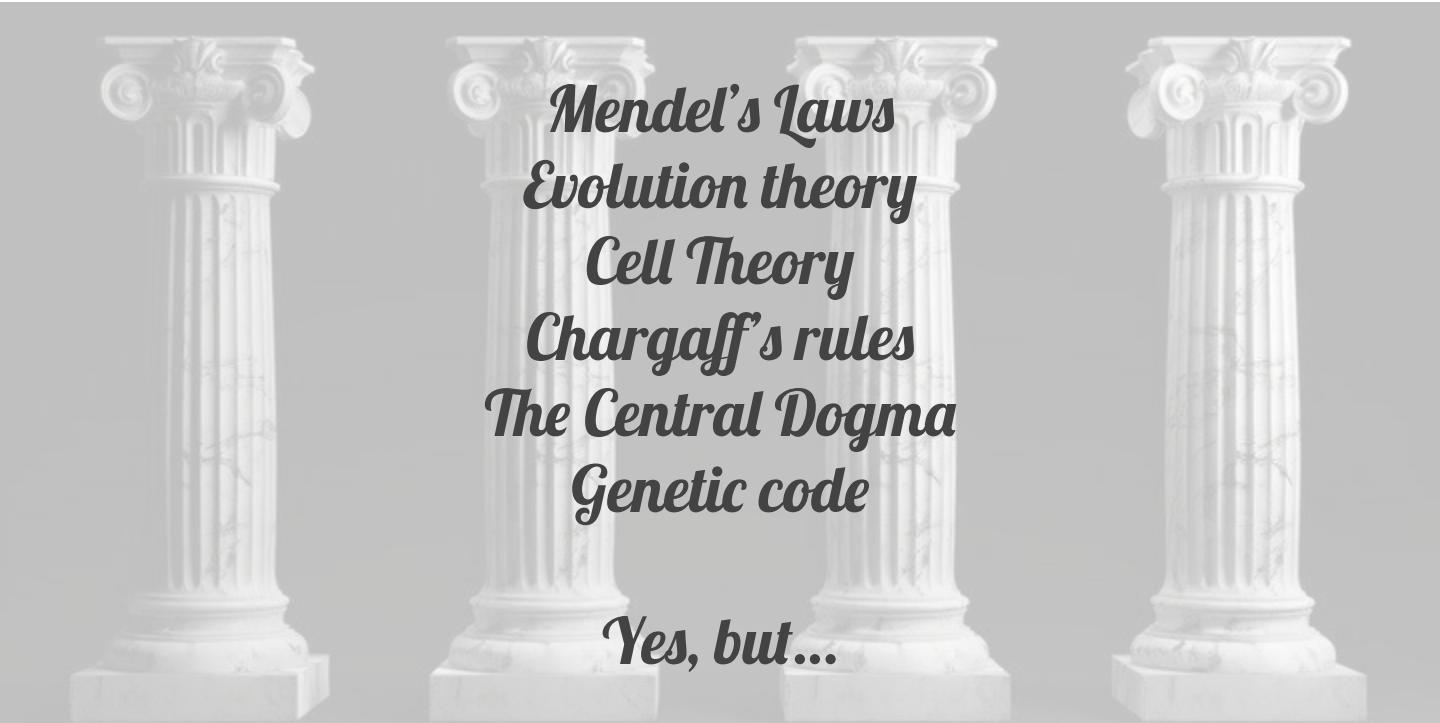


ECM Aging Atlas:
a database of time-resolved proteomic signatures

Biological first principles



*Mendel's Laws
Evolution theory*

Cell Theory

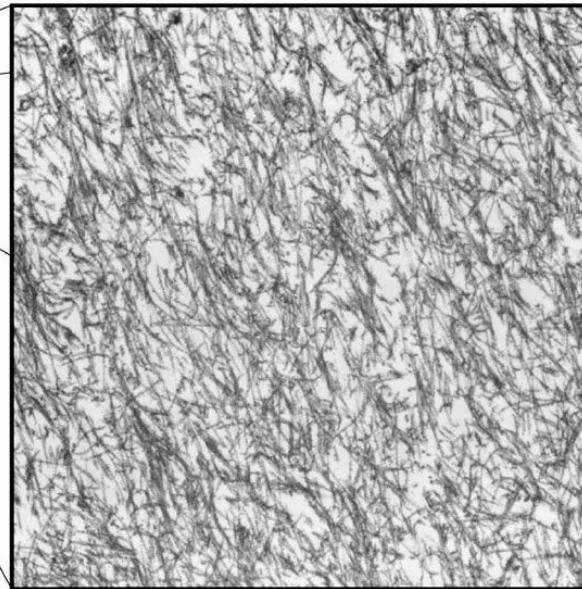
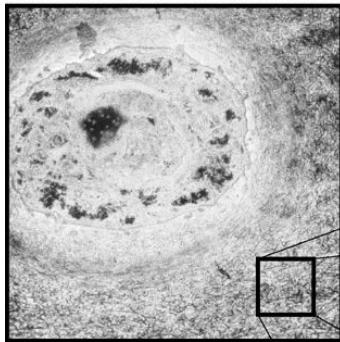
Chargaff's rules

The Central Dogma

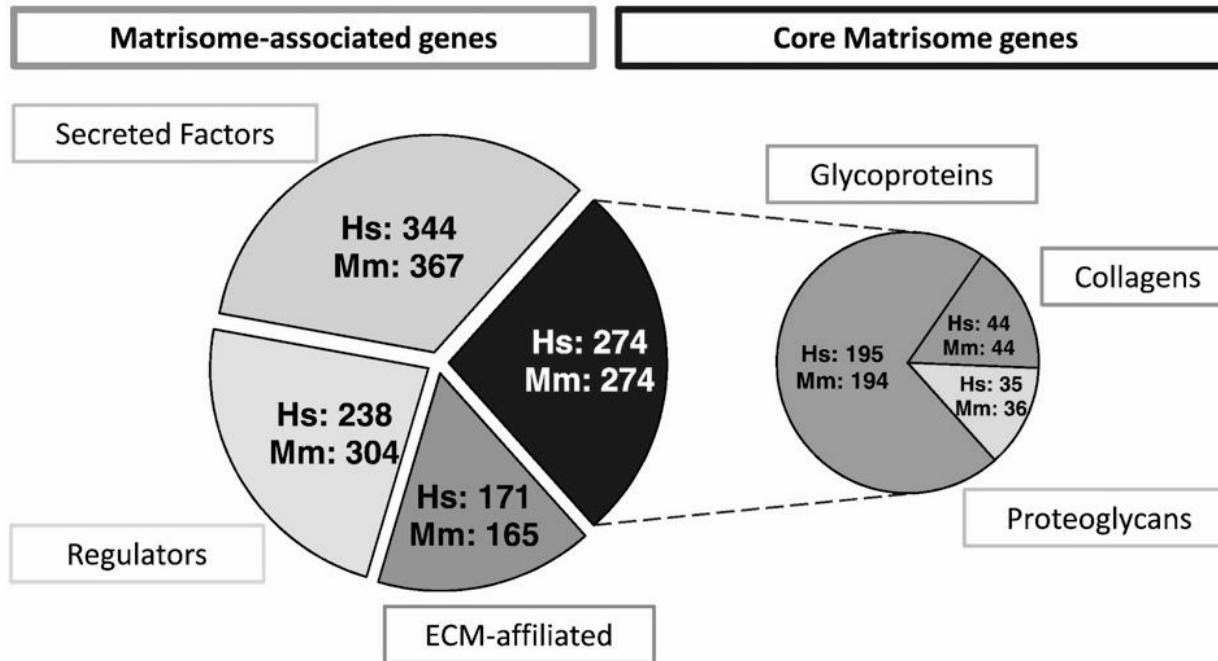
Genetic code

Yes, but...

Ground substance



Extracellular matrix



Problem and solution

Problem: *proteomics data are non-uniform!*

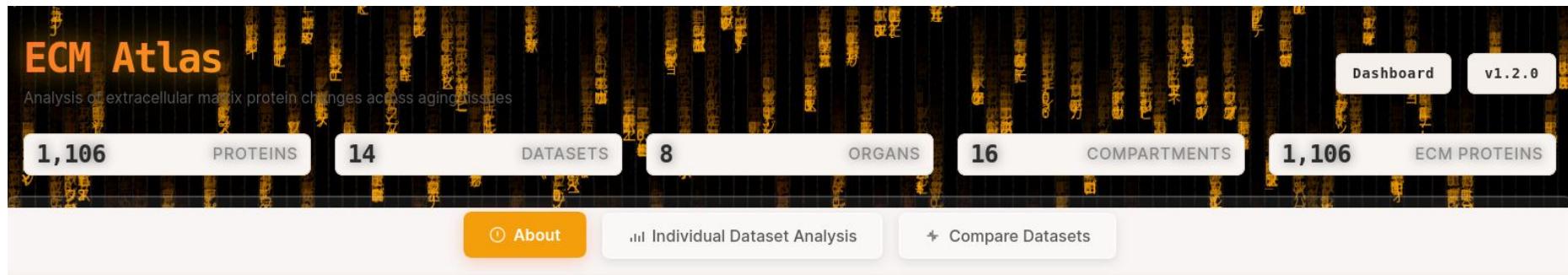
- different sample preparation workflows
- different quantification methodologies (*label-free vs. labeled*)
- different instruments (*Orbitrap, time-of-flight, etc.*)
- different data analysis pipelines and output dataset formats

Hard to compare studies directly.

Solution:

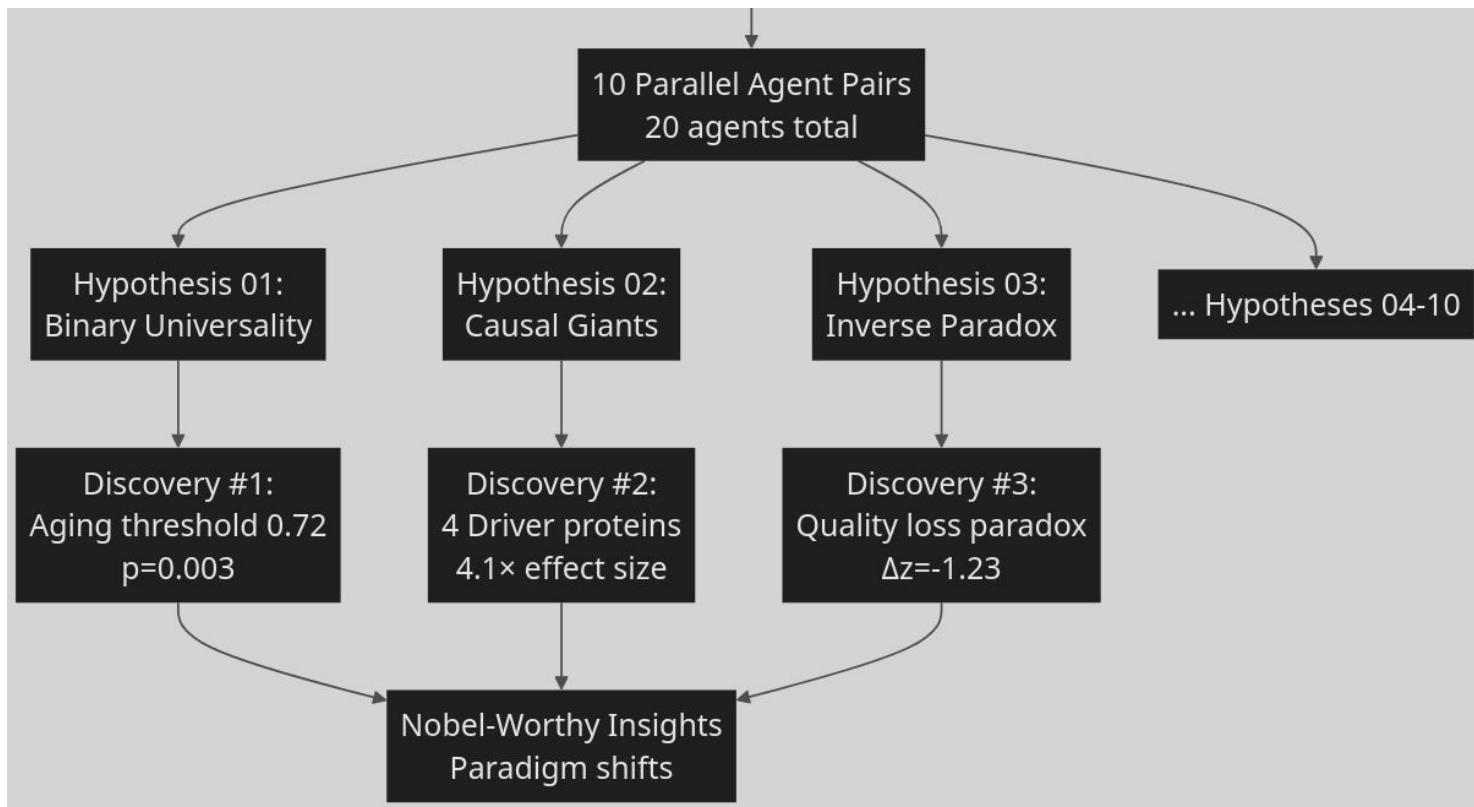
- *data ingestion agent* (*data wrangling, normalization, batch correction*)
- a *database for unbiased comparison between studies and compartments*

ECM Aging Atlas

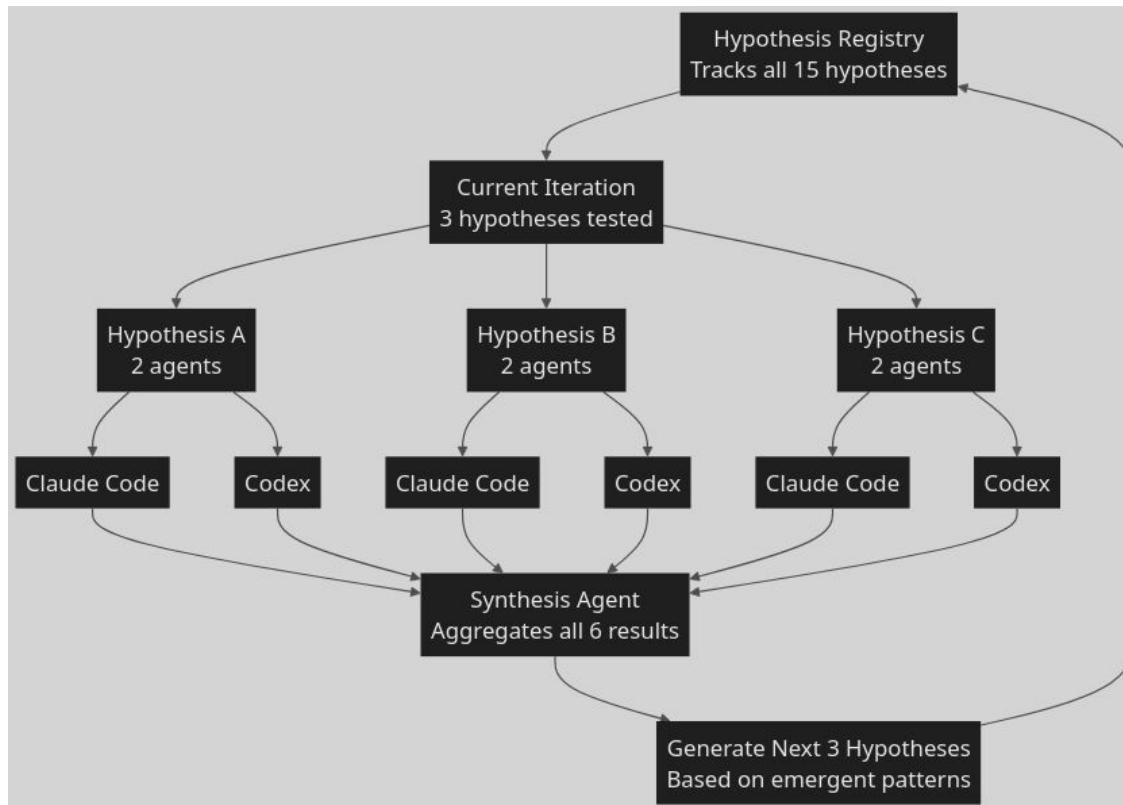


Live dashboard: <https://ecm-atlas.fly.dev/dashboard.html>

Hypothesis generation



Hypothesis validation



Team

Bob Walsh, data analyst, mathematician

Daniel Kravtsov, developer, entrepreneur

Rakhan Aimbetov, biologist

*“Truly, what is stiff and hard is a companion of death;
what is soft and weak is a companion of life.”*

-- *Tao Teh Ching, 76.*

