









Leading the search for tomorrow's cures



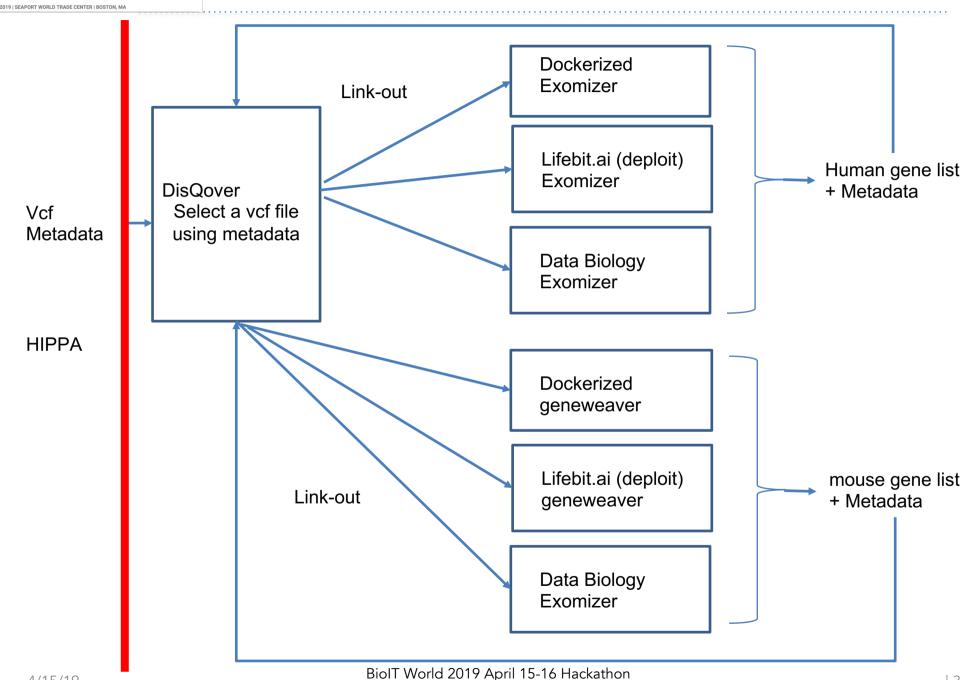




Complete Workflow

- Data are FAIR
- Applications are FAIR
- Data out are FAIR





JAX FAIR Beyond Data to Applications Team



Subteam formation

- 1. Exomiser dockerization lifebit.ai, databiology (leads: Juan/Pablo)
- 2. Geneweaver homology call (lead: Filipe)
- 3. DisQover (lead: Niels)
- 4. Data Input/Output gene, genesets (lead: Nolan)



Workplan

- 1. Initial/Baseline Assessments (Data and applications)
- 2. Conduct interventions
- 3. Re-Assessments (Data and Applications)



Challenges

 For Diagnostic purposes – VCF data is diagnostic information only available