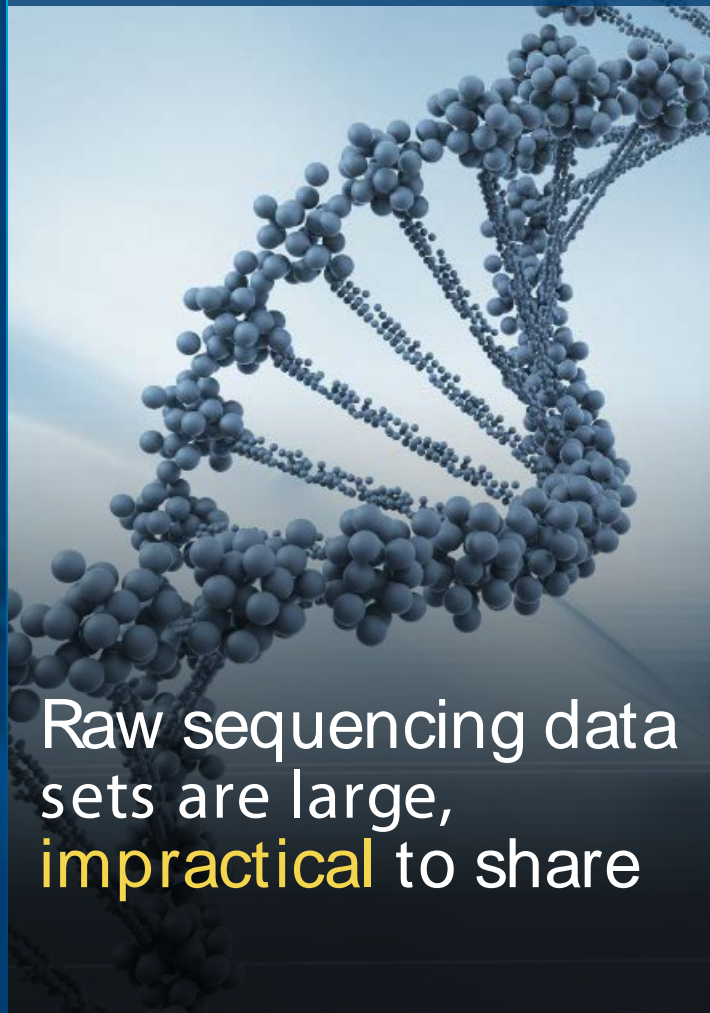


ALIGNED DAY BY 2020

MAIN COMPUTING CHALLENGES TO ACHIEVE 'ALL IN ONE DAY'

Size

Data management



Raw sequencing data sets are large, **impractical** to share

Speed

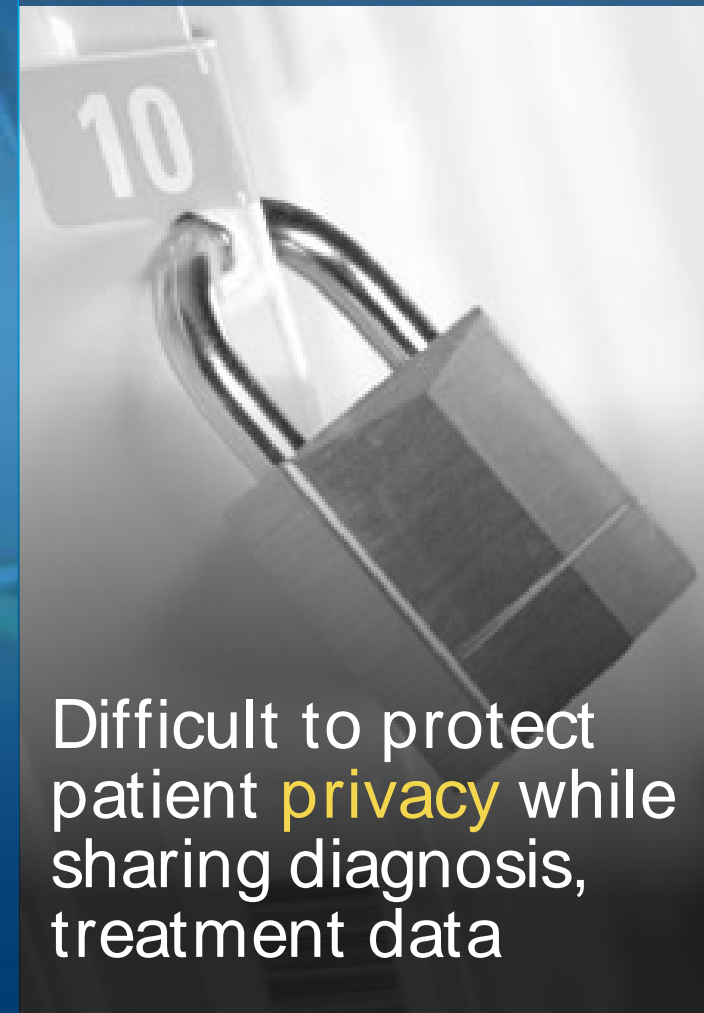
Workflow efficiency



Critical diagnosis, treatment pipelines can take **weeks**

Secure sharing

Privacy protection



Difficult to protect patient **privacy** while sharing diagnosis, treatment data

Scalability

Increasing requirements



Genomic sequencing **outpacing** capabilities of current architectures

An iceberg floating in a blue ocean. The small tip above the water represents 4% of data currently in research. The large, dark, and jagged mass submerged below the water represents 96% of data stuck in institutions.

4% of data currently in research

96% of data **stuck**
in institutions

- **Too large** to move
- **Protected**
(privacy laws, business model IP)

An iceberg floating in a blue ocean. The tip of the iceberg is above the water, while the much larger, submerged part is below the surface. The text '96%' is overlaid on the submerged part. To the right, text and a list are positioned. Lines connect the list items to the submerged part of the iceberg.

To achieve better research and clinical outcomes, we need

Collaborative analytics
across institutions

96%

- Sequencing data
- Treatments
- Outcomes

DESIGN CHALLENGES WE'VE BEEN FOCUSING ON

Get new insights in a secure, shared way without giving up control of your data

Move the
analytics,
not the data



Do it in an
open, secure
trust model



Do it at **scale**,
for
any researcher



Bring your own
tools and
infrastructure



Collaborative Cancer Cloud

With core DFC, OICR



- Jointly run **genomic pipelines**
- Drive **new analytic paradigms** against data sets using **integrated** platform
- Advance big data analytics in a **federated, secure, and scalable** environment

CCC is an open platform, enabling either on-premise or public clouds along with best practice precision medicine analytics

Genomictools



GATK

(The Genome Analysis Toolkit)



A wide variety of tools with a primary focus on variant discovery and genotyping

GenomicsDB

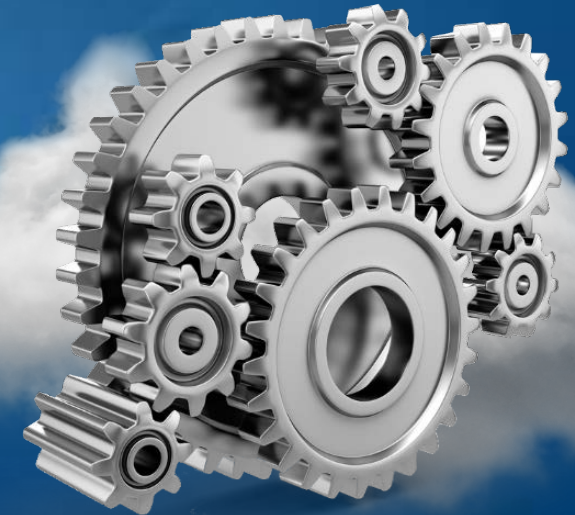
(Optimized for Genomics Data)



Store vast amounts of patient variant data and perform fast processing, at unprecedented scale

Cromwell

(Integrated workflow engine)



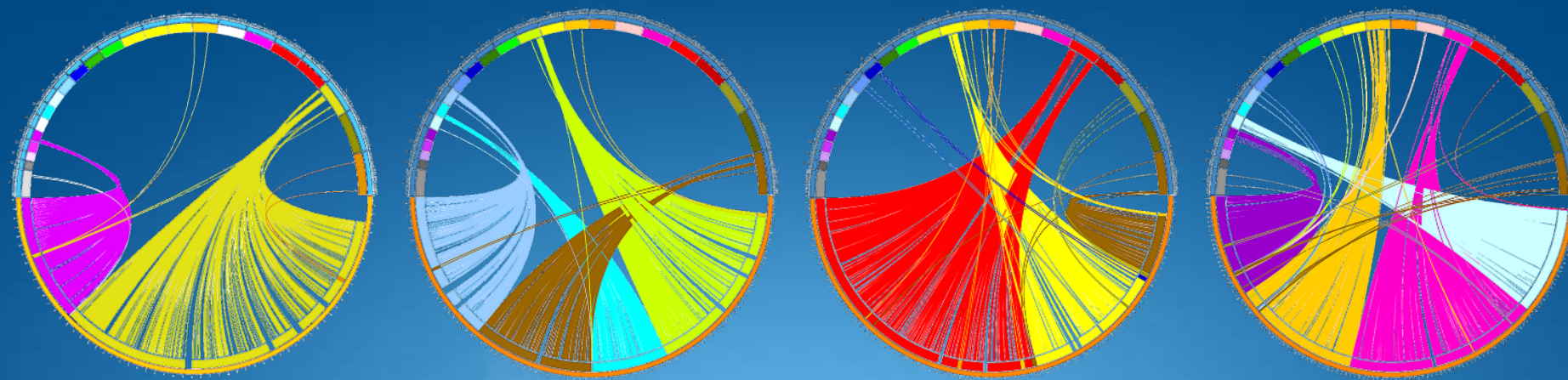
Launch genomic pipelines on private or public clouds in a portable and reproducible manner

AI in a day by 2020

- Need for federated, secure and scalable environments to enable precision medicine
- Tools –
 - <http://github.com/intel-hls/GenomicsDB>
 - <http://github.com/intel-hls/TileDB>
 - <http://github.com/intel-hls/Cromwell>

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ALIGNED DAY BY 2020

ALL-IN-ONE-DAY GOAL – A CHALLENGE FOR MEDICAL AND TECH INDUSTRIES

