SENTINELS OF THE HARVEST

GEOSPATIAL MONITORING OF CROPLAND HEALTH FOR A CHANGING CLIMATE

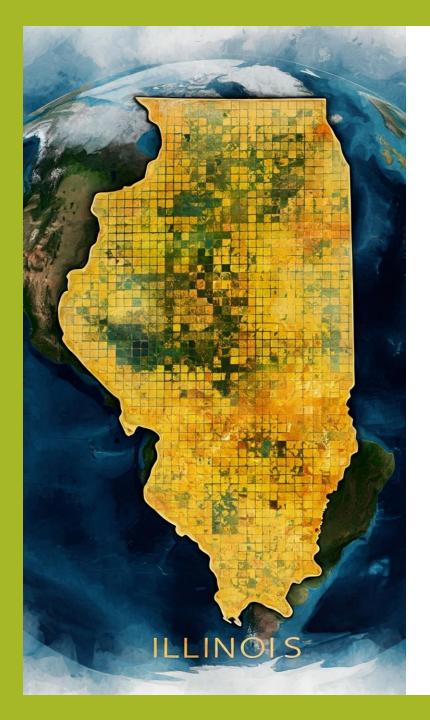
Team SaFe MeMe

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Why do we care?

- Cropland is an important thing to look at because of the US dependence on it
- Climate change presents a threat to cropland health
- Previous Research:
 - o Studies on climate change and vegetation in general





Phase 1: Data Collection and Preparation

Objective:

• Analyze the impact of climate change on cropland health and productivity in Illinois.

Methodology:

- Utilize Google Earth Engine API to access and analyze geospatial data.
- Focus on Illinois croplands.
- Analyze data at a 1km x 1km sptial resolution.
- Use data from 02/2000 to 12/2023.
- Use weekly resolution (1244 weeks in total)

Datasets:

- USDA NASS Cropland Data Layers
- MODIS (Normalized Difference Vegetation Index (NDVI), Enhanced Vegetation Index (EVI), Terra MODIS Vegetation Continuous Fields)
- PRISM (temperature and precipitation)

Parameters Obtained:

- Daymet: dayl, prep, srad, swe, tmax, tmin, vp
- MODIS: EVI, NDVI, ET, LE, PET, PLE, FPAR, LAI

Phase 1: Data Collection and Preparation

- Pipeline Overview:
 - Functional programming paradigms were essential to efficiently process large datasets from Google Earth Engine (GEE).
- Automation and Preprocessing:
 - Bash Scripts: Automated the retrieval of GeoTIFFs via GEE over an 8-hour runtime to manage large-scale data pulls. Downloaded 1244 GeoTIFF files (11.2GB) directly via API.
 - Object-Oriented Programming: Leveraged a custom GeoTIFFImage class in Python for preprocessing, including metadata extraction, band visualization, and flattening to pandas DataFrames.
 - Preprocessing ensured data quality by clipping to Illinois, aligning temporal spans, and removing non-relevant pixels.
 - Produced 205,151,722 worth of 20-dimensional data-points



Examples of Shell Scripting / Functional Programming

```
#!/bin/bash

# Run the Python script in the background via nohup; log terminal prints to home dir file
nohup python3 -u /home/katavga/code/safe-meme/gee_acquire_data.py > ~/gee_acquire_data.log 2>&1 &

# Confirm successful script instantiation
echo "Script is running in the background. Logs are being written to ~/gee_acquire_data.log"

# Verify that the process is running
ps -ef | grep gee_acquire_data.py | grep -v grep
```

```
# Ensure all images across all bands have consistent, common projection
# Since focusing on CONUS, use EPSG:3347 as default
target_projection_crs = ee.Projection('EPSG:3347')
target_scale = 1000  # meters

def reproject_image(image):
    return image.reproject(crs=target_projection_crs, scale=target_scale)

merged_dataset = merged_dataset.map(reproject_image)
```

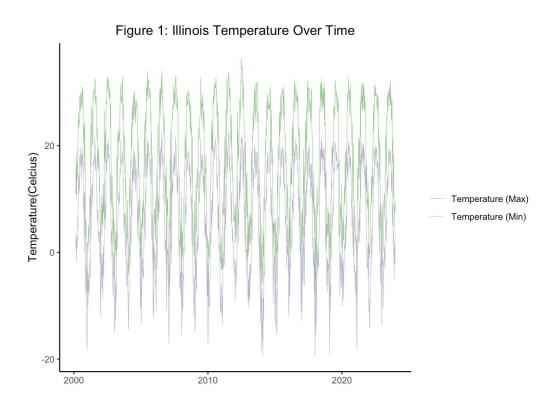
Example of OOP Paradigm Usage

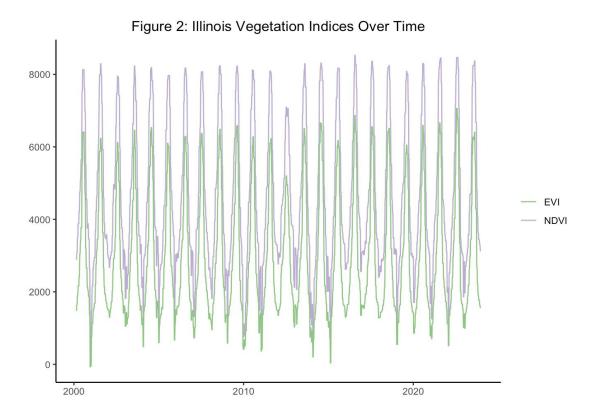
```
class GeoTIFFImage:
   Represents a GeoTIFF image with core attributes and data.
   def __init__(self, filepath):
        Initializes a GeoTIFFImage object.
       Args:
         filepath: Path to the GeoTIFF file.
       with rasterio.open(filepath) as src:
           self.data = src.read()
           self.crs = src.crs
           self.transform = src.transform
           self.width = src.width
           self.height = src.height
           self.count = src.count
           self.band_names = src.descriptions
           self.nodata = src.nodata
        basename = os.path.basename(filepath)
       parts = basename.split(' ')
       self.week number = int(parts[2])
       self.date = parts[3].split('.')[0] # Remove .tif extension
```

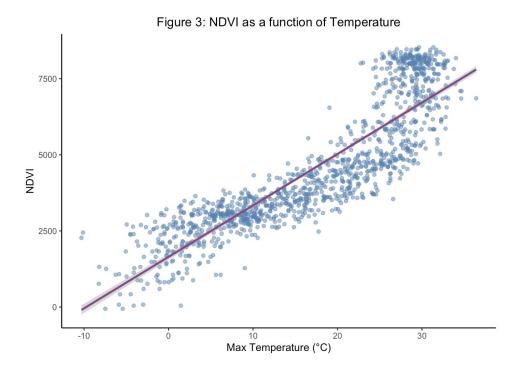
```
def flatten(self):
 Flattens the GeoTIFF data into a pandas DataFrame with specified columns.
  Returns:
   A pandas DataFrame with columns: week, date, lat, long, crop, dayl,
   prcp, srad, swe, tmax, tmin, vp, EVI, NDVI, ET, LE, PET, PLE,
   FPAR, LAI.
 # Create coordinate arrays
 rows, cols = np.meshgrid(np.arange(self.height), np.arange(self.width))
 xs, ys = rasterio.transform.xy(self.transform, rows, cols)
 # Reproject coordinates to lat/lon in degrees (epsq:4326)
 transformer = Transformer.from crs(self.crs, "epsg:4326")
 lats, lons = transformer.transform(xs, ys)
  data = {
      'week': np.full(lons.size, self.week number),
      'date': np.full(lons.size, self.date),
      'lat': lats.flatten(),
      'long': lons.flatten(),
 for i, band name in enumerate(self.band names):
      data[band name] = self.data[i].flatten()
 df = pd.DataFrame(data)
  return df
```

Phase 2: Exploratory Data Analysis

- Data Quality Assessment: checked for data completeness, removed duplicate rows, verified range validity for numeric columns, ensured consistency in units and data types and detected and addressed outliers where necessary.
- **Data Imputation:** Imputed missing values using group means and combined preprocessed data into a single cohesive dataset for downstream analysis.
- Exploratory Data Analysis (EDA)
 - Analyzed relationships between various features in the dataset.
 - Visualized the relationship between NDVI (the outcome of interest) and other variables to identify patterns.
- Correlation and Feature Selection
 - Computed correlation coefficients between predictors.
 - Removed highly correlated variables to reduce redundancy and avoid multicollinearity.
 - selected key features relevant to the analysis.
- Clustering for Feature Importance
 - Applied K-means clustering on scaled data to group observations and identify important features contributing to the outcome.







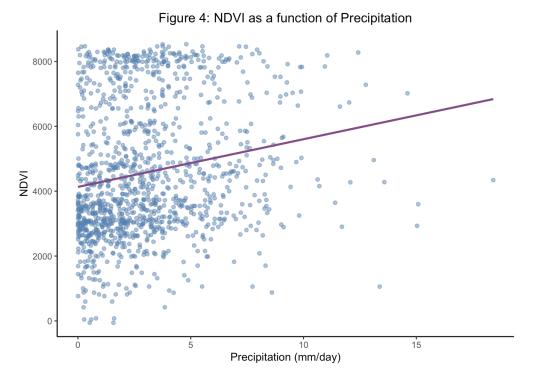
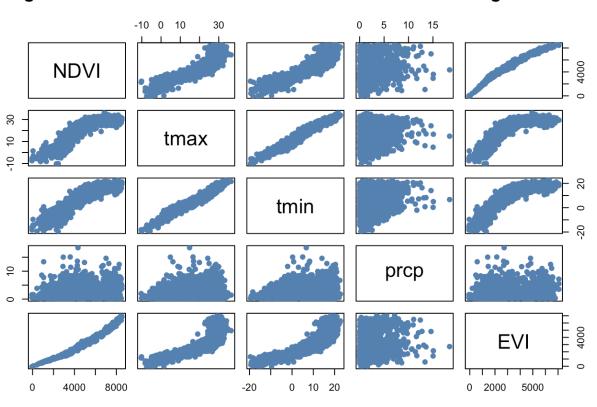
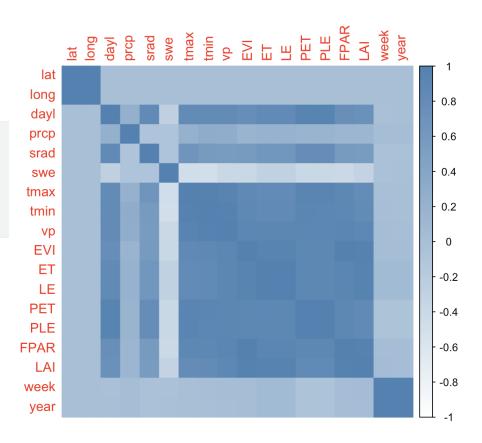


Figure 6: Pairwise Scatter Plots of NDVI and Climate/Veg Variables



```
#highly correlated features (> 0.85)
high_corr <- findCorrelation(corr_matrix, cutoff = 0.85, names = TRUE)
#features that are not highly correlated
selected_features_corr <- setdiff(colnames(numeric_cols), high_corr)
selected_features_corr</pre>
```

```
[1] "lat" "dayl" "prcp" "srad" "swe" "tmax" "LAI" "week"
```



```
# variables with high variance within clusters
feature_variances <- apply(scaled_data, 2, var)
selected_features_kmeans <- names(feature_variances)[order(-feature_variances)[1:15]]
final_selected_features <- selected_features_kmeans
# add NDVI
final_selected_features <- unique(c(final_selected_features, "NDVI"))
cat("Selected Features for Predictive Modeling:\n")</pre>
```

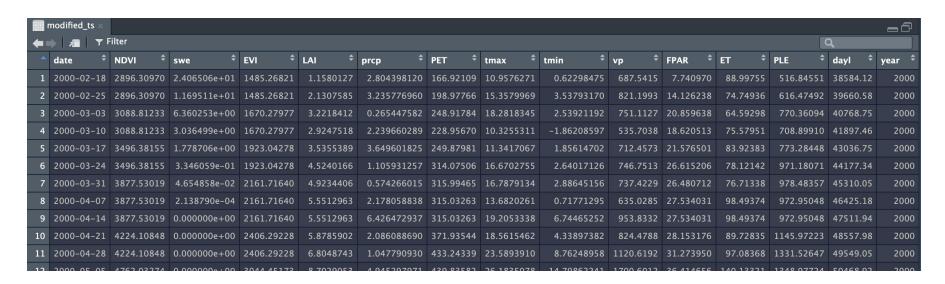
Selected Features for Predictive Modeling:

```
print(final_selected_features)

[1] "swe" "year" "EVI" "LAI" "prcp" "NDVI" "PET" "tmax" "vp" "week"
[11] "FPAR" "ET" "PLE" "tmin" "dayl"
```

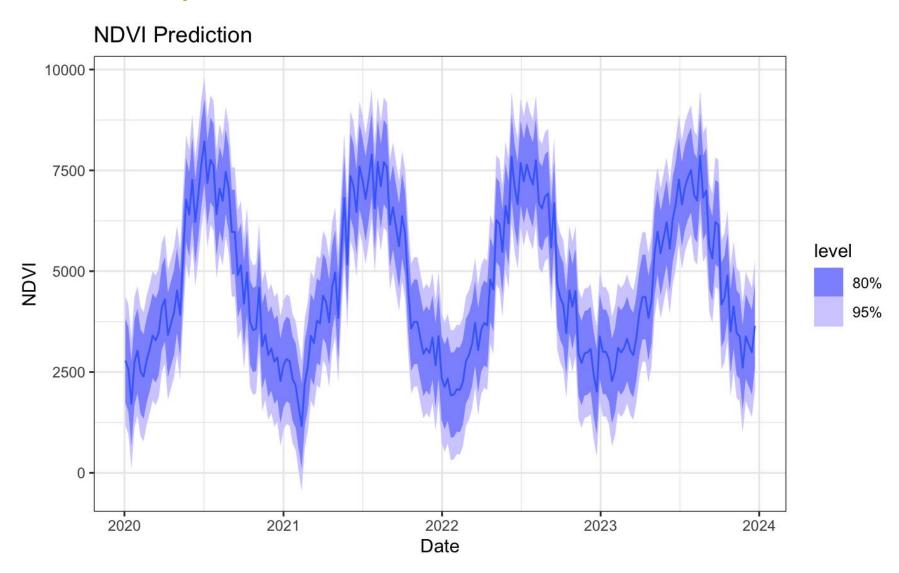
Phase 3: Predictive Modeling

- Predictive modeling to be used for future cropland health
 - Outcomes of interest: Normalized Difference Vegetation Index (NDVI), Enhanced Vegetation Index (EVI), Fraction of Photosynthetically Active Radiation (FPAR), Leaf Area Index (LAI), Evapotranspiration (ET)
 - o **Predictors:** snow water equivalent, precipitation, minimum and maximum temperature, water vapor pressure, and daylength
- Method of Analysis: Time series linear models using different outcomes of interest and predictor variables
 - Use of R packages tsibble and fable

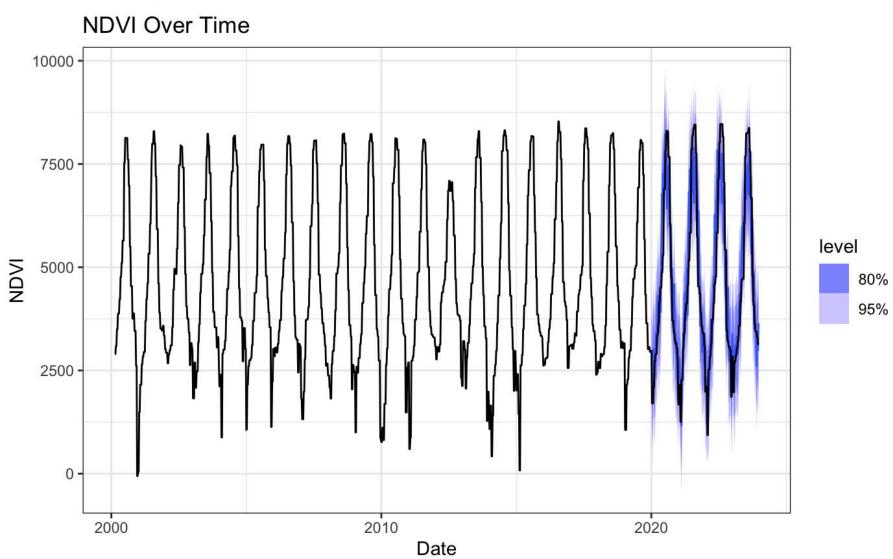




Phase 3: Analysis Results



Phase 3: Analysis Results



Website

- Created a website to explain our analysis and findings.
- Short demo of the website: WEBSITE

Website

