

ROMOPomics

NCBI Codeathon, Pittsburgh 2020

Nick Giangreco
Andrew Clugston
Ben Busby
Alex Francette
Anish Chakka
Yash Gokhale

GitHub:

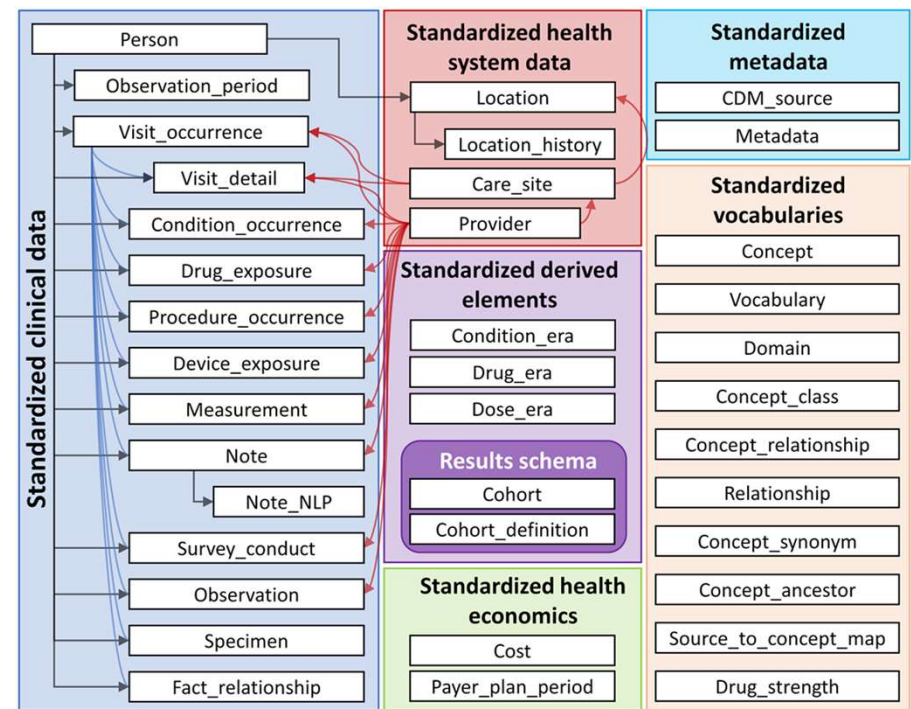
ngiangre
AndrewC160
DCGenomics
amfrancette
anishbhaswanth
yashgokhale

Twitter:

@AbsoluteClug
@DCGenomics

Patient data & OMOP

- Observational Medical Outcomes Partnership to transform patient data
- **Goal:** learning, evidence-based healthcare systems.
- **Problem:** patient data is separate, unharmonised within Hospitals, Medicare claims, [Clinical registries](#), etc.
- **Solution:** OMOP CDM
 - An established, patient-centric common data model already in clinical use.

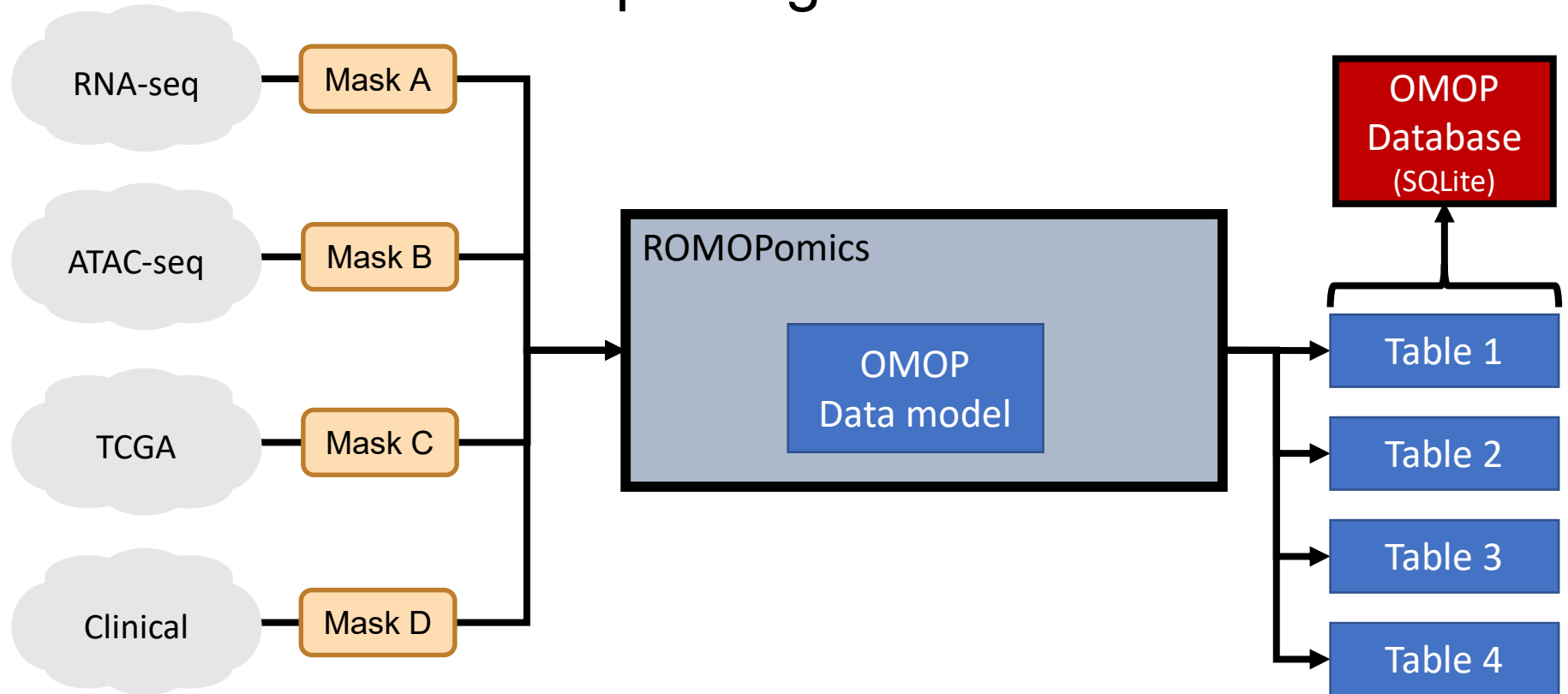


[The book of OHDSI](#)

OMOP with precision medicine and research.

- Precision medicine increasingly uses 'Omics' data (sequencing, etc.).
- Large amounts of 'omics data available (TCGA, GEO, clinics, etc.).
- **Goal:** Integrate 'omics data into patient clinical data.
- **Problem:** 'omics data is...
 - Detached from the patient.
 - Analyzed as a sample.
 - Difficult to retrieve and integrate.
- **Solution:** Use the OMOP common data model.

The ROMOPomics R package



Full integration into SQLite databases.

- The ROMOPomics R package combines sample-centric datasets from various sources
- Administrators build and implement the database, users interface via SQL.
- **Example:** Incorporate TCGA data, users can now query samples and treatments from patients of a certain age, sex, name, etc.

```
SELECT person_source_value, person.person_id, file_remote_repo_id, file_remote_repo_value  
FROM person INNER JOIN sequencing  
WHERE file_remote_repo_id IS NOT NULL  
AND person_source_value IS "tcga-3c-aaau"  
ORDER BY file_remote_repo_id
```

ROMOPomics

Dev. version available on GitHub:

`install_github("andrewc160/ROMOPomics")`

	<i>GitHub:</i>	<i>Twitter:</i>
Nick Giangreco	ngiangre	
Andrew Clugston	AndrewC160	@AbsoluteClug
Ben Busby	DCGenomics	@DCGenomics
Alex Francette	amfrancette	
Anish Chakka	anishbhaswanth	
Yash Gokhale	yashgokhale	