HTCondor for machine learning in biology

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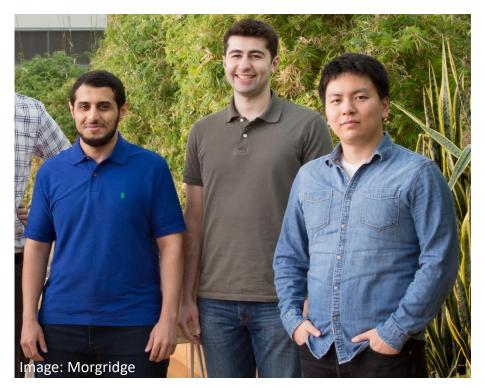


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Drug discovery and GPU computing



Moayad Alnammi Shengchao Liu Sam Gelman

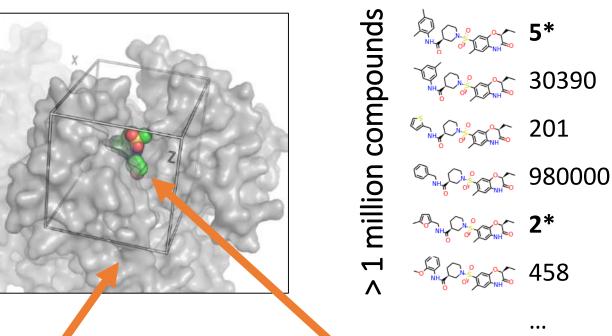
High-throughput chemical screening

Given a protein of interest, identify chemicals that may have the ability to control the protein



Computational chemical prioritization

New protein target

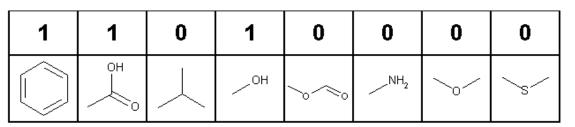


Suggested test order

Chemical compound ≈ key

Protein ≈ lock

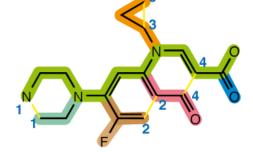
Chemical representations for machine learning



HN

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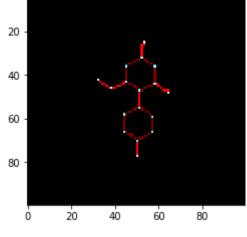
2-D searching tutorial



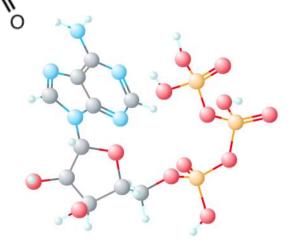
N1CCN(CC1)C(C(F)=C2)=CC(=C2C4=O)N(C3CC3)C=C4C(=O)O

Wikipedia SMILES

See Ching et al. 2018



Chemception



PubChem

OH

Training neural networks



Anthony Gitter @anthonygitter

We Academic hardware program is great registration.nvidia.com/ahr.aspx

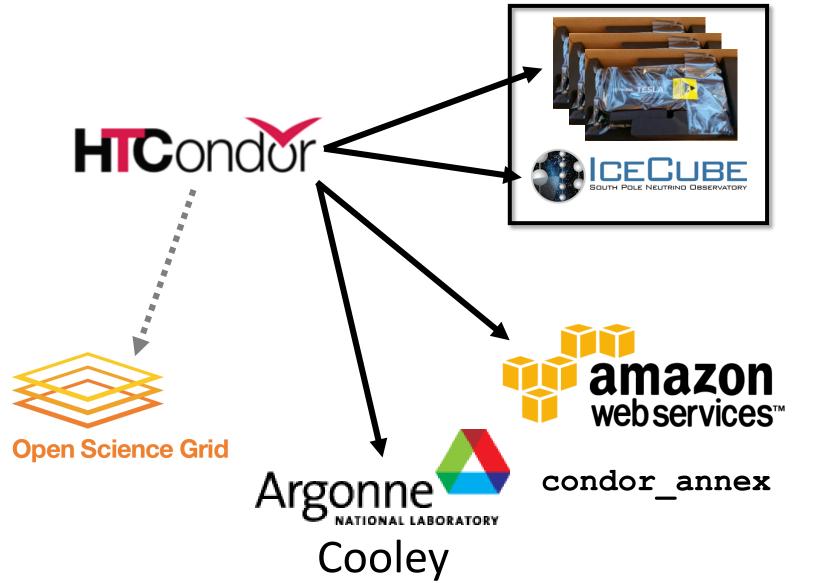
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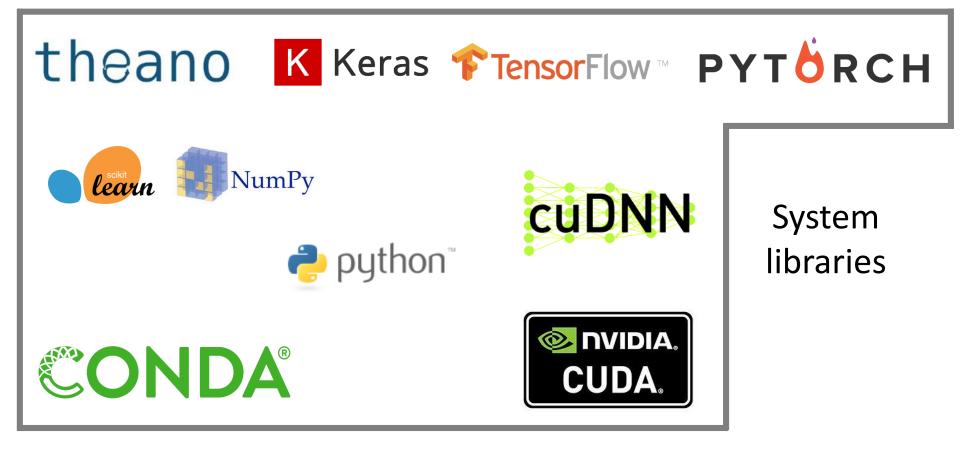
Expanding our GPU capacity



UW grid



Chemical screening software



NVIDIA driver XXX.YY

Tradeoffs of conda

Pros:

- Easy to install Anaconda
- Environments are shareable
- Easy to update packages in environment
 Cons:
- Still depend on system libraries
- conda_submit gpu-job.sub
- condor install numpy

Predictive models perform well in experimental tests

Train on <u>75k</u> chemicals, PriA-SSB inhibition Choose among many possible models

Models select 250 of 25k new chemicals

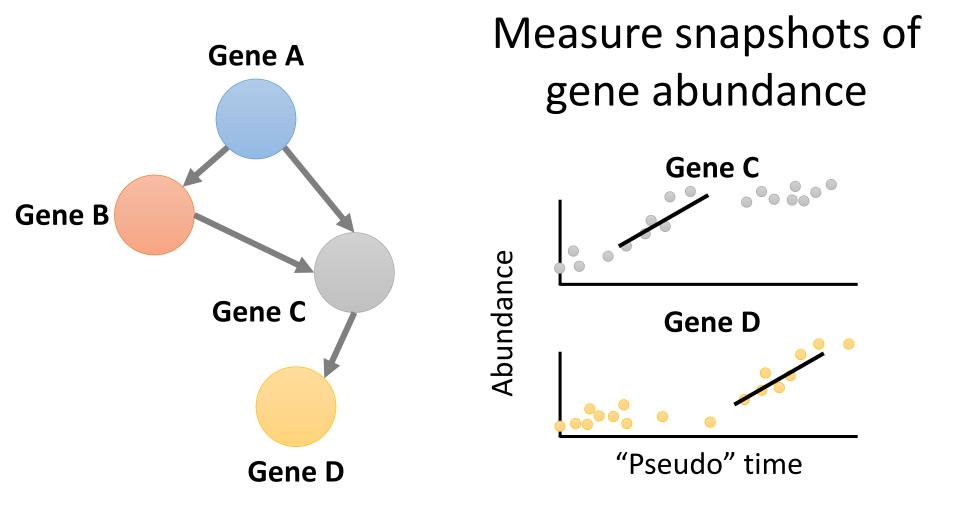
- 64 active chemicals in the 25k
- Model we selected is the best, finds 40 of the 62
- <u>Random forest outperforms the neural networks</u>
- Now testing on much larger chemical libraries

Gene networks and single cell expression

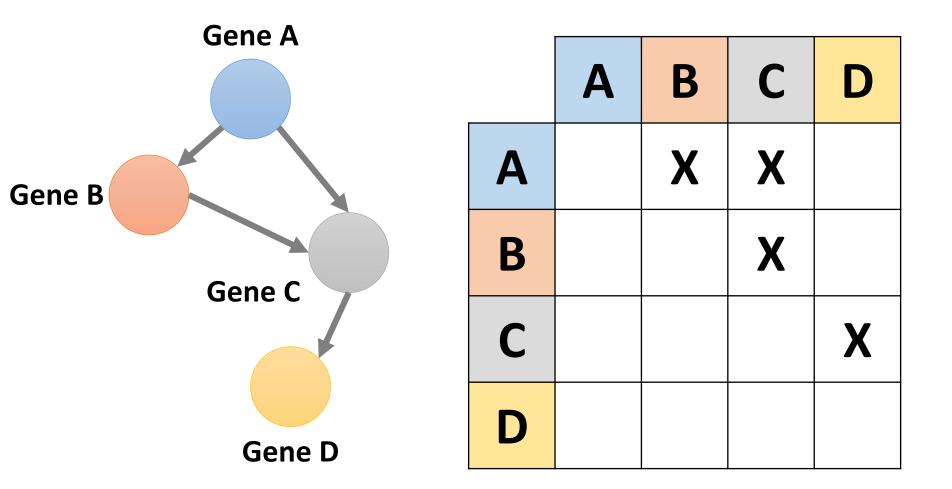


Atul Deshpande

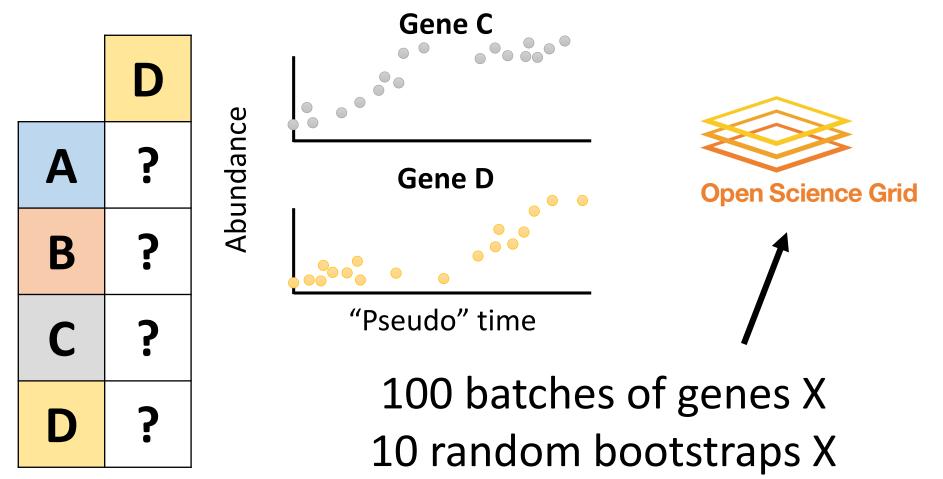
What are the relationships among genes inside a cell?



For each gene, learn which genes control it



Divide network inference into many small computational jobs



100 parameter combinations

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