# Download biodiversity data with galah:: CHEAT SHEET 🐥 python"



galah is an interface to biodiversity data hosted by the Atlas of Living Australia (ALA). It enables users to locate and download species occurrence records (observations, specimens, eDNA records, etc.), taxonomic information, or associated media such as images or sounds, and to restrict their queries to particular taxa or locations.

## Build a query

No matter what kind of data you want to return in Python, every download query consists of the same building blocks.

- 1. Start a guery with a galah.atlas\_function, which specifies the data your query will
- 2. Modify a query by adding filters, specifying taxa or other options
- 3. Output data to screen or store in variable

## AN EXAMPLE QUERY:

query data = galah.atlas\_occurrences( taxa = "reptilia", filters = ["year>2010", "cl22=Victoria"], Modify a data Data is contained in variable dat a

## FOR MORE PYTHON INFORMATION:

Check out our ALA galah Python package & guides:





Start a

## Download data

### COUNTS

## galah.atlas\_counts()

Return the number of records that match a guery

## Return number of observations



galah.atlas\_counts( taxa="reptilia" filters="year=2020"

## Return number of species



galah.atlas\_counts( taxa="reptilia", filters="year=2020",
group\_by= "species", expand=False. total\_group\_by=True

## Return grouped counts

galah.atlas\_counts( taxa="reptilia",
filters="year=2020",
group\_by="order", expand=False

order	count
Crocodylia	6388
Rhynchocephalia	1
Squamata	29679
Testudines	4304

Must use an ALA-

registered email

Register at

ala.org.au/

## **OCCURRENCES**

-43.1

## galah.atlas\_occurrences()

Return species occurrence records that match a query



ecimalLongitude	decimalLatitude	eventDate	scientificName
-43.2	148.	2023-01-06 12:46:00	Perameles gunnii
-43.1	147.	2022-10-07 10:38:26	Perameles gunnii

2022-09-18 10:13:00

... i = 1,658 more rows ... j = 4 more columns

#### **SPECIES LISTS**

## galah.atlas\_species()

Return species information for each species that match a guery



Must use an ALAregistered email

Register at

kingdom	phylum	class	order	family	genus	species
Animalia	Chordata	Mammalia	Pera melemor phia	Pera melidae	Pera meles	Perameles nasuta
Animalia	Chordata	Mammalia	Pera melemor phi a	Pera melidae	Pera meles	Parameles gunnii
Animalia	Chordata	Mammalia	Pera melemor phi a	Pera melidae	Pera meles	Perameles notina
Animalia	Chordata	Mammalia	Pera melemor phi a	Pera melidae	Perameles	Perameles fasciata
i = 1 more row	,	ļ	ļ	ļ	l .	

## **MEDIA & IMAGES**

## galah.atlas\_media()

Return species occurrence records with associated images, sounds or media that match a query. Return matching records with atlas\_media(), then add collect=True and specify a path to download locally





galah.atlas\_media( taxa="perameles" filters="year>2021",

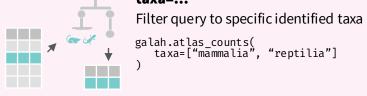
Returns output like atlas\_occurrences() with media metadata



galah.atlas\_media( taxa="perameles" filters="year>2021",
collect=True, path = "path-to-folder"

#### Download media to folder

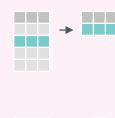
## MODIFY QUERY ON THE SERVER BEFORE DOWNLOADING:



## data profile=...

Apply a set of data quality filters to narrow

galah.galah\_config(data\_profile="ALA") galah.atlas\_counts( use\_data\_profile=True,



## filters=...

Filter query to rows that meet a logical criteria

galah.atlas\_counts( filters="year=2020"

### select=...

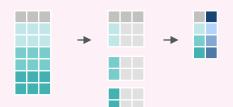
Filter query to return specified columns

galah.atlas occurrences(
 select=["scientificName", "eventDate"]

## group\_by=...

Filter guery to rows that meet a logical criteria

galah.atlas\_counts( group\_by=["year", "species"],





Specify the lowest taxonomic level in a query for a species list

galah.atlas\_species( taxa="fungi", rank="phylum"

## polygon=...|bbox =...

Specify the location of data returned with a polygon or bounding box

bbox = shapely.box(143, -29, 148, -28)galah.atlas counts( hhox = hhox





## Choose an atlas



The Global Biodiversity Information Facility

(GBIF) network consists of a series of 'nodes' — **Living Atlases**—that collate biodiversity data from their own countries. GBIF acts as an umbrella organisation to store data from all nodes.

## galah supports data downloads for 5 Living Atlases and GBIF.

See full list:

https://galah.ala.org.au/Python/galah user gui de/Choosing Atlas.html

```
galah.galah_config(atlas = "Austria")
galah.galah_config(atlas = "GBIF")
```

## Species list in your area

If you want to know what species are in your area, you can use **atlas\_species** to do this.

## Species list for year 2022 in Victoria

For this example, we know that our filters are cl22=Victoria and year=2022. The Python code then looks like this:

```
galah.galah config(email="your-email-here")
species_list = galah.atlas_species(
   filters=["cl22=Victoria","year=2022"]
```

Finally, write data to a csv file.

```
species_list.to_csv("NAME_OF_FILE.csv")
```

## Getting counts of species in your area

If you want to get the species name and counts, we will use the above example to get the initial species list. We will then get the species column, and use that as input for atlas\_counts:

```
species_list_counts=galah.atlas_counts(
  filters=["cl22=Victoria","year=2022"],
  group by="species".
  expand=False,
  counts=True
species_list_counts
```

species	count
Aacanthocnema dobsoni	3
Aades cultratus	8

<sup>...</sup> i = 9,579 more rows

Finally, write data to a csv file.

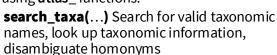
species\_list\_counts.to\_csv("NAME\_OF\_FILE.csv")

## Lookup information — galah provides look-up functions to help users find ways to modify their queries

The living atlases store a huge amount of information, above and beyond only occurrence records. This information can be useful for modifying queries.

### **Taxonomic information**

Look up taxonomic names before downloading data from the ALA using atlas\_functions.



```
galah.search_taxa(taxa=["reptilia", "mammalia"])
```

Specify taxonomic levels in a list using "specificEpithet"

```
galah.search_taxa(
   specific_epithet=["class=aves",
     "family=pardalotidae", "genus=pardalotus",
     "specificEpithet=punctatus"]
```

Specify taxonomic levels in a dictionary using "scientificName"

```
galah.search taxa(
   scientific name={
    "family": ["pardalotidae", "maluridae"],
     "scentificName": ["pardolatus striatus",
                     "malurus cyaneus"]}
```

Search for unique identifiers of a taxon. Identifiers are assigned by atlases to identify all taxonomic clades.

```
galah.search_taxa(
   identifiers=
   "https://id.biodiversity.org.au/node/apni/2914510"
```

## galah.show\_all(type=True)

Show all available options or categories for a specified type of information

## Configuration



## atlases

Show what atlases are available galah.show\_all(atlases=True)
galah.search\_all(atlases="Brazil")



Show what APIs & functions are available

galah.show\_all(apis=True)
galah.search\_all(apis="counts")

### reasons



Show what values are acceptable as "download reasons" for a specified atlas galah.show\_all(reasons=True)
galah.search\_all(reasons="research")

#### Filters fields



Show fields that are stored in an atlas

galah.show\_all(fields=True)
galah.search\_all(fields="state")

## assertions



Show data quality checks run by each atlas

galah.show\_all(assertions=True)
galah.search\_all(assertions="longitude")

### licenses



Show what copyright licenses are applied to media

galah.show\_all(licenses=True)
galah.search\_all(licenses="CC BY")

## galah.search\_all(type="query")

Search for a specific option or category for a specified type of information

## **Taxonomy**

## ranks



Show valid taxonomic ranks (eg Class)

galah.show\_all(ranks)
galah.search\_all(ranks="suborder")

## **Group filters**



### profiles

Show what data profiles (sets of data quality filters) are available

galah.show\_all(profiles=True)
galah.search\_all(profiles="ALA")



Show what species lists are available galah.show\_all(lists=True) ğalah.search\_all(lists="EPBC")

## **Data providers**



providers Show which institutions provide data

galah.show\_all(providers)
galah.search\_all(providers="botanic")

## u

#### collections

Show specific collections within institutions galah.show\_all(providers=True)
galah.search\_all(collections= "antarctic")

## datasets



Show all data groupings collections

galah.show\_all(datasets) galah.search\_all(datasets="river")

## **Values**

Users may wish to see what values are within a chosen field, profile or list to modify a query or understand more about the information of interest. The values functions get their inputs from show\_all(fields=True) or search all(fields="field-ID"). Feed this field ID to show values or search values to see this information.









Specified field

Field values

## show\_values(field="field-ID")

Search for valid taxonomic names, look up taxonomic information, disambiguate homonyms

galah.show values(field="cl22")

field	count
cl22	New South Wales
cl22	Victoria
cl22	Queensland
cl22	South Australia
cl22	Western Australia
cl22	Northern Territory
cl22	Australian Capital Territory
cl22	Tasmania

## search\_values(field="field-ID", value="value")

Search for valid taxonomic names, look up taxonomic information, disambiguate homonyms

galah.search\_ field="cl value="te

	field	count
_values( l22 <b>",</b>	cl22	Northern Territory
erritory"	cl22	Australian Capital Territory

## Want ideas on how to visualise your data?

For more information, including tutorials & examples, check out ALA LABS.



