

Integrating heterogeneous predictive models using Reinforcement Learning

[Ana Stanescu](#)¹ and [Gaurav Pandey](#)²

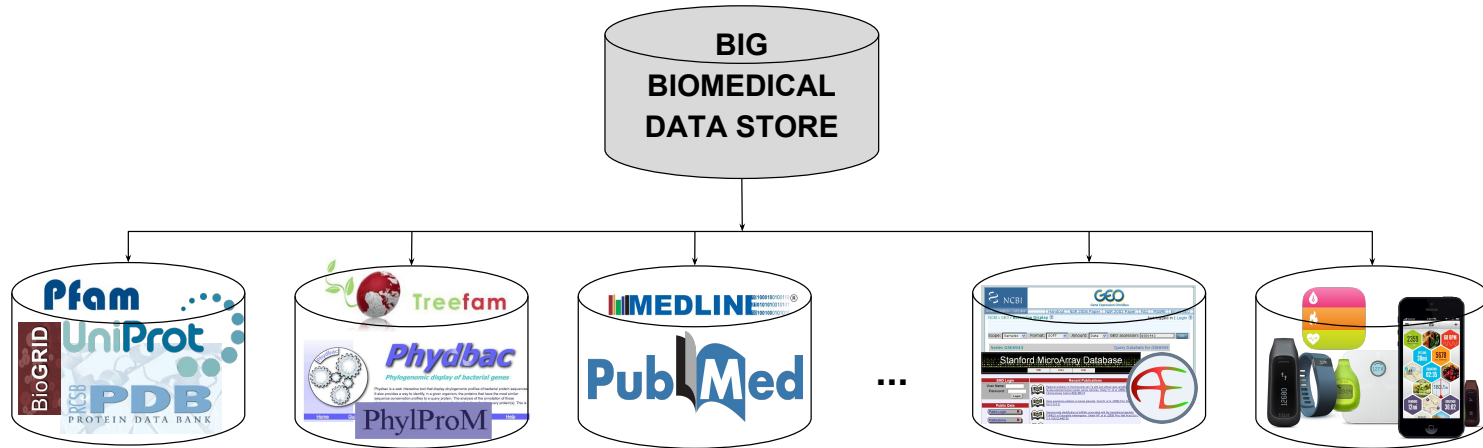
¹Department of Computer Science, University of West Georgia, Carrollton, GA, USA

²Department of Genetics and Genomic Sciences and Icahn Institute for Genomics and Multiscale Biology, Icahn School of Medicine at Mount Sinai, New York, USA

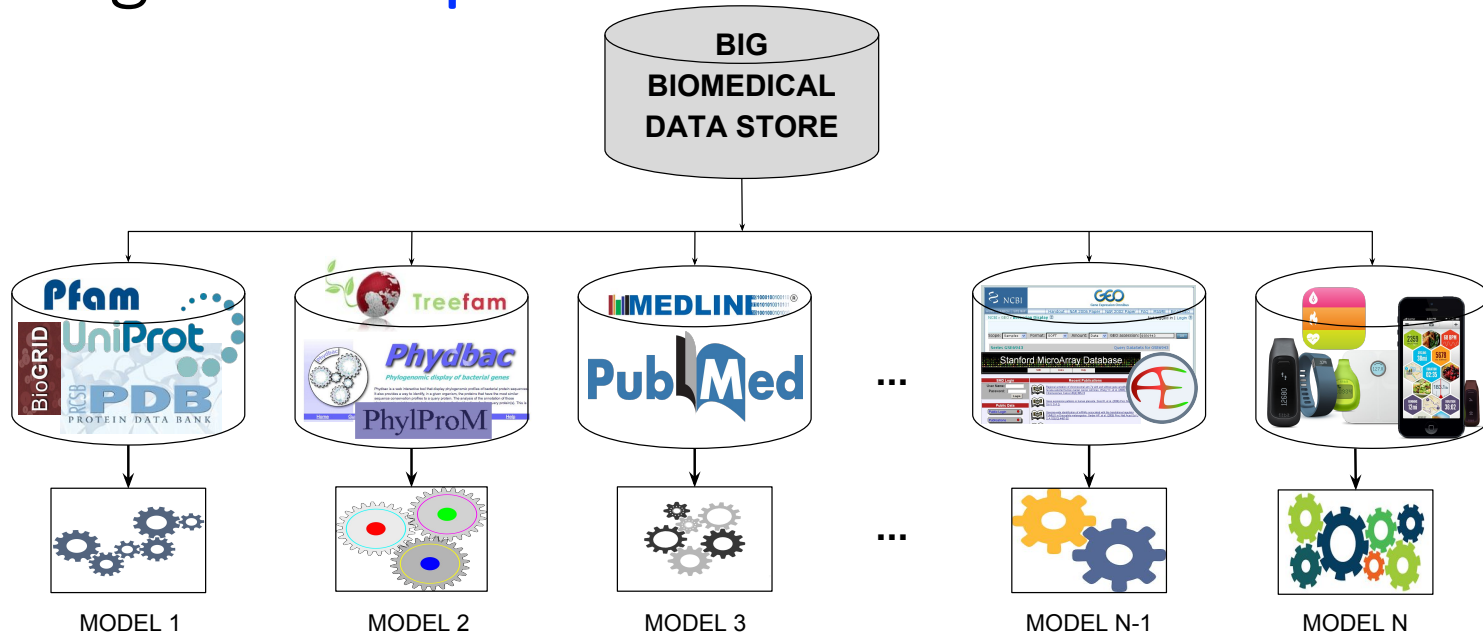
* Partially supported by the NIH BD2K Program



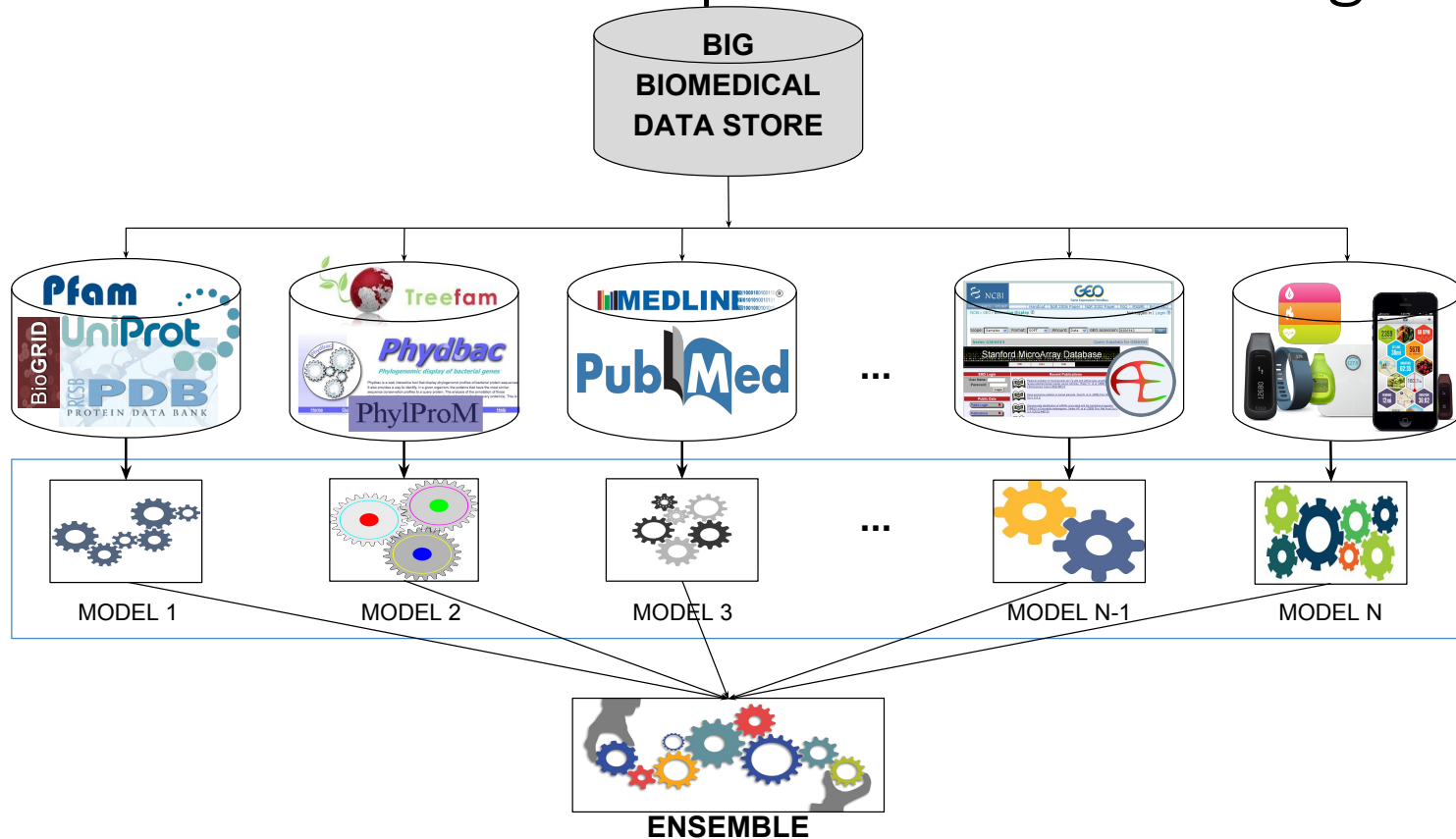
Biomedical data are abundant



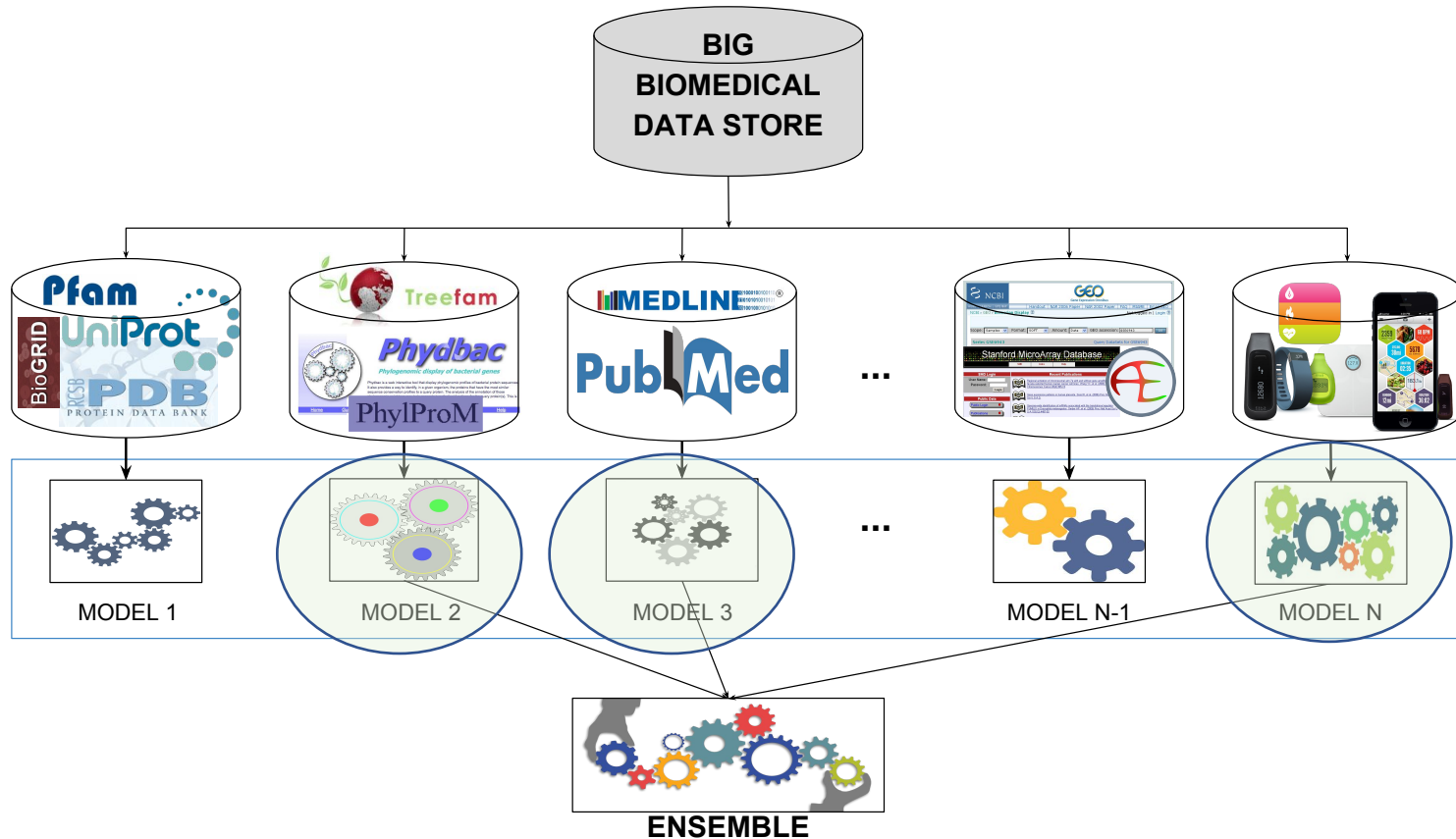
Systems biology and machine learning can generate **predictive models** from data



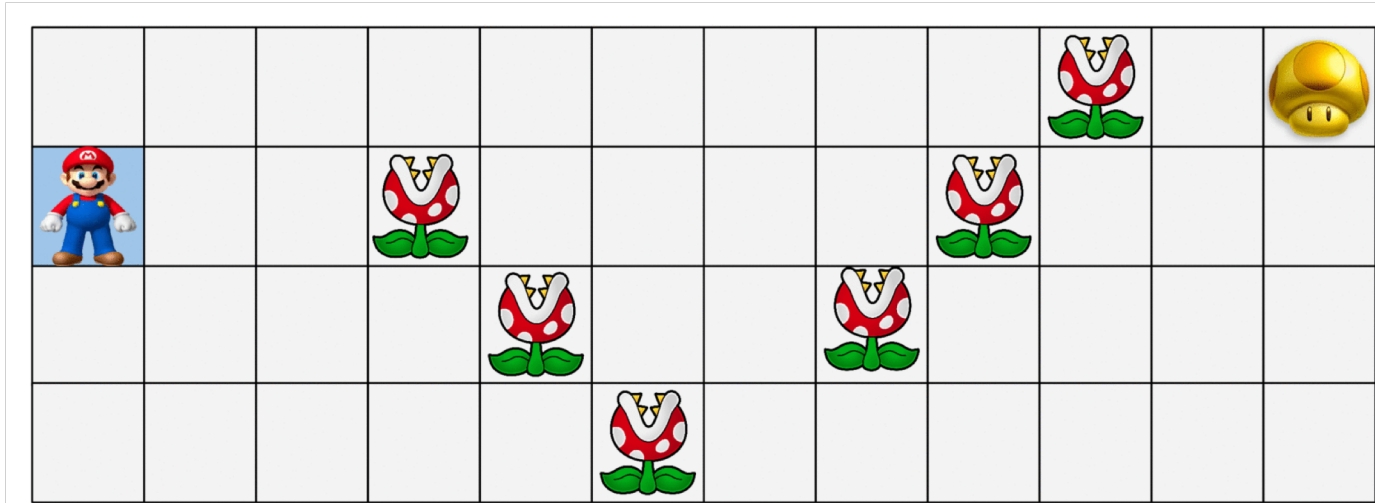
Heterogeneous ensembles can enhance effectiveness of predictive modeling



Selecting a parsimonious set of models into an ensemble can further advance predictive performance and interpretability

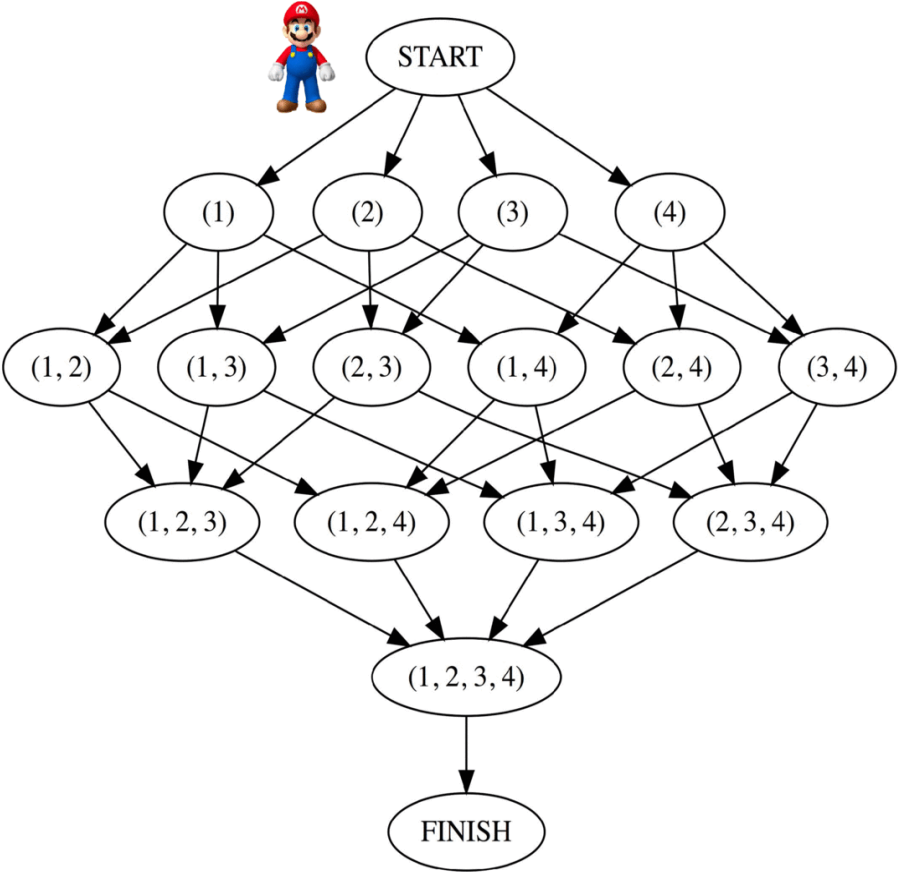


Reinforcement Learning: searching a large structured environment with rewards to find an optimal path (behavior) to reach the goal



An agent learns by interacting with its environment through “**exploitation-exploration**”.

Ensemble selection using Reinforcement Learning



Reward functions can be formulated in terms of ensemble performance and/or diversity

Fine balance between ensemble performance and ensemble diversity

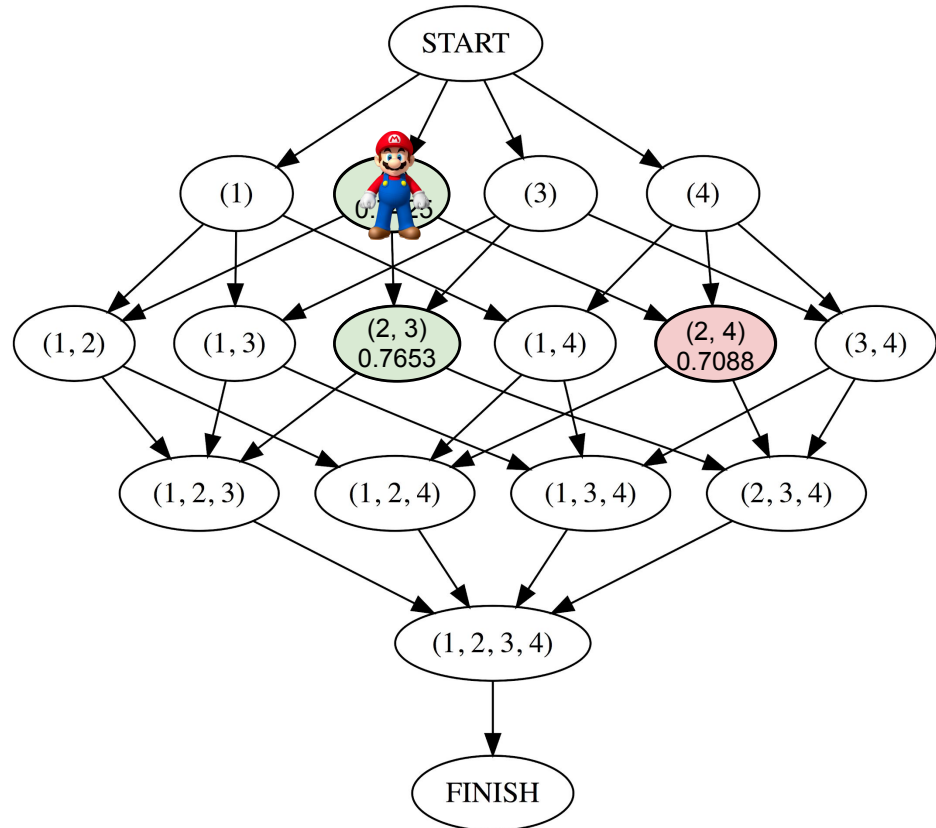
We have designed several search strategies focused on:

- **performance**

([Stanescu and Pandey, PSB 2017](#))

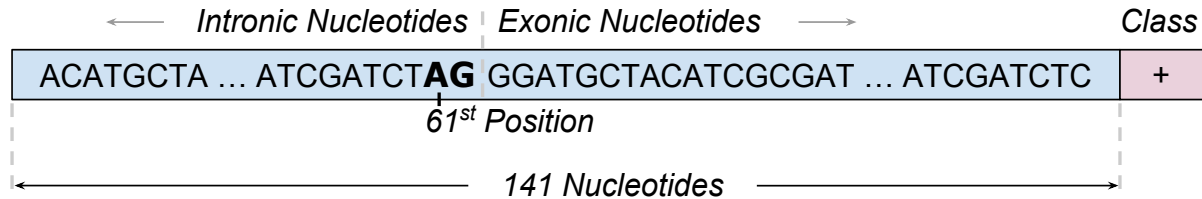
- **diversity**

([Stanescu and Pandey, arXiv 2018](#))



Target problem and evaluation methodology

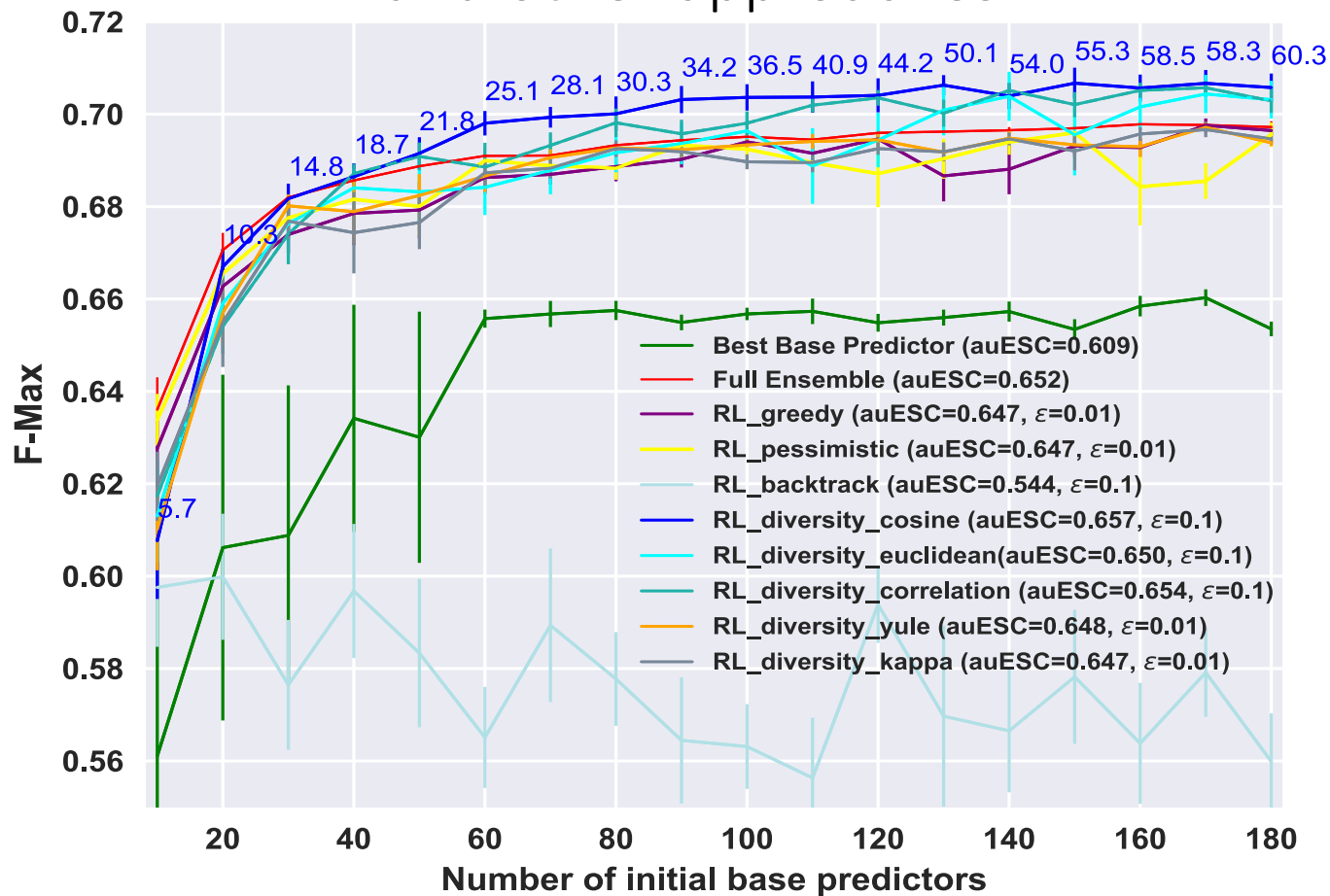
- Predict splice sites in various organisms based on nucleotide positional match representation using several public datasets.



Problem	<i>C. elegans</i>	<i>D. melanogaster</i>	<i>P. pacificus</i>	<i>C. remanei</i>	<i>A. thaliana</i>
#Features	141	141	141	141	141
#Positives	1,598	997	1,596	1,600	1,600
#Negatives	158,150	99,003	156,326	157,542	158,377
Total	159,748	100,000	157,922	159,142	159,977

- 10 bagged versions of 18 different classifiers: 180 base classifiers in a 5-fold cross-validation setup.

Performance of ensembles selected using RL and other approaches



Conclusions

- Reinforcement learning-driven ensembles are **competitive in predictive performance to** larger ensembles consisting of all base predictors, while being substantially smaller, *i.e.*, **more parsimonious**. ([Stanescu and Pandey, PSB 2017](#))
- Ensemble diversity, measured appropriately, can build **even more accurate and parsimonious ensembles**. ([Stanescu and Pandey, arXiv 2018](#))
- **Implementation available:** <https://github.com/GauravPandeyLab/LENS>
- **Future Work**
 - Test the RL ensemble framework on larger datasets, including non-biomedical ones.
 - Develop more efficient (parallel) implementations of the framework.

Acknowledgements

- NIH Grant R01-GM114434
- Pandey Lab at Mount Sinai
- Minerva supercomputing team at Mount Sinai
- Computer Science Department at UWG

Thank you!