

## Data From: TERRA-REF, An Open Reference Data Set From High Resolution Genomics, Phenomics, and Imaging Sensors

David LeBauer <sup>1</sup>	Maxwell A. Burnette <sup>2</sup>	Jeffrey Demieville <sup>3</sup>	Noah Fahlgren <sup>4</sup>
Andrew N. French <sup>5</sup>	Roman Garnett <sup>6</sup>	Zhenbin Hu <sup>7</sup>	Kimberly Huynh <sup>8</sup>
Rob Kooper <sup>9</sup>	Zongyang Li <sup>10</sup>	Maitiniyazi Maimaitijiang <sup>11</sup>	Jerome Mao <sup>12</sup>
Todd C. Mockler <sup>13</sup>	Geoffrey Morris <sup>14</sup>	Maria Newcomb <sup>15</sup>	Michael J Ottman <sup>16</sup>
Philip Ozersky <sup>17</sup>	Sidike Paheding <sup>18</sup>	Duke Pauli <sup>19</sup>	Robert Pless <sup>20</sup>
Wei Qin <sup>21</sup>	Kristina Riemer <sup>22</sup>	Gareth Scott Rohde <sup>23</sup>	William L. Rooney <sup>24</sup>
Vasit Sagan <sup>25</sup>	Nadia Shakoor <sup>26</sup>	Abby Stylianou <sup>27</sup>	Kelly Thorp <sup>28</sup>
Richard Ward <sup>29</sup>	Jeffrey W White <sup>30</sup>	Craig Willis <sup>31</sup>	Charles S Zender <sup>32</sup>

- <sup>1</sup>College of Agriculture and Life Sciences, University of Arizona, orcid:0000-0001-7228-053X
- <sup>2</sup>NCSA, University of Illinois, orcid:0000-0002-1786-3676
- <sup>3</sup>University of Arizona, orcid:0000-0002-7725-7379
- <sup>4</sup>Donald Danforth Plant Science Center, orcid:0000-0002-5597-4537
- <sup>5</sup>USDA ARS, orcid:0000-0002-4018-1817
- <sup>6</sup>Washington University in St. Louis, orcid:0000-0002-0152-5453
- <sup>7</sup>Kansas State University, orcid:0000-0002-1500-1255
- <sup>8</sup>University of Arizona, orcid:0000-0003-3360-4073
- <sup>9</sup>NCSA / University of Illinois, orcid:0000-0002-5781-7287
- <sup>10</sup>DDPSC, orcid:0000-0002-8601-378X
- <sup>11</sup>Saint Louis University, orcid:0000-0001-6153-1583
- <sup>12</sup>University of California, Irvine, orcid:0000-0003-2505-8666
- <sup>13</sup>Donald Danforth Plant Science Center, orcid:0000-0002-0462-5775
- <sup>14</sup>Kansas State University, orcid:0000-0002-3067-3359
- <sup>15</sup>Currently USDA Fores Service, Forest Health Protection. Formerly University of Arizona School of Plant Sciences, orcid:0000-0003-2974-9149
- <sup>16</sup>University of Arizona, orcid:0000-0003-0763-3073
- <sup>17</sup>Danforth Plant Science Center, orcid:0000-0002-8942-0524
- <sup>18</sup>Saint Louis University, orcid:0000-0003-4712-9672
- <sup>19</sup>University of Arizona, orcid:0000-0002-8292-2388
- <sup>20</sup>George Washington University, orcid:0000-0001-5775-8216
- <sup>21</sup>Amazon.com, orcid:NA
- <sup>22</sup>University of Arizona, orcid:0000-0003-3802-3331
- <sup>23</sup>University of Illinois at Urbana-Champaign, orcid:0000-0001-9030-0936
- <sup>24</sup>Texas A&M University, orcid:0000-0001-7953-1856
- <sup>25</sup>Geospatial Institute, Saint Louis University, 3694 West Pine Mall, St. Louis, MO 63108, USA, orcid:0000-0003-4375-2096
- <sup>26</sup>Donald Danforth Plant Science Center, orcid:0000-0002-2035-7117
- <sup>27</sup>Saint Louis University, orcid:0000-0002-4387-028X
- <sup>28</sup>USDA Agricultural Research Service, orcid:0000-0001-9168-875X
- <sup>29</sup>none, orcid:0000-0003-4436-0019
- <sup>30</sup>University of Florida, orcid:0000-0002-2051-4645
- <sup>31</sup>University of Illinois at Urbana-Champaign, orcid:0000-0002-6148-7196
- <sup>32</sup>University of California, Irvine, orcid:0000-0003-0129-8024

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## Abstract

The ARPA-E funded TERRA-REF project is generating open-access reference datasets for the study of plant sensing, genomics, and phenomics. Sensor data were generated by a field scanner sensing platform that captures color, thermal, hyperspectral, and active fluorescence imagery as well as three dimensional structure and associated environmental measurements. This dataset is provided alongside data collected using traditional field methods in order to support calibration and validation of algorithms used to extract plot level phenotypes from these datasets.

Data were collected at the University of Arizona Maricopa Agricultural Center in Maricopa, Arizona. This site hosts a large field scanner with fifteen sensors, many of which are capable of capturing mm-scale images and point clouds at daily to weekly intervals.

These data are intended to be re-used, and are accessible as a combination of files and databases linked by spatial, temporal, and genomic information. In addition to providing open access data, the entire computational pipeline is open source, and we enable users to access high-performance computing environments.

The study has evaluated a sorghum diversity panel, biparental cross populations, and elite lines and hybrids from structured sorghum breeding populations. In addition, a durum wheat diversity panel was grown and evaluated over three winter seasons. The initial release includes derived data from from two seasons in which the sorghum diversity panel was evaluated. Future releases will include data from additional seasons and locations.

The TERRA-REF reference dataset can be used to characterize phenotype-to-genotype associations, on a genomic scale, that will enable knowledge-driven breeding and the development of higher-yielding cultivars of sorghum and wheat. The data is also being used to develop new algorithms for machine learning, image analysis, genomics, and optical sensor engineering.

# Introduction

## General information

- Title: TERRA-REF, An Open Reference Data Set From High Resolution Genomics, Phenomics, and Imaging Sensors
- Dates of Data Collection: 2017 and 2018
- Geographic Location: Maricopa, Arizona
  - Center of the field: 33.07549° N 111.9749° W
  - Field is approximately 0.4 ha (200 m x 20 m). The scannable area of the field is 22.1 m on East - West axis and 205.5 m on N/S axis.
- Keywords: Sensor, Phenomics, Sorghum, TERRA-REF
- Funding: The work presented herein was funded in part by the Advanced Research Projects Agency-Energy (ARPA-E), U.S. Department of Energy, under Award Number DE-AR0000598. Computational support was provided by the National Center for Supercomputing Applications and XSEDE.

The data pipeline used to process sensor data and generate derived data was described in the [Burnette et al. \(2018\)](#) ACM PEARC 2018 proceedings paper “TERRA-REF data processing infrastructure”.

## Data Use, Sharing, and Access

To cite this dataset: ...

This data publication consists of data and metadata, as described below. Plot level phenotypes, experimental meta-data, and a catalog of large files are available on Dryad. The file catalog provides the location of larger sensor files that can be accessed using the Globus file transfer service.

Because the dataset contains over 400TB of sensor data files, these large files are stored at the National Center for Supercomputing Applications and are made available using the Globus file transfer service.

In addition to hosting an archival copy of data on Dryad, the documentation includes instructions for browsing and accessing these data through a variety of online portals. These portals provide access to web user interfaces as well as databases, apis, and R and Python clients. In some cases it will be easier to access data through these portals using web interfaces and software libraries.

Except where clearly indicated for sensor data, the structure of directories that contain data and meta-data refer to the contents of the Dryad archive. There are four directories: **data**, **metadata**, **code**, and **documentation**.

The TERRA-REF documentation is hosted at [docs.terraref.org](https://docs.terraref.org). The section “How to Access Data” provides an overview of methods that can be used to access data beyond what is provided in this repository. There is also a PDF copy of the documentation in the file `docs.terraref.org_2020_04_06.pdf` in the **metadata/** directory.

Tutorials for getting started with TERRA-REF data are available at [terraref.org/tutorials](https://terraref.org/tutorials) and on GitHub at [github.com/terraref/tutorials](https://github.com/terraref/tutorials).

The TERRA-REF YouTube channel hosts 1) video walkthroughs of the tutorials [https://www.youtube.com/channel/UComeQAqYR5aZrXN\\_3K5iFGw](https://www.youtube.com/channel/UComeQAqYR5aZrXN_3K5iFGw) and 2) a playlist of videos related to the project <https://www.youtube.com/playlist?list=PLNgRX4VLed8213stlJp60MvVx2p6VTv6N>.

All data are released to the public domain under the [CC-0 license](#). All original software are licensed with the BSD 3-clause or MIT/BSD compatible license. All software used for data processing have been archived on Zenodo and are available on GitHub in the **terraref** organization: [github.com/terraref](https://github.com/terraref).

This software was created specifically for the field scanner data processing pipeline that is described by [Burnette et al. \(2018\)](#). The file `code/source_code_dois.txt` provides the DOIs for code archives that contain the state of the software at the time the data were processed.

Component	Github Organization / Repository	Archive Citation
TERRA-REF Documentation	<a href="#">terraref/documentation</a>	<a href="#">LeBauer et al. (2020b)</a>
Reference Data	<a href="#">terraref/reference-data</a>	<a href="#">LeBauer et al. (2020a)</a>
Sensor Metadata	<a href="#">terraref/sensor-metadata</a>	<a href="#">Willis et al. (2020)</a>
Computing Pipeline	<a href="#">terraref/computing-pipeline</a>	<a href="#">Burnette et al. (2020)</a>
terrautils Python Library	<a href="#">terraref/terrautils</a>	<a href="#">Burnette et al. (2019e)</a>
Metadata Processing	<a href="#">terraref/extractors-metadata</a>	<a href="#">Burnette et al. (2019c)</a>
Laser 3D Scanner	<a href="#">terraref/extractors-3dscanner</a>	<a href="#">Burnette et al. (2019f)</a>
Environmental Logger	<a href="#">terraref/extractors-environmental</a>	<a href="#">Burnette et al. (2019d)</a>
Hyperspectral	<a href="#">terraref/extractors-hyperspectral</a>	<a href="#">Mao et al. (2019)</a>
Multispectral, Thermal, PSII	<a href="#">terraref/extractors-multispectral</a>	<a href="#">Burnette et al. (2019a)</a>
Stereo RGB	<a href="#">terraref/extractors-stereo-rgb</a>	<a href="#">Burnette et al. (2019b)</a>

Other Software used in this project:

Software	Github Organization / Repository	Software Archive
Clowder	<a href="#">clowder-framework/clowder</a>	<a href="#">Marini et al. (2019)</a>
BETYdb Trait Database	<a href="#">pecanproject/bety</a>	<a href="#">Rohde et al. (2016)</a>

## Datasets

In this README, we define the following types of data:

1. **Sensor Data** from five thermal, light, and shape imaging sensors scanning a 4000 m<sup>2</sup> field at hourly to weekly intervals at 1 mm<sup>2</sup> scale resolution. Additional details in the [section on sensors](#) and sensor technical descriptions that can be found in the [metadata/sensors/](#) directory of this repository.
2. **Phenotypes** include both sensor-derived and standard field measurements required to validate and calibrate algorithms that compute plant phenotypes from remote sensing data.
3. **Environmental data** include time series of meteorological variables including temperature, relative humidity, precipitation, wind direction and speed, photosynthetically active radiation, and downwelling spectral radiance.
4. **Genomics** data include whole-genome resequencing data for 384 varieties from the sorghum Bioenergy Association Panel (BAP) and genotyping-by-sequencing data for 768 sorghum Recombinant Inbred Lines (RILs). Contains raw and derived sorghum genome sequencing data. Raw data includes DNA sequence files in compressed FASTQ format. Derived data are available in Variant Call Format (VCF) and Hapmap files.

## Sensor Data

### Field Scanner Sensors

This publication includes the following data generated by sensors. Detailed sensor and system details can be found in the file [metadata/sensors\\_information.zip](#) as well as browsed online through the Clowder Interface [terraref.org/clowder](#) in a space named “Maricopa Agricultural Center Device and Sensor Information”.

The sensor information folder contains extensive documentation for each of the sensors, the field scanner, calibration targets, and the results of sensor validation tests.

Additional sensors not represented in this version of the data are listed in the section on sensors in the section on [additional sensors](#).

Sensor Name	Model	Technical Specifications
<b>Imaging Sensors</b>		
Stereo RGB Camera	Allied Vision Prosilica GT3300C	
Laser Scanner	Custom Fraunhofer 3D)	Spatial Resolution: 0.3 to 0.9 mm
Thermal Infrared	FLIR A615)	Thermal Sensitivity < 50mK @ 30°C
PS II Camera	LemnaTec PS II Fluorescence Prototype)	Illumination 635nm x 4000 $\mu\text{mol}/\text{m}^2/\text{s}$ , Camera 50 fps
<b>Environmental Sensors</b>		
Environmental Sensors	Thies Clima 4.9200.00.000)	
VNIR Spectrometer	Spectral Evolution PSR+3500	Range 350 to 800 nm
Environmental Sensors	Thies Clima 4.9200.00.000)	
VNIR Spectrometer	Spectral Evolution PSR+3500	Range 350 to 800 nm
PAR Sensor	Quantum SQ-300	Spectral Range 410 to 655 nm

## Sensor Data Products

The total size of raw (Level 0) data generated by these sensors is 60 TB. Combined, the Level 1 and Level 2 sensor data products are 490 TB. This size could be substantially reduced through compression and removal of duplicate data. For example, the same images at the same resolution appear in the georeferenced Level 1 files, the full field mosaics, and the plot level clip.

Sensor data are stored on the Storage Condo at the National Center for Supercomputing Applications in Urbana, Illinois. We make them available for download with the Globus file transfer system. The following steps are required to access them: 1) get an account at [globus.org](https://globus.org); 2) search for the **terra-public** endpoint; 3) install the [Globus Personal Connect application](#) and transfer data. Further information is provided in the data access chapter of the [TERRA-REF documentation](#). As an alternative, the data can be provided on hard drives for the cost of supplies, labor, and shipping.

## Sensor Data Catalog

Globus provides the easiest way to navigate the data. This archive also contains a catalog listing all of the files in the dataset. The catalog is one compressed .zip file per season, currently seasons 4 and 6, found in the **data/** directory in files named **season\_[n]\_catalog.zip**. The compressed catalogs are 373 MB total, and expand to 5.4 GB when uncompressed. After uncompressing the catalog files, the catalog contains one directory per data product and one file per day named **[data product]/file\_catalog\_season[n]\_[data product]\_[filetype]\_[YYYY-MM-DD].json**.

These catalog files contain the following information in JSON format:

```
collections: 'collection name [Data Product Name] - [YYYY-MM]'
  datasets: 'dataset name [Data Product Name] - [YYYY-MM-DD]'
    files:
      path: 'season-[n]/sites/ua-mac/Level_[m]/[data_product]/[filename]'
      checksum: '[checksum_string]'
      name: '[data_product]_L[m]_[YYYY-MM-DD]_[Scan Name]'
      size: 'bytes'
```

There is one collection per data product per month, and one dataset per data product per day. This structure of collections and datasets refers to the organization of files in the the Clowder database and web interface [see data access documentation](#).

Below is a summary of the sensor data products included in the first release of TERRA-REF data. Sensor-derived phenotypes described in the [Phenotype Data](#) section were generated from the 3D laser scanner and RGB camera sensors as described in the file `metadata/methods.csv`.

Data Product	Sensor	Algorithm	File Format	Plot Clip	Full Field
Environment	Thies Clima	envlog2netcdf	netcdf	NA	NA
Thermal Image	FLIR	ir_geotiff	geotiff	+	
Point Cloud	Fraunhofer Laser 3D	laser3d_las	las	+	
Point Cloud	Fraunhofer Laser 3D	scanner3DTop	ply		
Images Time-Series	PSII Camera	ps2png	png		
Color Images	RGB Stereo	bin2tiff	geotiff	+	+
Plant Mask	RGB Stereo	rgb_mask	geotiff		x

Figure 1 shows the number of files for each data type across seasons 4 and 6.

### Sensor Data Directory Contents

The following list describes the organization and contents of the Storage Condo server. These can be accessed at the `#terra-public` endpoint on Globus. Directory names have a leading `/` while file names do not.

- **Environment Logger**
  - `/envlog_netcdf`
    - \* Daily aggregated files named `envlog_netcdf_L1_ua-mac_[YYYY-MM-DD].nc`.
    - \* There are also 24 hourly files for each day named `[YYYY-MM-DD_HH-MM-SS]_environmentlogger.nc`.
- **Laser3D**
  - `/laser3d_las`
    - \* One merged file per scan across the short (E-W) axis with names ending in `_merged.las`. There are typically 50-100 of these each day.
  - `/laser3d_las_plot`
    - \* Each directory has the name of one plot, and there is one LAS file clipped to the plot boundaries for each scan (there may be more than one scan per day).
- **RGB Stereo:**
  - `/rgb_geotiff`
    - \* File names ending in `_left.tif` and `_right.tif` represent simultaneous images from left and right stereo pair cameras.
  - `/rgb_mask`
    - \* These images have the soil represented as black pixels. For each file ending in `*_left_mask.tif` in the RGB Geotiff dataset, an image with black pixels representing areas that contain soil and not plants.
  - `/rgb_geotiff_plots`
    - \* For each RGB Geotiff image, a Geotiff file with the same dimensions as the plot. It contains the image clipped to the plot boundaries as well as fill values for parts of the plot not in the image.
  - `/rgb_fullfield`
    - \* Key data product is one full resolution full-field image per scan.
    - \* Other files include: lower resolution versions of the full field (files with names ending in `_10pct.tif`, `_thumb.tif` and `.png`); CSV files containing canopy cover values for each plot; a JSON file listing images contained in the fullfield mosaic; a VRT file that is a “virtual geotiff” that was used to generate the full-field mosaic.
    - \* These full field Geotiff images are RGB images and image masks tiled together to make up a full-field view. These full field images are *not* orthomosaics since they are not stitched together because doing so causes geometric aberrations.

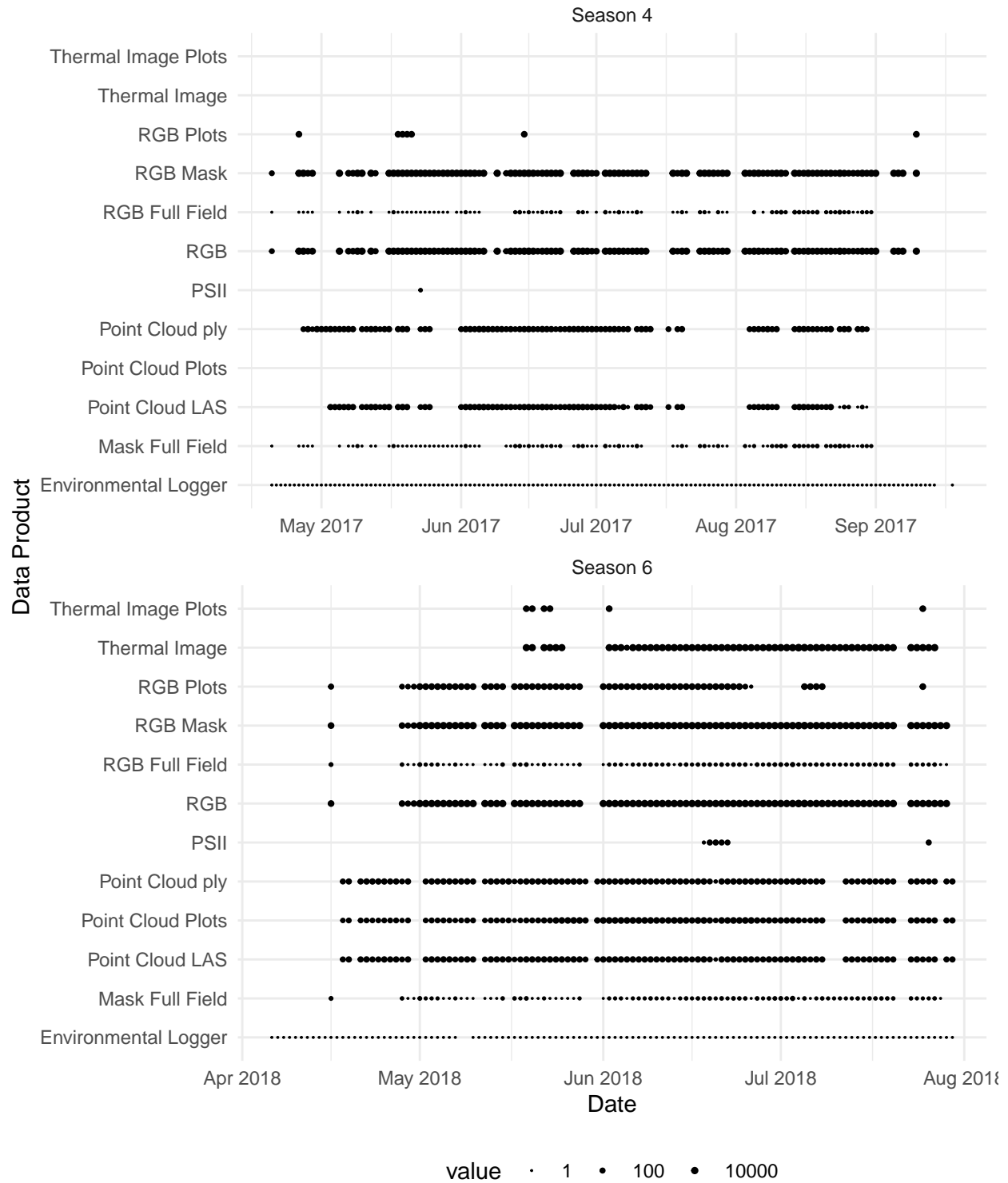


Figure 1: Counts of individual files for each type of sensor data product.



- **PSII Camera:**

- /ps2\_png:
  - \* 101 .png files per folder. The order of the images is indicated by the last four digits of the file name, i.e. \_0000.png to \_0100.png.
  - \* 101 georeferenced Geotiff files otherwise identical to the PNG counterparts.
  - \* These files represent a time series of images captured at a rate of 50 frames per second.

### Sensor Data Directory Structure and File Naming

File names follow consistent patterns based on data product and date of collection. These are intended to be easily understood. The hierarchy is season, data product level, date, date-time, and files.

Specifically, under the directory /season\_[4,6]/sites/ua-mac we have:

```
| -Level
|   | - Data product name
|   |   | -Date (YYYY-MM-DD)
|   |   |   | -Date-Time (YYYY-MM-DD__HH-MM-SS-SSS)
|   |   |   |   | - File name(s)
```

e.g.

```
| -Level_1
|   | - envlog_netcdf
|   |   | -2017-08-26
|   |   |   | - 2017-08-26_13-34-54-321_environmentlogger.nc
|   | - laser3d_las
|   |   | -2017-08-26
|   |   |   | -2017-08-26__12-34-54-321
|   |   |   |   | - scanner3DTop - 2017-08-26__12-34-54-321 MergedPointCloud.las
```

For convenience, we have pre-processed some images and point clouds to plot boundaries and have organized them by Date and then Plot name, e.g.:

```
| -Level_1_plots
|   | - rgb_geotiff
|   |   | - 2017-04-26
|   |   |   | - MAC Field Scanner Season 4 Range 21 Column 16
|   |   |   |   | - rgb_geotiff_L1_ua-mac_2017-04-26__12-56-14-907_right.tif
|   |   |   |   | - rgb_geotiff_L1_ua-mac_2017-04-26__12-56-14-907_left.tif
|   |   |   |   | - rgb_geotiff_L1_ua-mac_2017-04-26__12-53-34-106_right.tif
```

## Phenotype Data

### Raw Phenotype Data

Tables of phenotypes can be found in the compressed files `data/season_4_traits.zip` and `data/season_6_traits.zip`. Once uncompressed, each directory will contain one CSV file for each combination of trait and measurement method. The names of these CSV files help identify the contents because they follow the pattern `season_[n]_[trait]_[measurement_type].csv`. For example, the file `season_6_aboveground_biomass_manual.csv` contains manual measurements of above-ground biomass taken during season 6. Figure @ref(fig:trait\_counts) provides a summary of available traits and the dates on which they were collected.

These CSV files have one measurement per row for a specific date, location, genotype, and measurement. The first line is a header that contains the names of the fields:

- **plot** (text) Plot name, using the format `<field site> Season <n> Range <m> Column <k>`.
- **scientificname** (text) Latin name for the crop species. This will always be *Sorghum bicolor* until future versions with data from additional crops are published.
- **genotype** (text) Genotype or accession identifier.
- **treatment** (text) Name of experimental treatment.
- **date**: (YYYY-MM-DD) Date of measurement.
- **trait**: (text) Name of the trait measured. Defined in the file `metadata/variables.csv`.
- **method**: (text) The method used to measure the trait. Defined in the file `metadata/methods.csv`.
- **mean**: (numeric) Value of the phenotype data.
- **checked**: (boolean) 0 = unchecked and 1 = checked: has the data been independently reviewed?
- **author**: (text) name of scientist who collected the data or who wrote the algorithm used to derive phenotypes from sensor data.
- **season**: (text) Name of season: one of ‘Season 4’ or ‘Season 6’.
- **method\_type**: (text) Type of measurement: one of ‘manual’ or ‘sensor’.

## Code

For the purposes of reproducibility, the code used to run and query data from a copy of the TERRA REF trait database [terraref.org/bety](http://terraref.org/bety) is provided. This uses Docker to run the database and R to query the data and prepare the CSV files. The `code/` directory contains the following:

- An R script `s4s6_sql2csv.R` that connects to the database and generates the phenotype data files described above. This requires a copy of the database. Instructions for installing the database using Docker are found in `betydb_docker`. A much slower implementation that uses the API and the R traits package [Chamberlain et al. \(2019\)](#) is provided in `code/s4s6_slow_api.R`.
- The `betydb_docker/` folder that contains files named `Docker`, `docker-compose.yml`, `README.md` and `.pgpass`. This allows a user to run a copy of the TERRA REF PostgreSQL database containing phenotype data and agronomic metadata. Instructions are in the `README.md` file and database connection parameters are in the `.pgpass` file. This database can be queried using the included R script or from any programming language using the credentials stored in the `.pgpass` file.

## Environmental Data

### Weather

We have data from two stations, AZMet and the field scanner system.

### Field Scanner System “Environment Logger”

These data are from the Theis-Clima weather station mounted on the field scanner. The Environment Logger has more sensors and provides high temporal resolution data than the weather station. In the NetCDF files, data is recorded at five seconds intervals. This logger also collects the downwelling solar radiation every five seconds at a spectral resolution of 0.5 nm. The sensors and data collection frequency are described in the [section on sensor data](#). We describe both the full resolution data provided as NetCDF files in the sensor data product named “`envlog_netcdf`” and the 5-minute aggregated data provided here as JSON files that were accessed using the Geostreams API as described in the TERRA REF tutorials. These time series are not continuous, as can be seen in [Figure 3](#).

### Environment Logger Aggregated JSON Files

The files `metadata/weather/envlog_aggregated/season_[n]_envlog.json` were downloaded from the Geostreams API on April 21 2020. The API is described in the tutorials, and the specific calls used to download these files are in `code/envlog_curl.sh`.

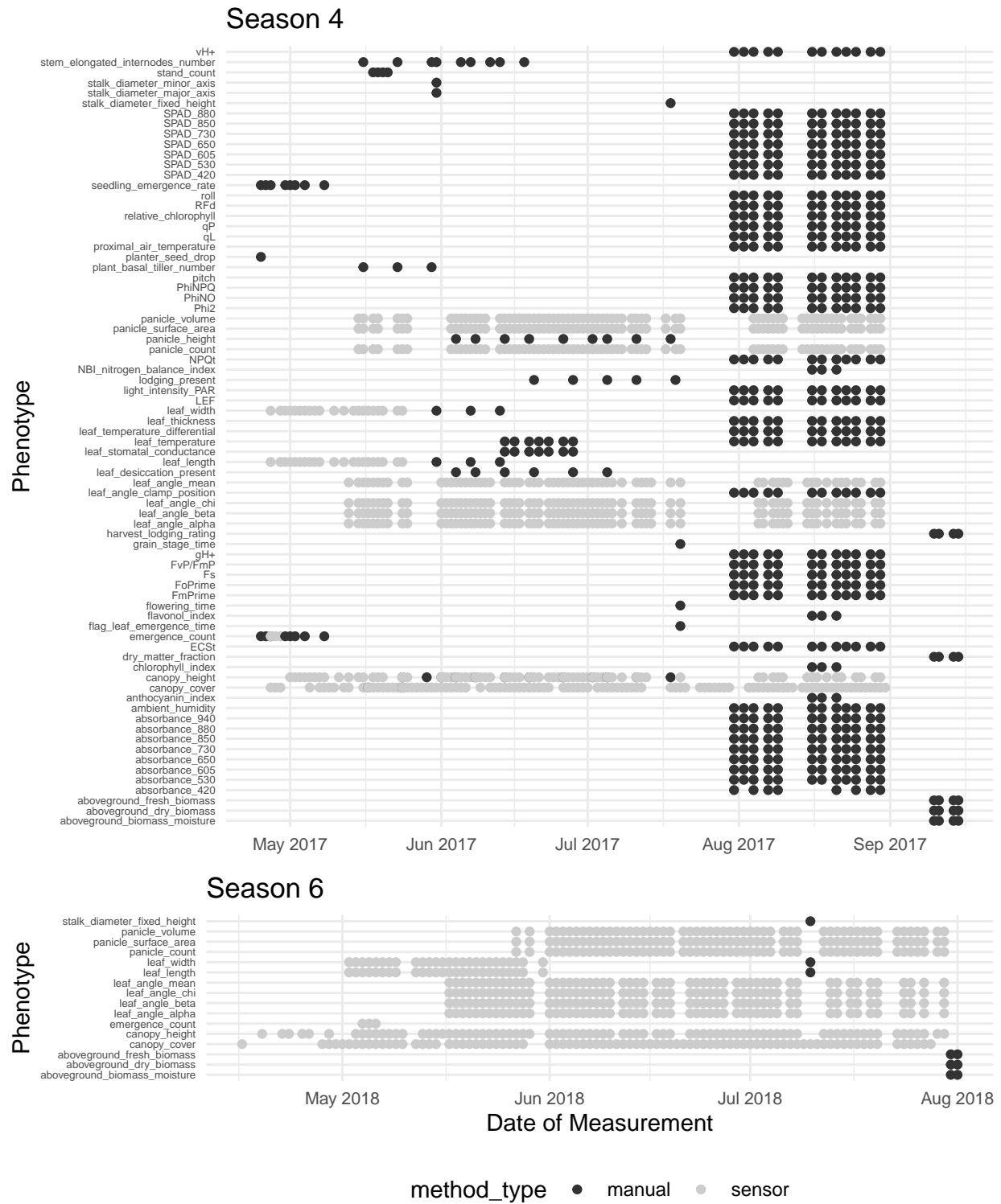


Figure 2: (#fig:trait\_counts)Counts of individual plot level measurements for each trait.

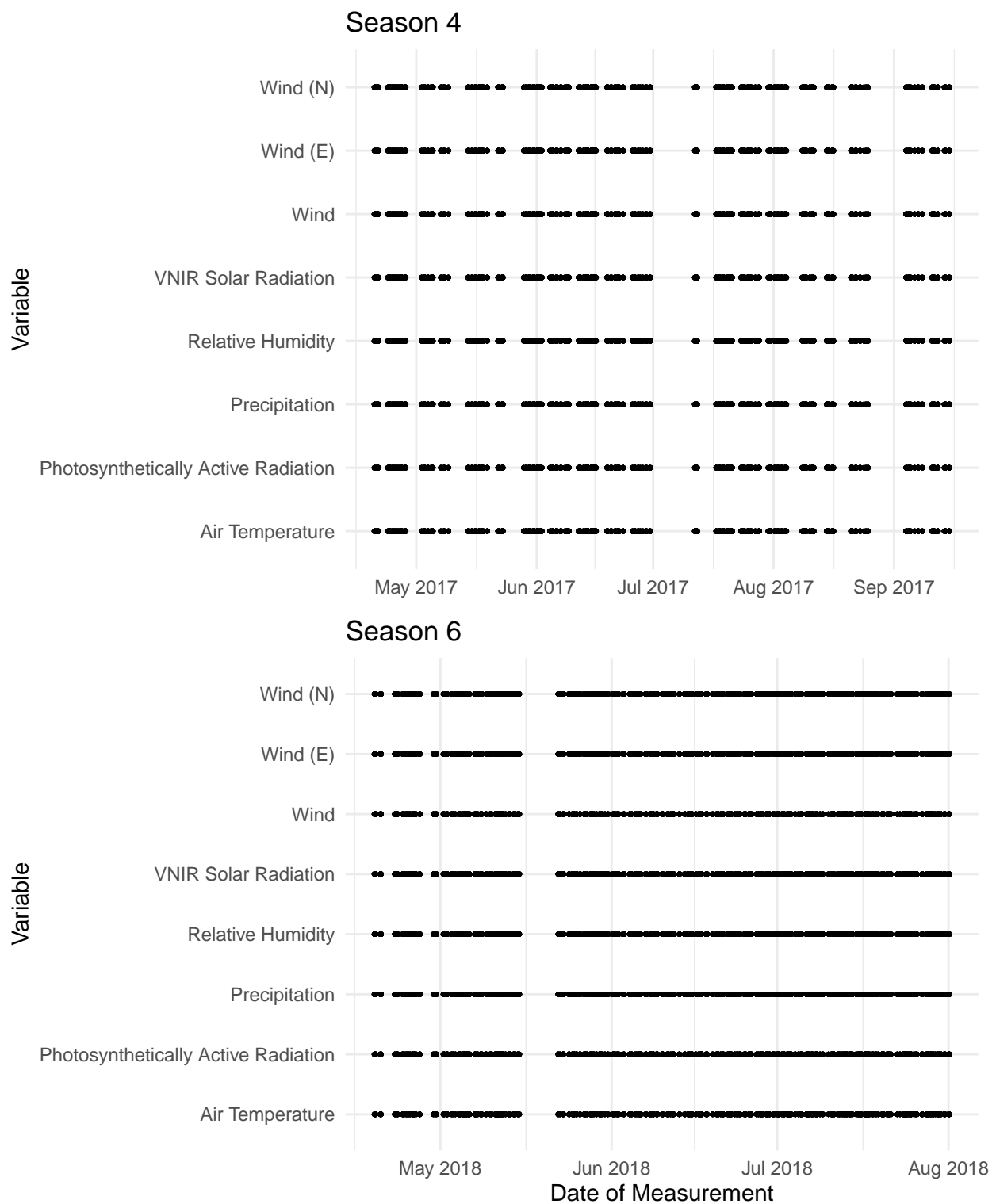


Figure 3: Dates of aggregated weather data - available through the Geostreams API and in the `metadata/weather` folder.

## Environment Logger NetCDF Files

These are even higher resolution time series.

The files for each season are approximately 40 GB and, like sensor data, are provided using methods such as Globus transfer or the Clowder API (see “How To Access Data” in the [documentation](#)). On Globus, they are in the `/ua-mac/Level_1/envlog_netcdf/` directory.

The data are described in the TERRA REF documentation under the environmental data section of the [data products chapter](#).

The primary reason that these files are so large is that they contain the downwelling spectral radiance measurements (350-800 nm) at five second intervals.

On Globus, each date will include hourly files named `[YYYY-MM-DD_HH-MM-SS]_environmentlogger.nc` and as a single daily file named `envlog_netcdf_L1_ua-mac_[YYYY-MM-DD].nc`. The daily file contains all of the hourly files concatenated into a single file for ease of use.

## AZMET

These gap-filled and corrected data have been provided for convenience. When using these data, **users must cite** [Brown and Russell \(1996\)](#): > Brown, P. W., & Russell, B. (1996). AZMET, The Arizona Meteorological Network. Arizona Cooperative. website: <https://cals.arizona.edu/AZMET/>.

Both hourly and daily values are available, and can be found in the `/metadata/weather/azmet` folder of the Dryad repository. There are four CSV files – one daily and one hourly for each season, and a README file that provides a description of their content. These files have not been modified from the collection on the website other than to subset it to the time range of the two seasons of sensor and phenotype data.

## Soils Data

A comprehensive analysis of soil physical properties will be published alongside the TERRA-REF datasets ([Babaeian et al., 2020](#)).

## Genomics Data

The Biomass Association Panel planted in Seasons 4 and 6 was described by [Brenton et al. \(2016\)](#). These genotypes have been sequenced and the sequence data and SNPs are provided as an archive on Dryad. A copy of these data is available on the CyVerse Data Store <https://datacommons.cyverse.org/browse/iplant/home/shared/terraref>.

The genomics dataset includes raw and derived sorghum genome sequencing data from the TERRA-REF project. Raw data includes DNA sequence files in compressed FASTQ format. Derived data is available for whole-genome resequencing and genotyping-by-sequencing.

The data is contained in the following directory structure:

```
| -terraref
|   | -genomics
|   |   | -raw_data
|   |   |   | -bap
|   |   |   |   | -resequencing
|   |   |   |   | -ril
|   |   |   |   | -gbs
|   |   | -derived_data
|   |   | -bap
```

```
| | | | | -resequencing
| | | | | -danforth_center
| | | | | -ril
| | | | | -gbs
| | | | | -kansas_state
```

## Germplasm

A list of cultivars and the experiments in which they are planted are listed in the file `metadata/germplasm.csv`.

The `germplasm.csv` file contains the following fields:

- **germplasmName** Genotype identifier used by TERRA REF; typically equal to Genesys Accession Number
- **germplasmPUI** UUID from [Genesys](#)
- **season\_4** A boolean which is *TRUE* if the accession was planted in Season 4 experiments.
- **season\_6** A boolean value which is *TRUE* if the accession was planted in Season 6 experiments.

For user convenience, we also provide this information and additional metadata in a format that follows the Breeder's API (BrAPI) specification v1.3 [Selby et al. \(2019\)](#) in files named `season_[n]_germplasm.json` and in the `terraref_sorghum_bap_samples.txt` table alongside raw sequencing data (described below).

## Whole-genome resequencing data for the sorghum Bioenergy Association Panel (BAP)

### Raw data

There are 384 available samples. Each sample is stored as a pair of files in bzip2 FASTQ format with names ending in `*_R1.fastq.bz2` and `*_R2.fastq.bz2`. R1 and R2 in these names represent read pairs one and two. Each file in a pair has the same number of DNA sequences that line up and are read from opposite ends of the same sequenced molecule.

A table of sample information (`terraref_sorghum_bap_samples.txt`) can be found in the [raw data directory](#) `genomics/raw_data/bap/resequencing`.

Each table contains the following fields:

- **identifier**: The unique genotype identifier that will be used in the derived data outputs
- **accession**: Genotype identifier used by TERRA REF
- **taxid**: Taxonomy identifier from the [NCBI taxonomy database](#)
- **organism common name**: Genbank common name from the [NCBI taxonomy database](#)
- **subspecific genetic lineage rank**: Cultivar
- **subspecific genetic lineage name**: Unique germplasm identifier
- **ploidy**: Source, [Phytozome](#)
- **number of replicons**: Number of chromosomes, source [Phytozome](#)
- **estimated size**: Estimated genome size, source [Phytozome](#)
- **source material identifiers**: UUID from [Genesys](#)
- **sample name**: Internal sample identifier
- **raw file 1**: File name for read-pair 1
- **raw file 2**: Filename for read-pair 2

### Derived data

Data derived from analysis of the raw resequencing data at the Danforth Center (version1) are available as gzipped, genotyped variant call format (gVCF) files and the final combined hapmap file in the [derived data directory](#), `genomics/derived_data/bap/resequencing/danforth_center/version1`.

## Genotyping-by-sequencing (GBS) data for sorghum recombinant inbred lines (RIL)

### Raw data

Raw data are in gzip FASTQ format. There are 768 available samples.

A key file (`Key_ril_terra`) with sample information can be found in the [raw data directory, genomics/raw\\_data/ril/gbs](#).

### Derived data

Combined genotype calls are available in VCF format in the [derived data directory, genomics/derived\\_data/ril/gbs/kansas\\_state/version1](#).

## Future Releases

This data release does not contain all of the data that was collected during the TERRA-REF project. The field scanner was operated as part of the TERRA-REF program with the mission of generating open access data from April 2016 to September of 2019. Although this release focuses on two seasons of data, all raw data and metadata is available in the public domain. Some derived products from the entire project is available for use. We will not put this in the public domain until it has been processed and validated - please contact the authors for more information.

For the first public release of data we have focused on a subset of the data that we collected, processed, and subjected to quality assurance and control. We expect that this initial data release and subsets of these data curated for specific research projects will provide users and funders with sufficient information to justify processing, validating, and storing data from additional seasons and sensors.

Any use of datasets other than those made available under the CC-0 license must adhere to the [data use](#) and [authorship](#) guidelines outlined in the TERRA-REF documentation at [docs.terraref.org](#) and the file [documentation/docs.terraref.org\\_2020\\_04\\_06.pdf](#).

## Additional Seasons Not Included in the Current Data Release

Season	Crop	Experiments	Populations	Planting Date	Harvest
1	Sorghum	Density	BAP, RIL	2016-04-20	2016-07-16
2	Sorghum	Uniformity Trials	Stay Green RILs F10	2016-07-27	2016-12-02
3	Durum Wheat	Late Season Drought	Diversity Panel	2016-12-15	2017-04-05
4	Sorghum			2017-04-13	2017-09-21
5	Durum Wheat		Diversity Panel	2017-11-20	2018-04-05
6	Sorghum		BAP	2018-04-20	2018-08-02
7	Sorghum	Hybrid Uniformity Blocks	Stay Green RILs, Mutants, F2 families	2018-08-23	2018-11-01
8	Durum Wheat	Uniformity Trials	Diversity Panel	2019-01-01	2019-03-31
9 S	Sorghum		GRASSL x RIO RILs	2019-05-01	2019-07-28

Season	Crop	Experiments	Populations	Planting Date	Harvest
9 N	Sorghum		SAP	2019-04-29	2019-09-05

table caption: ”\*RIL = Recombinant Inbred Lines, BAP = Bioenergy Association Panel, SAP = Sorghum Association Panel, Uniformity Trial = same lines planted in strips across field. In season 9 a second field ‘North’ was added, and separate trials were conducted.”

### Additional Sensors

Sensor Name	Model	Technical Specifications
<b>Multi-spectral Radiometers</b>		
Dedicated NDVI Multispectral Radiometer	<a href="#">Skye Instruments SKR 1860D/A</a>	650 nm, 800 nm +/- 5 nm; 1 down, 1 up
Dedicated PRI Multispectral Radiometer	<a href="#">Skye Instruments SKR 1860ND/A</a>	531nm +/- 3nm; PRI = <a href="#">Photochemical Reflectance Index</a>
Active Reflectance VNIR Spectroradiometer	Holland Scientific Crop Circle ACS-430	670 nm, 730 nm, 780 nm
	Ocean Optics STS-Vis	Range: 337-824 nm @ 1/2 nm
<b>Hyper-spectral Cameras</b>		
VNIR Hyper-spectral Imager	<a href="#">Headwall Inspector VNIR</a>	380-1000 nm @ 2/3 nm resolution
SWIR Hyper-spectral Imager	<a href="#">Headwall Inspector SWIR</a>	900-2500 nm @ 2/3 nm resolution
<b>Environmental</b>		
SWIR Spectrometer	<a href="#">Spectral Evolution PSR+</a>	Range 800-2500nm; Installed 2018
Open Path CO <sub>2</sub> Sensor	<a href="#">Vaisala CARBOCAP Carbon Dioxide Probe GMP343</a>	Range: 0-1000 ppm



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