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2021-2022

# LEARNING THE GOVERNING EQUATIONS OF ECOLOGICAL DYNAMICS

# Markus Bauer

B.S. Computer Science - Data Science

Role: Machine Learning Developer

Focus: Balanus and Mytilus Growth Rate



# Samson Mont

B.S. Computer Science - Artificial Intelligence

Role: Machine Learning Developer

Focus: Prey Density and Temperature  
Predicting Feeding Rate



# Nour Rahal-Arabi

B.S. Computer Science - Cybersecurity

Role: Machine Learning Developer

Focus: Symbolic Regression Output Equation  
Interpretation/Parsing





# Madelyn Smith

B.S. Computer Science - Custom Applied Plan

B.S. Sustainability

Role: Lead Researcher

Research Focus: Relevant Work (Symbolic  
Regression in Ecological Studies)



# Jinshui Wang

B.S. Computer Science - Machine learning

Role: Machine Learning Developer

Focus: Growth rate of mussel



Our Task: Answer a fundamental question in Ecology

How and why do species abundances vary over space and time?

## PROJECT DESCRIPTION

Create a machine learning model to describe abundance dynamics as influenced by predator-prey interactions in the Oregon marine intertidal which is easy to understand and accurate, based on time-series data from a three-year ecological field experiment



# An Introduction to gplearn

A genetic programming package in python

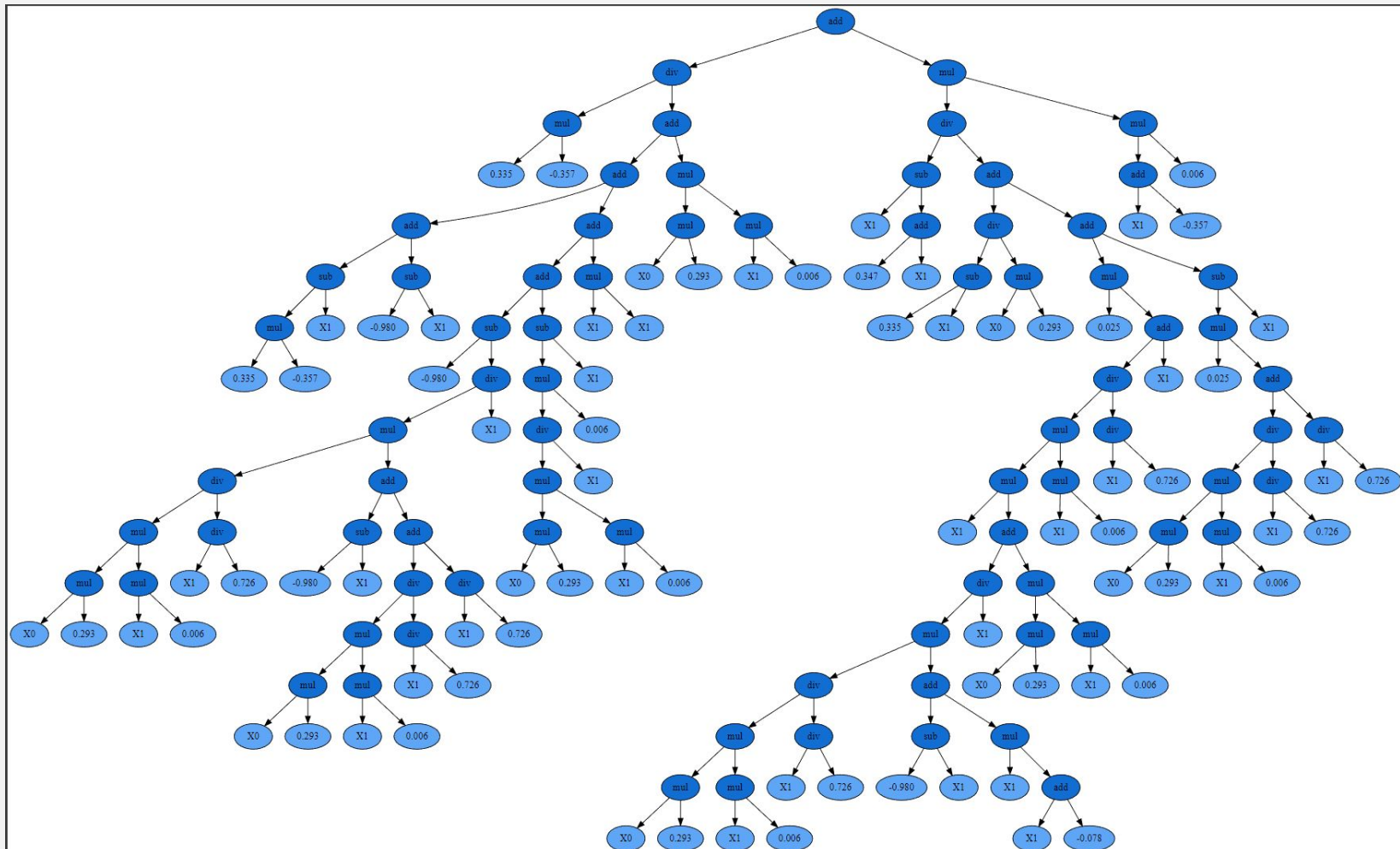
## Symbolic Regressor

The Tool of Choice

```
classgplearn.genetic.SymbolicRegressor(*,  
population_size=1000, generations=20,  
tournament_size=20, stopping_criteria=0.0,  
const_range=(- 1.0, 1.0), init_depth=(2, 6),  
init_method='half and half', function_set=('add',  
'sub', 'mul', 'div'), metric='mean absolute error',  
parsimony_coefficient=0.001, p_crossover=0.9,  
p_subtree_mutation=0.01,  
p_hoist_mutation=0.01,  
p_point_mutation=0.01, p_point_replace=0.05,  
max_samples=1.0, feature_names=None,  
warm_start=False, low_memory=False,  
n_jobs=1, verbose=0, random_state=None)
```

## I/3 Parsimony Coefficient

- Most important parameter – and the most frustrating to work with
- Encourages the model to pick smaller programs to avoid equation bloat
- High Parsimony means short equations
  - Can get “stuck”
- Must be relative to the results, high y values will need much higher parsimony
  - Was a problem during our project



## 2/3 Trials and Runtime

- Stopping Criteria – If the model reaches this threshold it will stop executing
- Generations – The number of generations to run if the stopping criteria is never met
- Population Size – The number of equations created per generation
- Tournament Size – A subset of the population where the fittest equation is selected to move on to the next generation

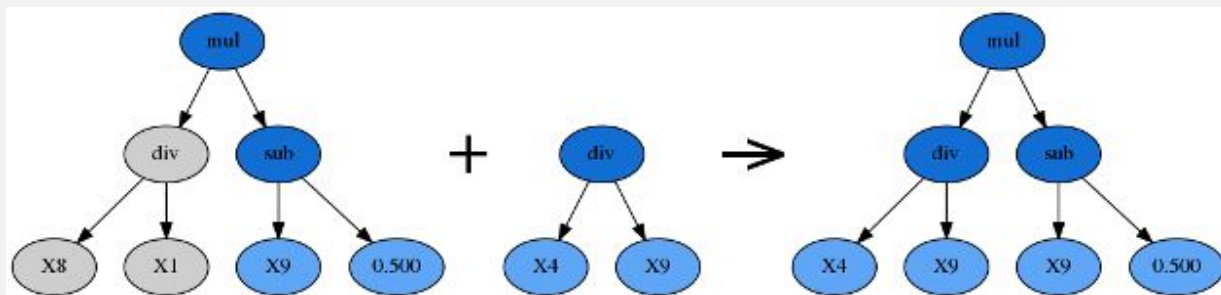
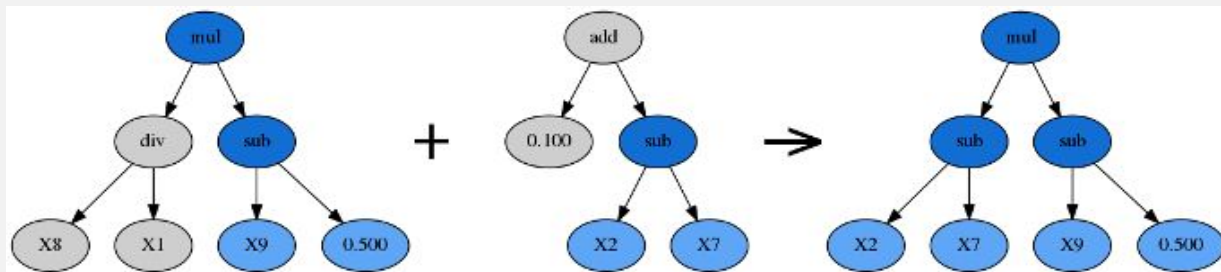
## 3/3 Evolution (4-P's)

### Run on Tournament Winners

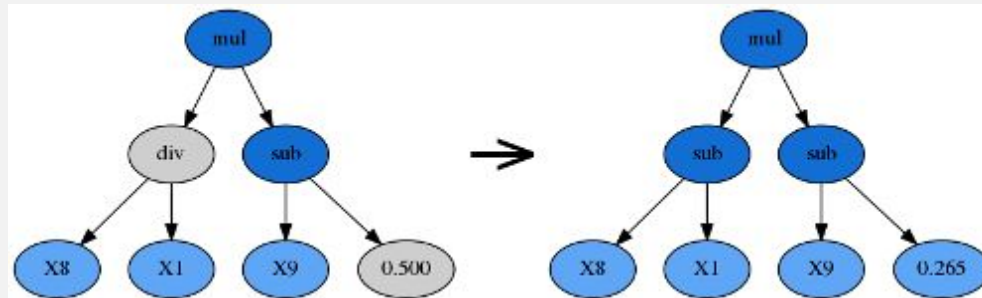
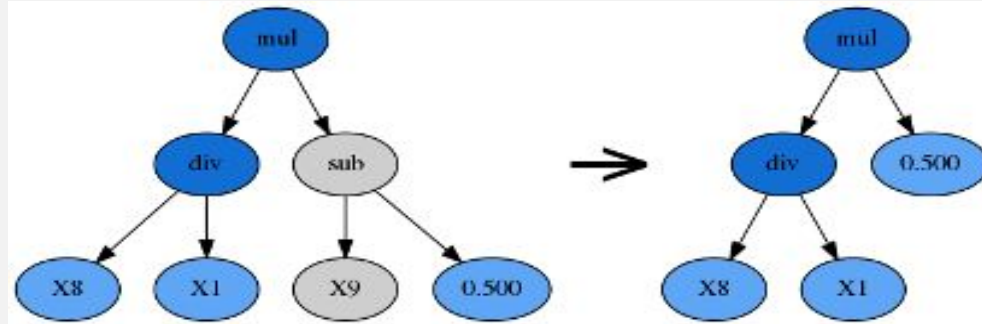
- Crossover – Running two tournaments and replacing a branch from the first with one from the second
- Subtree Mutation – Replaces branches with randomly generated subtrees
- Hoist Mutation – Deletes sections of a subtree to fight bloat
- Point Mutation – Selects random nodes and replaces them with random functions and terminals
- Reproduction – If the above parameters do not sum to 1, the remainder is the probability of the selected winner advancing unchanged to the next generation



## Crossover vs Subtree Mutation



## Hoist vs Point Mutation



## Honorable Mentions

- Function Set – The functions that the regressor uses
- Low Memory – If set to true, will not store previous equations when no longer in use
  - Very helpful for running through Google Colab which had memory issues when we would run long tests
- N\_Jobs – Allows for concurrent runs of the fitness function, only helpful with large function sets or input variables
- Warm Start – If true, will continue running where the previous function stopped
  - Didn't get it to work in colab, but would have been very helpful in saving overnight runs that timed out

# Conclusion and Future Development

- Another regression
- Another development tool
- Extract immediate equation
- Produce more graph or visual

Thank you.