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The Data Mine

Modeling Crop Gene-Environment Interactions under Climate Change

Project Overview and Motivations

Overview:

The human population is growing rapidly and the demand for food continues to increase. In the figure to the right, the corn yield has increased dramatically over the years. The increase began in the 1930s due to the double-cross hybrid corn. The demand for corn will continue to increase which is a driving force in the motivation for this project.

Motivations:

- Cope with climate change
- Reduce efforts in predictions
- Resource management
- Competitive advantages in crop growing



derived from annual USDA-NASS crop production reports.



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DATA OVERVIEW

• Around 3000 genetic markers for parent and progeny • 1 represents dominant homozygous, 0 represents heterozygous, and -1 represents recessive homozygous

• Copernicus ECMWF Reanalysis 5 (ERA5) • 2030, 2040, 2050 benchmarks Coupled Model Intercomparison

• Snow Fall/Snow Melt Rates and Snow Thickness

Cross Pollination (Hybrids)





Inbred

hybrid

Inbred

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MODELING APPROACHES

M	Ietrics:
-	RMSE: 1

- ISE: mean squared error MAPE: mean absolute percentage error
- R²: percentage of the variation in yield explained by the model

Model comparison:

Comparing the models, Light gradient Boost has the best scores. For RMSE, the root mean square error, lower scores represents a better fit to the data. Additionally, for MAPE, the mean absolute percentage error, a lower score is also preferred. The higher the R², the better the model fits the data. When looking at LGBoost we can see its RMSE, MAPE, R² scores are most suitable for our goals.

CONCLUSIONS and **PREDICTIONS**

- We have created a new data frame that implements genomic and climate data
- We used new models and worked on creating more accurate predictions
- Integrating a user interface that allows us to look at predictions based on different variables

REFERENCES

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