AGROSTAC APIs
This document gives an overview of the available AGROSTAC API services.

Registration
To access the harmonized data in AGROSTAC a user first needs to register. For this purpose a registration form is available. The user is asked to register providing an email address, together with information on organization and function. By registering the user gives WUR the permission to store the email address, function and organisation with the purpose to send AGROSTAC related information.

REGISTRATION FORM:
https://agrostac.wenr.wur.nl/register.html

RESPONSE:
The response returns the following attributes:
- SUCCESS
- MESSAGE

The user will receive an email with a personal access token.

EXAMPLE:
{"success":true,"message":"API registration succeeded, please check your email"}

The user will receive an e-mail with the following content:

    Dear visitor,

    Thank you for your registering for the Agrostac API. 
    Please use this access token for your API requests: xxx
    Example: https://agrostac.wenr.wur.nl/agrostac/crops?accesstoken=xxx

    Best regards

    The agrostac team.

In case the user would like to receive data via a download link, instead of a JSON-based API response, the email address is used to send such download link.

Access token
One service (API request 3), providing general metadata of the available data sets, is fully public and does not need an access token.

Requests for retrieving available crops, available quantities per crop, available locations and requests for retrieving the data itself need a personal access token. See section registration.

Most data are open and made available under the original licenses (DATA_ACCESS = OPEN). The data is accessible via the personal access token received after registration. Other data can only be shared within certain user groups (specific DATA_ACCESS code). The AGROSTAC moderator can grant access on demand.
1. Retrieve crop codes
A list of available crop codes can be requested.

REQUESTS:
https://agrostac.wenr.wur.nl/agrostac/crops&amp;accesstoken=<youraccesstoken>

RESPONSE:
The response is an array of crops in JSON format, with the following attributes per crop:
- CROP_CODE
- CROP_NAME
- DATASET_ACCESS

EXAMPLE:
```json
{"Crops": [
{"crop_code":"SBN","crop_name":"Soy bean (Glycine max)","dataset_access":"OPEN"},
{"crop_code":"OAT","crop_name":"Oats (Avena sativa)","dataset_access":"OPEN"},
{"crop_code":"RYE","crop_name":"Rye (Secale cereale)","dataset_access":"OPEN"}
,...
]}
```

2. Retrieve quantities per crop
For each crop the available quantities (variables) can be requested. The crop is indicated by parameter <CROP_CODE>.

REQUEST:
https://agrostac.wenr.wur.nl/agrostac/cropquantities/&lt;CROP_CODE&gt;?accesstoken=&lt;youraccesstoken&gt;
e.g. https://agrostac.wenr.wur.nl/agrostac/cropquantities/MAZ?accesstoken=<youraccesstoken>

RESPONSE:
The response is an array of quantities in JSON format, with the following attributes (including time and data range coverage):
- CROP_CODE
- QUANTITYCODE
- QUANTITYDESCRIPTIONUK
- QUANTITYDATATYPE
- QUANTITYID
- QUANTITYUNIT
- DATEMAX
- DATEMIN
- VALUemin
- VALUemax
- DATASET_ACCESS

EXAMPLE:
```json
{"Cropquantities": [
```
3. Retrieve description of data sets

The description of one or all data sets can be requested. As this is general request no access token is required.

**REQUESTS:**

https://agrostac.wenr.wur.nl/agrostac/datasets/<datasetid>

or

https://agrostac.wenr.wur.nl/agrostac/datasets

**RESPONSE:**

The response is an array of dataset codes in JSON format, with the following attributes per data set:

- **DATASET_CODE** (a short meaningful unique code)
- **DATASETID** (unique code of data set)
- **SOURCE_URL** (unique persistent identifier to the dataset)
- **WIKI_URL** (reference to web page that explains data curation of the original dataset)
- **TITLE** (a short clear description of the dataset)
- **LICENSE** (license of the original dataset)
- **RELATED_PUBLICATION** (how to cite the original dataset)
- **ORGANIZATION_NAME** (name of organization and country of the original data set)
- **ORGANIZATION_WEB_ADDRESS** (URL of organization of the original data set)

**EXAMPLE:**

```json
[[
  "license":"Creative Commons Attribution 4.0 International License; CC0 - Public Domain
  "wiki_url":"http:\/\/wiki.agrostac.geodesk.nl\/index.php\/ODJAR_DE_WIT_ET_AL_2018","related_publication":"De Wit et al., 2018, Open Data journal for Agricultural Research, vol. 4, pg. 22-27 (10.18174/odjar.v4i0.15925).","organization_name":"Wageningen Environmental Research (WENR), The Netherlands","title":"A dataset of spectral and biophysical measurements over Russian wheat fields","dataset_code":"ODJAR_DE_WIT_ET_AL_2018","source_url":"https:\/\/doi.org\/10.7910\/DVNY CG0UQB","organization_web_address":"http:\\/\/www.wur.nl\/environmental\-research"
];
]]
```

4. Retrieve description of available crop locations

Get description of crop locations.

**REQUESTS:**

https://agrostac.wenr.wur.nl/agrostac/locations2?objecttypecode=CROP_CULTIVATION&accesstoken=<youraccesstoken>

**OPTIONAL PARAMETERS:**

- `lond` (decimal degrees)
- `latdd` (decimal degrees)
- `distancedd` (decimal degrees)
- `email`
The first three optional parameters enable selections for a certain user defined bounding box. The last parameter can be used to make the API response available as CSV-file via a download link sent to the specified e-mail address.

https://agrostac.wenr.wur.nl/agrostac/locations2?objecttypecode=CROP_CULTIVATION&accesstoken=<youraccesstoken>&londd=32&latdd=-13&distancedd=1

RESPONSE:

The response is an array of crop locations in JSON format, with the following attributes per crop location:

- **OBJECTID** (unique ID of location and its observations)
- **NAMEORIG** (unique name of location and its observations)
- **LONGITUDEDD** (decimal degrees)
- **LATITUDEDD** (decimal degrees)
- **GEOG_ACPCETERACY** (indication of the spatial accuracy of a point location)
- **ALTITUDEM** (meters above sea level)
- **DATEMIN**
- **DATEMAX**
- **NUMBEROFDATAROWS**
- **DATASET_CODE** (a short meaningful unique code)
- **DATASET_ACCESS**
- **FIELDMANAGEMENTTYPE** (see Annex A)
- **WATERMANAGEMENTTYPE** (see Annex A)
- **NUTRIENTS MANAGEMENTTYPE** (see Annex A)
- **NUTRIENTSSTYPE** (see Annex A)
- **PESTSDISEASESMANAGEMENTTYPE** (see Annex A)
- **OBJECTIVE** (brief description of the objective of the experiment)

The OBJECTID is required to retrieve data for a certain crop location. The attribute NUMBEROFDATAROWS provides the total number of days available for a specific crop location.

**EXAMPLE:**

```
```

5. **Retrieve description of specific crop location**

Get description of a crop location.

**REQUESTS:**

https://agrostac.wenr.wur.nl/agrostac/cropcultivationlocation/<OBJECTID>?accesstoken=<youraccesstoken>

**RESPONSE:**

The response is an array of crop locations in JSON format, with the following attributes per crop location:

- **OBJECTID** (unique ID of location and its observations)
6. Retrieve crop data for a crop location

Crop data can be retrieved for one crop location identified by the OBJECTID.

REQUEST:


For example:

https://agrostac.wenr.wur.nl/agrostac/cropcultivationdata2/5?accesstoken=<youraccesstoken>&pagenumber=1&pagesize=50

OPTIONAL PARAMETER:

- pagenumber
- pagesize
- email

Retrieval can be organized in parts via paging. Paging works with the parameters pagenumber and pagesize. If both parameters are omitted all the records are returned. Parameter pagenumber activates
the paging (starting from 1 till a maximum value\(^1\)). The default pagesize is 100 and can be changed by passing a different number.

The e-mail parameter can be used to make the API response available as CSV-file via a download link sent to the specified e-mail address.

**RESPONSE:**

The response is an array with data in JSON format, with for each timestamp the following attributes:

- **OBJECTID** Unique identifier of crop location
- **DATASET_ACCESS** Keyword on access
- **DATEMIN** First date of the observation period
- **DATEMAX** Last date of the observation period
- **CANOPY_HEIGHT_M** Canopy height in m
- **CROP_CODE** Crop code
- **CROP_DEV_BBCH** Crop phenological development according to BBCH scale
- **CUL_NAME** Cultivar name
- **CUL_NOTES** Cultivar notes
- **LAIG** Leaf area index of green leaves (area leaves per surface area)
- **LAIT** Leaf area index of green and dead leaves (area leaves per surface area)
- **PLANT_DENSITY_CNT_M2** Planting density
- **SO_DWT_KGHA** Economic product (oven dry wt) in kg/ha, can be from intermediate harvest
- **SO_FWT_KGHA** Economic product (fresh wt) in kg/ha, can be from intermediate harvest
- **SO_MOISTURE_FWT_FR** Moisture content of fresh yield (e.g., grain, fruit, leaves) in kg[water]/kg[harvest]
- **TOPS_DWT_KGHA** Above ground biomass (oven dry wt) in kg/ha, can be from intermediate harvest
- **TOPS_FWT_KGHA** Above ground biomass (fresh wt) in kg/ha, can be from intermediate harvest

**EXAMPLE:**

```json
{"CropCultivationData"):
[{
```

7. Retrieve crop data for specific crop and quantity

To access data for a certain crop and quantity (see API request 2 for available quantities of crop of interest), the following request can be used.

**REQUEST:**

---

\(^1\) The maximum value can be calculated by dividing NUMBEROFDATAROWS of an OBJECTID (from request ../locations?objecttypecode=CROP_CULTIVATION) by the pagesize
For example:


**OPTIONAL PARAMETERS:**

- minlondd (decimal degrees)
- maxlondd (decimal degrees)
- minlatdd (decimal degrees)
- maxlatdd (decimal degrees)
- pagenumber
- pagesize
- email

The first four optional parameters enable selections for a certain user defined bounding box. Retrieval can be organized in parts via paging. Paging works with the parameters pagenumber and pagesize. If both parameters are omitted all the records are returned. Parameter pagenumber activates the paging (starting from 1 till a maximum value). The default pagesize is 100 and can be changed by passing a different number. The email parameter can be used to make the API response available as CSV-file via a download link sent to the specified e-mail address.

**RESPONSE:**

The response is an array with data in JSON format, with for each timestamp the following attributes:

- **OBJECTID** Unique identifier of crop location
- **LON** Longitude (decimal degrees)
- **LAT** Latitude (decimal degrees)
- **DATASETID** Unique identifier of dataset
- **DATEAVG** Average date
- **CROP_CODE** Crop code
- **<QUANTITYCODE>** Value of selected quantity

**EXAMPLE:**

```json
{"CropDataByArea":
["cropcode":"WHB","dateavg":"1982-10-25","tops_dwt_kgha":null,"datasetid":22,"lon":5.5,"lat":52.5,"objectid":1757}
,"cropcode":"WHB","dateavg":"1982-10-25","tops_dwt_kgha":null,"datasetid":22,"lon":5.5,"lat":52.5,"objectid":1755}
,"cropcode":"WHB","dateavg":"1982-10-25","tops_dwt_kgha":null,"datasetid":22,"lon":5.5,"lat":52.5,"objectid":1756}
,...
}]
```

---

2 Average date in case observed timestep spans a larger period than days
Annex A Management codes

During the harmonization of data, we made an assessment of the management in support of the use and interpretation.

The following management types are considered.

- **FieldManagementType = UNKNOWN** (or enter FARMER or FIELD_TRIAL). Extra information on the context of the crop observation. Does the field management relate to a FARMER or a FIELD_TRIAL? (if not specified, UNKNOWN will be set).

- **WaterManagementType = UNKNOWN** (or enter OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or NOT). Extra information on the context of the crop observation. Was the water management OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or NOT (no management/irrigation)? (if not specified, UNKNOWN will be set).

- **NutrientsManagementType = UNKNOWN** (or enter OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or EXCESSIVE or NOT). Extra information on the context of the crop observation. Was the nutrients management OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or EXCESSIVE (more than required by the crop) or NOT (no nutrients)? (if not specified, UNKNOWN will be set).

- **NutrientsNType = UNKNOWN** (or select OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or EXCESSIVE or NOT). Extra information on the context of the crop observation. Was the nutrients N-application OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or EXCESSIVE (more than required by the crop) or NOT (no N-fertilization)? (if not specified, UNKNOWN will be set).

- **NutrientsPType = UNKNOWN** (or select OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or EXCESSIVE or NOT). Extra information on the context of the crop observation. Was the nutrients P-application OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or EXCESSIVE (more than required by the crop) or NOT (no P-fertilization)? (if not specified, UNKNOWN will be set).

- **NutrientsKType = UNKNOWN** (or select OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or EXCESSIVE or NOT). Extra information on the context of the crop observation. Was the nutrients K-application OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or EXCESSIVE (more than required by the crop) or NOT (no K-fertilization)? (if not specified, UNKNOWN will be set).

- **PestsDiseasesManagementType = UNKNOWN** (or select OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM or NOT). Extra information on the context of the crop observation. Was the pest and disease management OPTIMAL or SUBOPTIMAL or EXT_SERV_RECOMM (extension service recommendations) or NOT (no pest and disease control)? (if not specified, UNKNOWN will be set).