# Software to Streamline Sharing of Agricultural Algorithms and Data

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### Introduction

### Challenges

How to make appropriate data deposition commonplace?

Solution



# **UA Digital Agriculture Group**

Mission: Providing solutions for computing in agriculture, so that we can engineer crops and sustainable agricultural landscapes.

Vision: Faster, more collaborative agricultural science and engineering through shared software and data.



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#### Collaborators, Too Many to List:

Tyson Swetnam, Brian Heidorn, CyVerse, Todd Mockler, Nadia Shakoor, Geoff Morris, Vasit Sagan, Robert Pless, Rob Kooper, Max Burnette, Steve Long

Others to be listed as we go ...

# Outline: Our Approach in 4 Ts

- Tools
- Translators
- Tutorials
- Templates

# Open Software as Germplasm: Aligned with CG and Land Grant Missions







CIMMYT genebank

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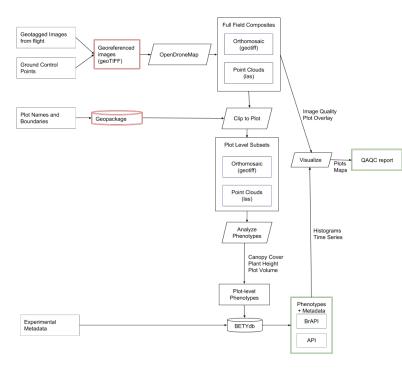




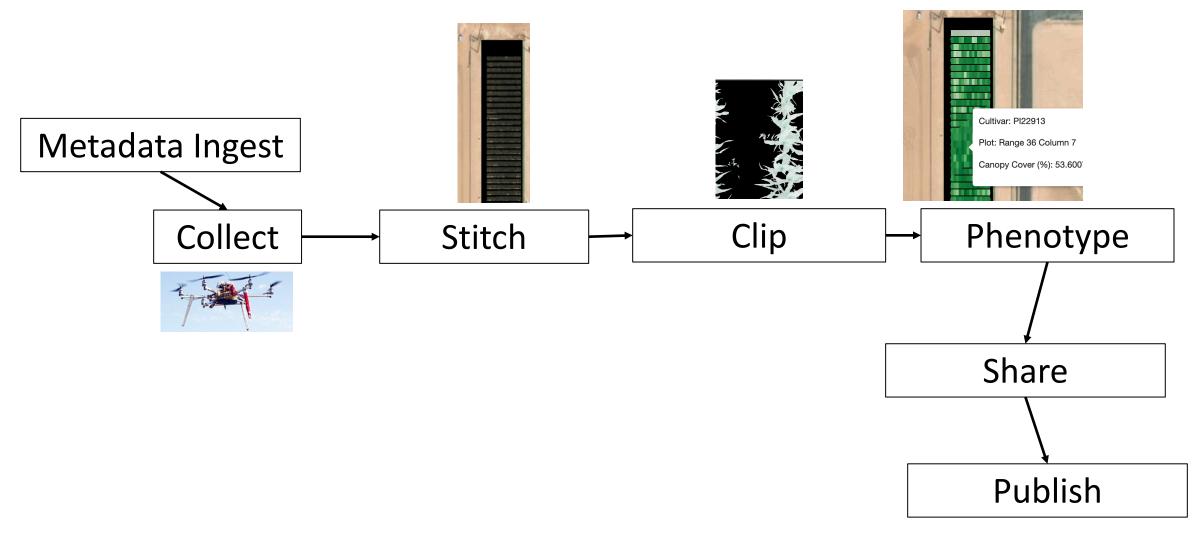




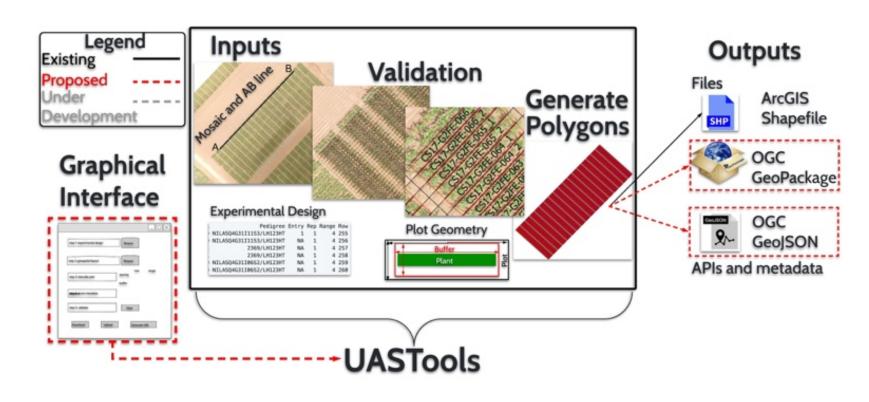
# High Throughput Phenomics (images → phenotypes) Pipelines



### **A Simple Drone Pipeline**



# **Metadata Ingest: Plot Boundaries**

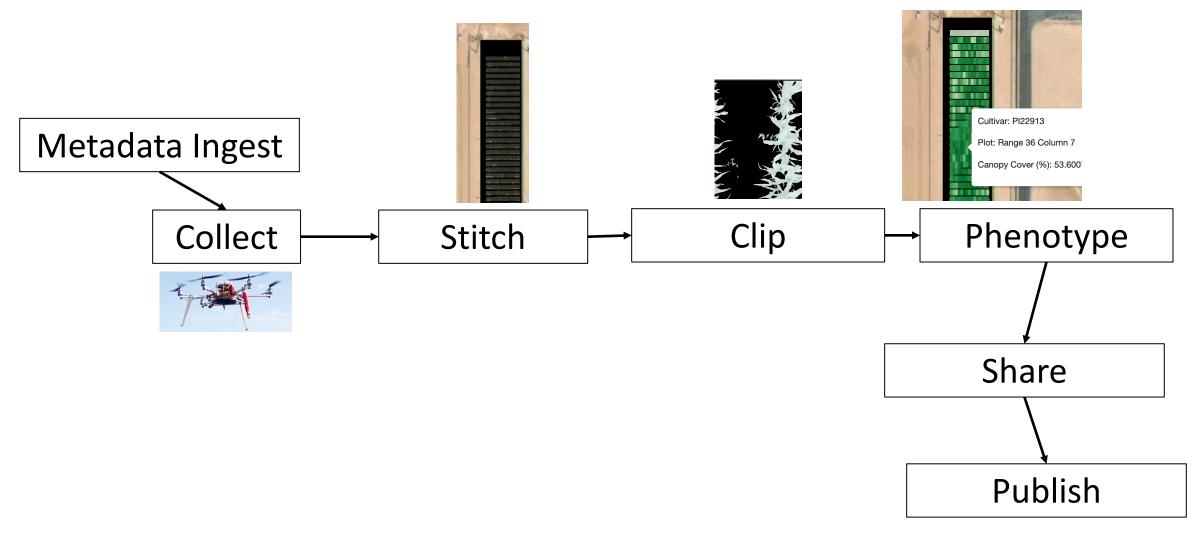


Works

Open Source ✓\*

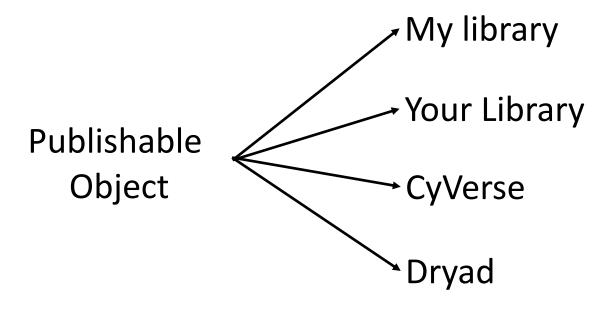
Standard API ✓\*

### **A Simple Drone Pipeline**



#### Full Field Composites Orthomosaic (geotiff) Point Clouds (las) Image Quality Clip to Plot Plot Overlay Plot Level Subsets Visualize Plots Orthomosaic Maps (geotiff) Point Clouds (las) Analyze Histograms Phenotypes Time Series Canopy Cover Plant Height Plot Volume Plot-level Phenotypes Phenotypes + Metadata **BrAPI** BETYdb E SCIEN**API** COLLEGE OF AGRICULTURE AND LIF

### **Data Publication**

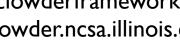


#### **Data Publication**



#### Name My Publication Creator(s): 6 One or more creators are required SELECT REPOSITORY Edit Metadata Note: The links in this page redirect to the live objects. Add metadata Select field No metadata available for this resource **SUBMIT TO Select Repository** REPOSITORY > **Candidate Repositories** The results below are based on an analysis of the dataset's properties and metadata and the preferences you specified. Inter-university Consortium for Political and Social Research Match: all requirements are satisfied.

terraref.org/clowder/



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### **Translators**

HOW STANDARDS PROLIFERATE: (SEE: A/C CHARGERS, CHARACTER ENCODINGS, INSTANT MESSAGING, ETC.)

SITUATION: THERE ARE 14 COMPETING STANDARDS.



SOON: SITUATION: THERE ARE 15 COMPETING STANDARDS. Alternative to a 'new' standard:

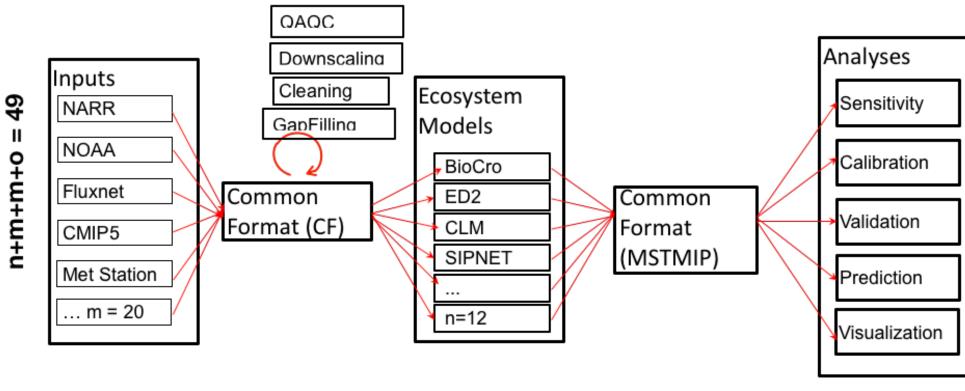
Combine, translate among existing standards.

# **Ecosystem Models Without Standards**

Converter Inputs Analyses Ecosystem NARR  $n^*m+m^*k = 300$ Models Sensitivity NOAA BioCro Calibration Fluxnet ED2 CLM CMIP5 Validation SIPNET Met Station Prediction ... m = 20 n=12 Visualization



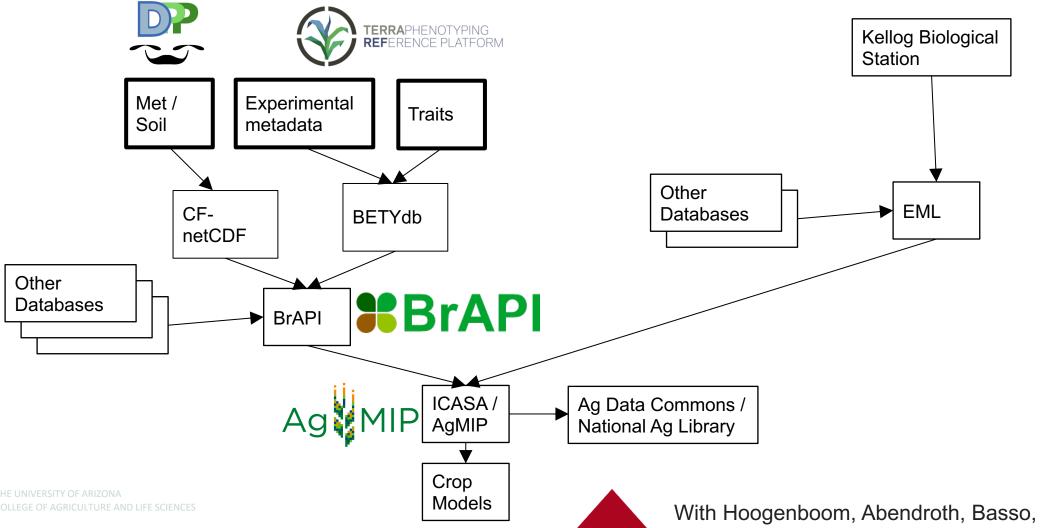
# PEcAn + Standards: benefits of interoperability







# Agricultural Research Data Network (ARDN)

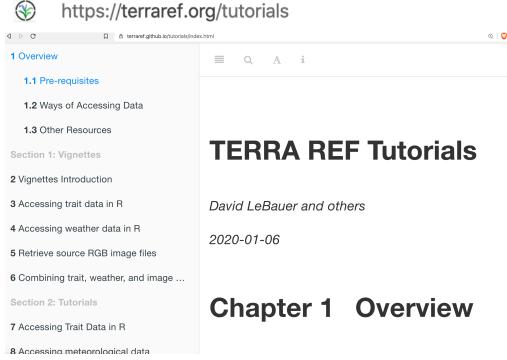




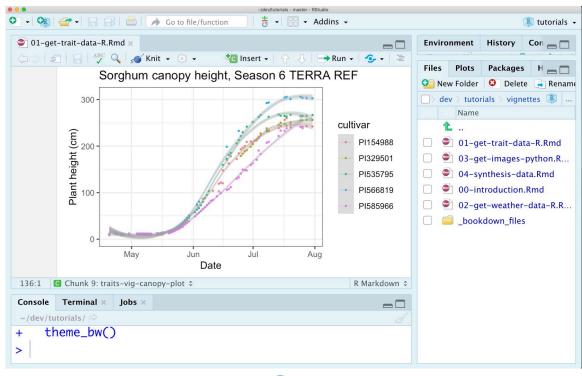
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# Tutorials: Self Guided, Live Coding Webinars, Videos

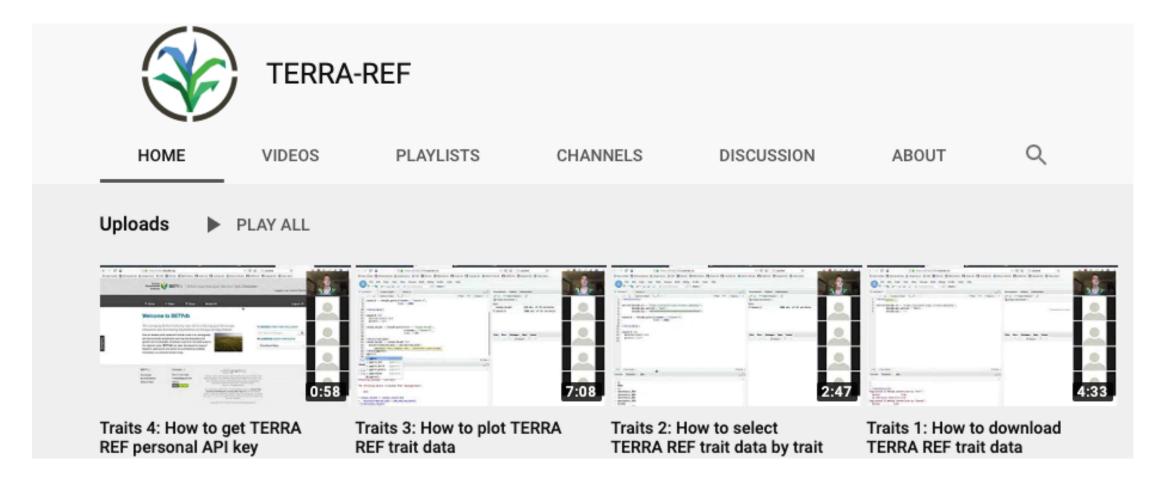


Kristina Riemer, UA





### YouTube Videos





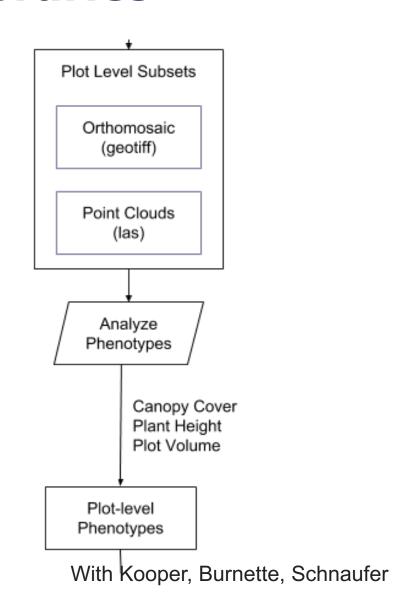
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### **'Transformer' libraries**

vI: github.com/terraref

v2: github.com/agPipeline



# **Templates**

### Guide users to best practices

- Standard formats and vocabularies
- Best practices
- Testing
- Documentation

#### Minimize Overhead

- Quickly integrate a new algorithm into a pipeline
- Versioning
- Sharing

## **Example template**

```
def calculate(pxarray: np.ndarray):
    """Calculates one or more values from plot-level RGB data
    Arguments:
        pxarray: Array of RGB data for a single plot
    Return:
        Returns one or more calculated values
    111111
    # ALGORITHM: replace the following lines with your algorithm
    channel_size = pxarray[:, :, 1].size
    # RETURN: replace the following return with your calculated values.
    return channel_size
   github.com/AgPipeline/template-rgb-plot
   algorithm rgb.py
                                                       Schnaufer
```

# A Template for Plot level RGB data



- 1. Setup: Click the Use this template button in GitHub to make a copy of this repository (or run git clone)
- 2. Definitions: Fill in and modify the definitions in the algorithm\_rgb.py file
- 3. Algorithm: Replace the code in the calculate function with your algorithm
- 4. Test: Run the testing.py script to run your algorithm and validate the results
- 5. Generate: Run generate.py to create a Dockerfile
- 6. Docker: Create a Docker image for your algorithm and publish it
- 7. Finishing: Finish up your development efforts

### **Conclusions**

Reuseable software and data will accelerate science and engineering

Tools, Translators, Templates, and Tutorials enable open, synthetic science

#### **More Information**

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