
arcovid19

Release 0.6b

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Libs is a package to use the COV-19 data analysis tool. It has a module written in python called arcovid19 dedicated to the analysis of all real cases of COVID-19 in Argentina.

Arcovid19 is made up of two classes CasesFrame and CasesPlot, these tools are used to report the analysis of COVID-19 data in Argentina from tables and visualizations. Furthermore, the CasesFrame class inherits and adds functionalities to the DataFrame class of Pandas.

CHAPTER 1

License

arcovid19 is under [The BSD-3 License](#)

The BSD 3-clause license allows you almost unlimited freedom with the software so long as you include the BSD copyright and license notice in it (found in [Fulltext](#)).

Please acknowledge arcovid19 in any research report or publication that requires citation of any author's work. Our suggested acknowledgment is:

The authors acknowledge the arcovid19 project that contributed to the research reported here. <<https://github.com/ivco19/libs/>>

ABOUT THE DATA

Please cite:

Luczywo, N. A., Daza, V., Koraj, M., Dominguez, M., Lares, M., Paz, D. J., Quiroga, R., Rios, M. E. D. L., Sánchez, B. O., Stasyszyn, F., & Cabral, J. B. (2020). Infecciones de COVID-19 en Argentina. Unpublished. <https://doi.org/10.13140/RG.2.2.22519.78246>

BibText:

```
@misc{https://doi.org/10.13140/rg.2.2.22519.78246,  
doi = {10.13140/RG.2.2.22519.78246},  
url = {http://rgdoi.net/10.13140/RG.2.2.22519.78246},  
author = {  
Luczywo, Nadia Ayelen and Daza, Vanessa and Koraj, Mauricio and  
Dominguez, Mariano and Lares, Marcelo and Paz, Dante Javier and  
Quiroga, Rodrigo and Rios, Martín Emilio De Los and  
Sánchez, Bruno Orlando and Stasyszyn, Federico and  
Cabral, Juan Bautista},  
language = {es},  
title = {Infecciones de COVID-19 en Argentina},  
publisher = {Unpublished},  
year = {2020}  
}
```


CHAPTER 3

Raw Data

- Viewer
- CSV
- XLSX

CHAPTER 4

Contact:

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5.1 Installation

This is the recommended way to install arcovid-libs.

5.1.1 Installing with pip

Make sure that the Python interpreter can load arcovid-libs code. The most convenient way to do this is to use virtualenv, virtualenvwrapper, and pip.

After setting up and activating the virtualenv, run the following command:

```
$ pip install arcovid19
```

That should be it all.

5.1.2 Installing the development version

If you'd like to be able to update your corral code occasionally with the latest bug fixes and improvements, follow these instructions:

Make sure that you have Git installed and that you can run its commands from a shell. (Enter *git help* at a shell prompt to test this.)

Check out arcovid-libs main development branch like so:

```
$ git clone https://github.com/ivco19/libs.git
```

This will create a directory libs in your current directory.

5.2 Command line interface (CLI)

After install arcovid19 you also have a command line app to download the data as csv.

```
$ arcovid19 --help
Usage: arcovid19 [OPTIONS]

Retrieve and store the database as an as CSV file.

Options:
--url=STR      str The url for the excel table to parse. Default is ivco19 team table.
↳ (default: https://github.com/ivco19/libs/raw/master/databases/cases.xlsx)
--out=STR      PATH (default=stdout) The output path to the CSV file. If it's not
↳ provided the data is printed in the stdout.
--nocached     If you want to ignore the local cache or retrieve a new value.

Other actions:
-h, --help    Show the help
```

5.3 Cases Tutorial

This section contains a step-by-step tutorial with examples for using the **arcovid19** tools.

In part 1 of the tutorial, we explained loading data using arcovid19 and some of the methods and attributes inherited from the DataFrame class. Part 2 shows the implementation of the *Cases Frame* class methods on the data basis. Lastly, in part 3 of this tutorial, we share the visualization of the COVID-19 database using the tools provided by the *Plot Cases* class from arcovid19.

5.3.1 Part 1 - Load cases

The first step is to import **arcovid19**, then we load the COVID-19 database from Argentina (**CASES_URL**). As a result, a table is obtained that presents a multiple pandas index, with the following hierarchy:

level 0: cod_provincia - Argentina states

level 1: cod_status - Four states of disease patients (**R** = Recovered, **C** = Confirmed, **A** = Active, **D** = Dead)

```
[1]: import arcovid19 as ac
```

```
[2]: cases = ac.load_cases()
```

```
[3]: cases.head(4)
```

```
[3]:
```

cod_provincia	cod_status	provincia_status	Pcia_status	\
CABA	C	CABA_C	CABA Casos Confirmados	
	R	CABA_R	CABA Casos Recuperados	
	D	CABA_D	CABA Casos Muertos	
	A	CABA_A	CABA Casos Activos	
		2020-03-03 00:00:00	2020-03-04 00:00:00	\
cod_provincia	cod_status			
CABA	C	1.0	1.0	
	R	0.0	0.0	

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	D	0.0	0.0	
	A	1.0	1.0	
		2020-03-05 00:00:00	2020-03-06 00:00:00	\
cod_provincia	cod_status			
CABA	C	1.0	5.0	
	R	0.0	0.0	
	D	0.0	0.0	
	A	1.0	5.0	
		2020-03-07 00:00:00	2020-03-08 00:00:00	\
cod_provincia	cod_status			
CABA	C	6.0	8.0	
	R	0.0	0.0	
	D	1.0	1.0	
	A	5.0	7.0	
		2020-03-09 00:00:00	2020-03-10 00:00:00	... \
cod_provincia	cod_status			...
CABA	C	9.0	10.0	...
	R	0.0	0.0	...
	D	1.0	1.0	...
	A	8.0	9.0	...
		2020-05-19 00:00:00	2020-05-20 00:00:00	\
cod_provincia	cod_status			
CABA	C	3566.0	3823.0	
	R	1139.0	1139.0	
	D	135.0	137.0	
	A	2292.0	2547.0	
		2020-05-21 00:00:00	2020-05-22 00:00:00	\
cod_provincia	cod_status			
CABA	C	4202.0	4606.0	
	R	1139.0	1139.0	
	D	139.0	146.0	
	A	2924.0	3321.0	
		2020-05-23 00:00:00	2020-05-24 00:00:00	\
cod_provincia	cod_status			
CABA	C	5006.0	5500.0	
	R	1139.0	1678.0	
	D	152.0	155.0	
	A	3715.0	3667.0	
		2020-05-25 00:00:00	2020-05-26 00:00:00	\
cod_provincia	cod_status			
CABA	C	5875.0	6202.0	
	R	1678.0	1678.0	
	D	163.0	168.0	
	A	4034.0	4356.0	
		2020-05-27 00:00:00	2020-05-28 00:00:00	
cod_provincia	cod_status			
CABA	C	6564.0	6989.0	
	R	1678.0	1963.0	
	D	172.0	175.0	

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```
A                4714.0                4851.0

[4 rows x 89 columns]
```

```
[4]: cases.shape
```

```
[4]: (101, 89)
```

```
[5]: cases.describe()
```

```
[5]:
```

	2020-03-03	2020-03-04	2020-03-05	2020-03-06	2020-03-07	2020-03-08	\
count	100.000000	101.000000	101.000000	101.000000	101.000000	101.000000	
mean	0.040000	0.039604	0.089109	0.346535	0.357673	0.478548	
std	0.196946	0.196000	0.349257	1.359674	1.432113	1.941601	
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	
25%	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	
50%	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	
75%	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	
max	1.000000	1.000000	2.000000	8.000000	9.000000	12.000000	

	2020-03-09	2020-03-10	2020-03-11	2020-03-12	...	2020-05-19	\
count	101.000000	101.000000	101.000000	101.000000	...	101.000000	
mean	0.677393	0.753640	0.832725	1.232438	...	348.871805	
std	2.603514	2.926856	3.252652	4.689155	...	1176.302303	
min	0.000000	0.000000	0.000000	0.000000	...	0.000000	
25%	0.000000	0.000000	0.000000	0.000000	...	2.000000	
50%	0.000000	0.000000	0.000000	0.000000	...	14.000000	
75%	0.000000	0.000000	0.000000	0.000000	...	89.000000	
max	17.000000	19.000000	21.000000	31.000000	...	8809.000000	

	2020-05-20	2020-05-21	2020-05-22	2020-05-23	2020-05-24	\
count	101.000000	101.000000	101.000000	101.000000	101.000000	
mean	367.644097	393.307622	421.743290	449.624417	478.258056	
std	1248.414401	1348.455749	1460.412128	1572.673190	1664.493107	
min	0.000000	0.000000	0.000000	0.000000	0.000000	
25%	2.000000	2.000000	2.000000	2.000000	2.000000	
50%	14.000000	15.000000	16.000000	16.000000	16.000000	
75%	89.000000	90.000000	90.000000	90.000000	90.000000	
max	9283.000000	9931.000000	10649.000000	11353.000000	12076.000000	

	2020-05-25	2020-05-26	2020-05-27	2020-05-28
count	101.000000	101.000000	101.000000	101.000000
mean	500.119264	523.881659	551.802508	582.257972
std	1752.146024	1823.126460	1933.166375	2034.023400
min	0.000000	0.000000	0.000000	0.000000
25%	2.000000	2.000000	3.000000	3.000000
50%	17.000000	17.000000	17.000000	14.000000
75%	90.000000	90.000000	89.000000	110.000000
max	12628.000000	13228.000000	13933.000000	14702.000000

```
[8 rows x 87 columns]
```

Remember that the *cases* object has the properties of a DataFrame.

5.3.2 Part 2 - Functions

After loading the database we can use some of the functions of the class object *Cases Frame*.

```
[1]: import arcovid19 as ac
```

```
[2]: cases = ac.load_cases()
```

Using the method *last_growth_rate* and specifying the *provincia* argument of the function, we obtain the latest growth rate available for the whole country or by province.

If *provincia = None*, the method will yield the value of Argentina's last growth rate.

```
[3]: cases.last_growth_rate(provincia=None)
```

```
[3]: 0.05519270795952047
```

If you want to get the latest growth rate for any particular province, you must replace *None* for the full name of the province.

```
[4]: cases.last_growth_rate(provincia='córdoba')
```

```
/home/docs/checkouts/readthedocs.org/user_builds/arcovid19/envs/latest/lib/python3.7/
↪site-packages/arcovid19-0.6b0-py3.7.egg/arcovid19/cases.py:731: RuntimeWarning:
↪divide by zero encountered in true_divide
  growth_rate = np.array((I_n / I_n_1) - 1)
/home/docs/checkouts/readthedocs.org/user_builds/arcovid19/envs/latest/lib/python3.7/
↪site-packages/arcovid19-0.6b0-py3.7.egg/arcovid19/cases.py:731: RuntimeWarning:
↪invalid value encountered in true_divide
  growth_rate = np.array((I_n / I_n_1) - 1)
```

```
[4]: -0.002178649237472796
```

In the case that you want to estimate the growth rate in the period where we have data by province or country, you can use the *grateful_full_period* method.

Again, if the province argument equals *None*, the series shown corresponds to Argentina.

```
[5]: cases.grate_full_period(provincia=None).head(5)
```

```
[5]: 2020-03-04      0
     2020-03-05      1
     2020-03-06      3
     2020-03-07    0.125
     2020-03-08    0.333333
     Name: (ARG, growth_rate_C), dtype: object
```

```
[6]: cases.grate_full_period(provincia='córdoba').head(5)
```

```
[6]: 2020-03-04    NaN
     2020-03-05    NaN
     2020-03-06    NaN
     2020-03-07    0.0
     2020-03-08    0.0
     dtype: float64
```

In the case that you want the observations by date, that is, not cumulative, you can use the *restore_time_serie* method as shown in the next cell.

```
[7]: cases.restore_time_serie().head(4)
```

```
[7]:
```

		provincia_status	Pcia_status	\
cod_provincia	cod_status			
CABA	C	CABA_C	CABA Casos Confirmados	
	R	CABA_R	CABA Casos Recuperados	
	D	CABA_D	CABA Casos Muertos	
	A	CABA_A	CABA Casos Activos	
		2020-03-03 00:00:00	2020-03-04 00:00:00	\
cod_provincia	cod_status			
CABA	C	1.0	0.0	
	R	0.0	0.0	
	D	0.0	0.0	
	A	1.0	0.0	
		2020-03-05 00:00:00	2020-03-06 00:00:00	\
cod_provincia	cod_status			
CABA	C	0.0	4.0	
	R	0.0	0.0	
	D	0.0	0.0	
	A	0.0	4.0	
		2020-03-07 00:00:00	2020-03-08 00:00:00	\
cod_provincia	cod_status			
CABA	C	1.0	2.0	
	R	0.0	0.0	
	D	1.0	0.0	
	A	0.0	2.0	
		2020-03-09 00:00:00	2020-03-10 00:00:00	... \
cod_provincia	cod_status			...
CABA	C	1.0	1.0	...
	R	0.0	0.0	...
	D	0.0	0.0	...
	A	1.0	1.0	...
		2020-05-19 00:00:00	2020-05-20 00:00:00	\
cod_provincia	cod_status			
CABA	C	224.0	257.0	
	R	262.0	0.0	
	D	5.0	2.0	
	A	-43.0	255.0	
		2020-05-21 00:00:00	2020-05-22 00:00:00	\
cod_provincia	cod_status			
CABA	C	379.0	404.0	
	R	0.0	0.0	
	D	2.0	7.0	
	A	377.0	397.0	
		2020-05-23 00:00:00	2020-05-24 00:00:00	\
cod_provincia	cod_status			
CABA	C	400.0	494.0	
	R	0.0	539.0	
	D	6.0	3.0	
	A	394.0	-48.0	

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```

                2020-05-25 00:00:00  2020-05-26 00:00:00  \
cod_provincia cod_status
CABA          C          375.0          327.0
              R           0.0           0.0
              D           8.0           5.0
              A          367.0          322.0

                2020-05-27 00:00:00  2020-05-28 00:00:00
cod_provincia cod_status
CABA          C          362.0          425.0
              R           0.0          285.0
              D           4.0           3.0
              A          358.0          137.0

[4 rows x 89 columns]
```

5.3.3 Part 3 - Visualization of cases

In this part of the tutorial you can find some of the functions of the * Cases Plot * class, in particular this class inherits the properties of the Matplotlib class.

As usual the first step will be to import the arcovid19 module.

```
[1]: import arcovid19 as ac
import matplotlib.pyplot as plt
```

The next cell performs a cache cleanup, as arcovid19 saves a one-hour cache for every request made to the online database. With this action, we make sure that we are using an updated database.

```
[2]: ac.CACHE.clear()
```

```
[2]: 2
```

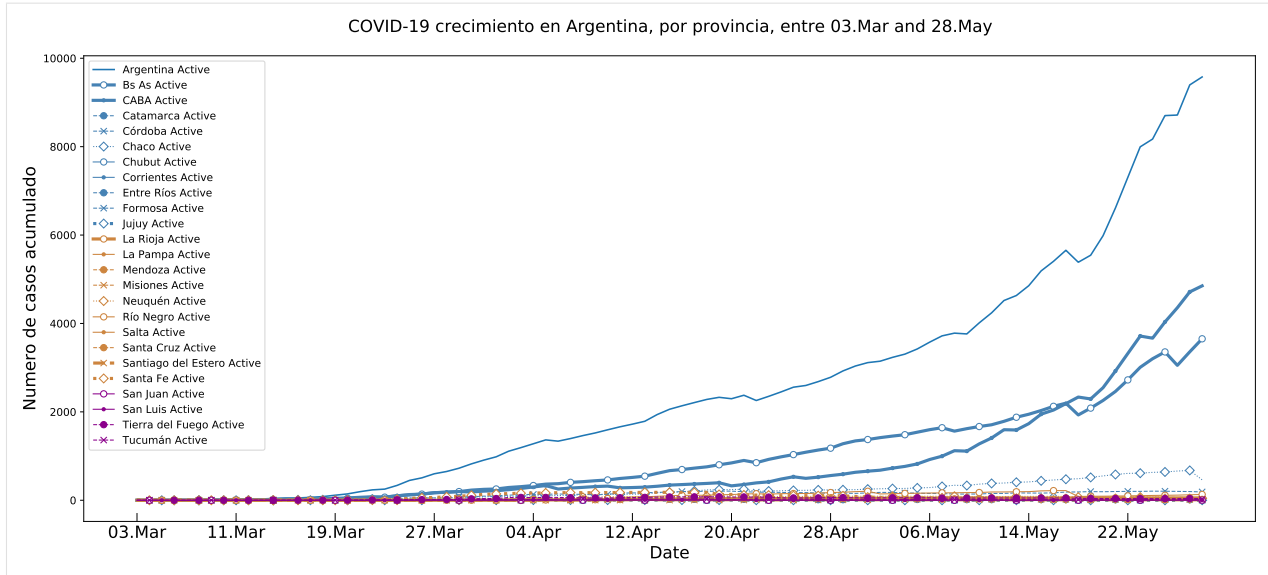
We loaded the COVID-19 database from Argentina.

```
[3]: cases = ac.load_cases()
```

The *plot* method shows the trend to date of each of the provinces along with that of Argentina. By default, it shows confirmed cases, but you can also choose to view active, recovered and dead cases.

```
[4]: fig, ax = plt.subplots(figsize=(20, 8))
cases.plot(ax=ax, confirmed=False, active=True, recovered=False, deceased=False)
ax.legend(loc=2)
```

```
[4]: <matplotlib.legend.Legend at 0x7f611f3a0080>
```

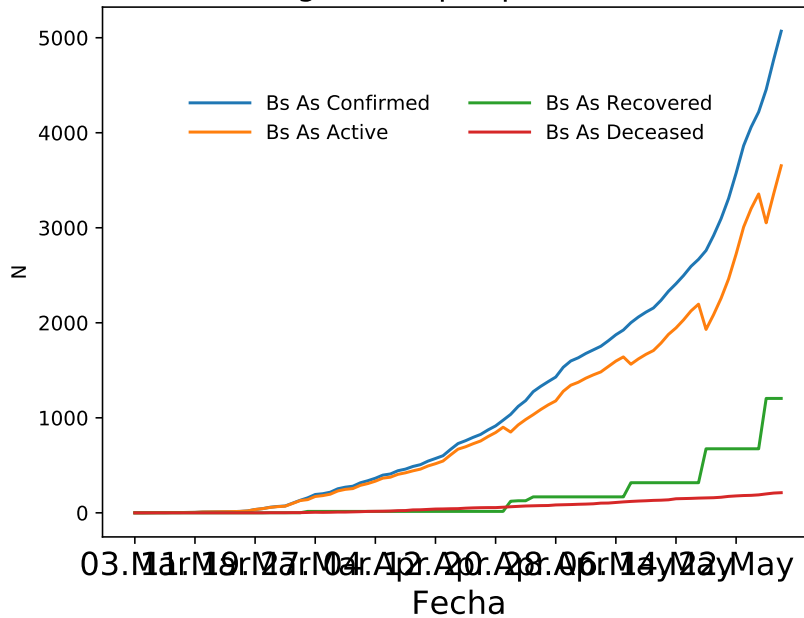


Then if you are interested in a particular province, you just have to add `grate_full_periode` and specify the name of the province. If you do not enter the name of any province the tool will show you the cases of Argentina.

```
[5]: ax = cases.plot.grate_full_periode('Bs As', confirmed=True, active=True,
    ↪ recovered=True, deceased=True)

/home/docs/checkouts/readthedocs.org/user_builds/arcovid19/envs/latest/lib/python3.7/
    ↪ site-packages/ipykernel_launcher.py:1: DeprecationWarning: Call to deprecated
    ↪ method grate_full_periode. (use growth_provincia instead) -- Deprecated since
    ↪ version 0.5.
    """Entry point for launching an IPython kernel.
```

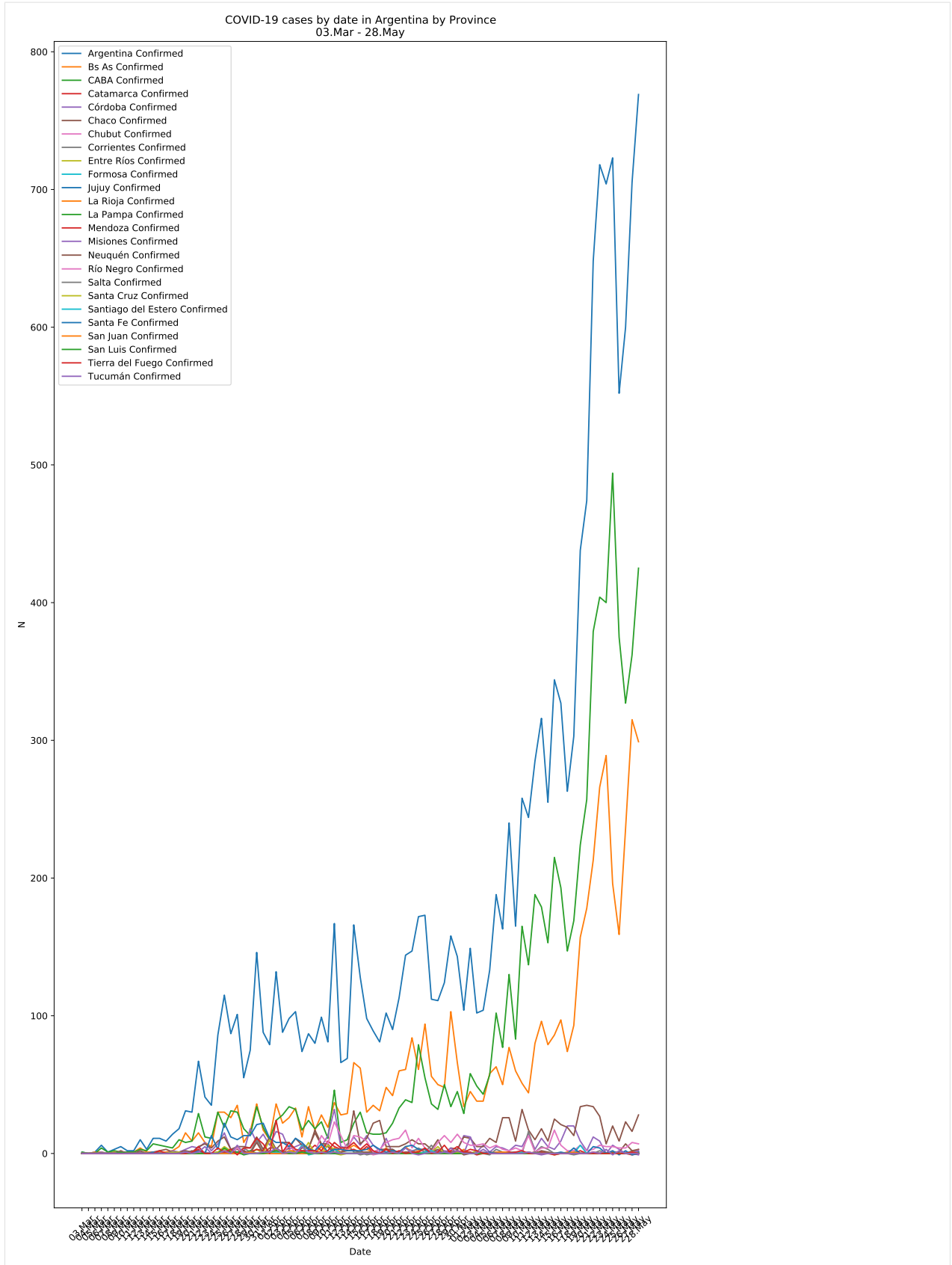
COVID-19 crecimiento en Argentina, por provincia, entre 03.Mar and 28.May



Also, if you want to observe the behavior of observations by date, you can use the following graphical representations.

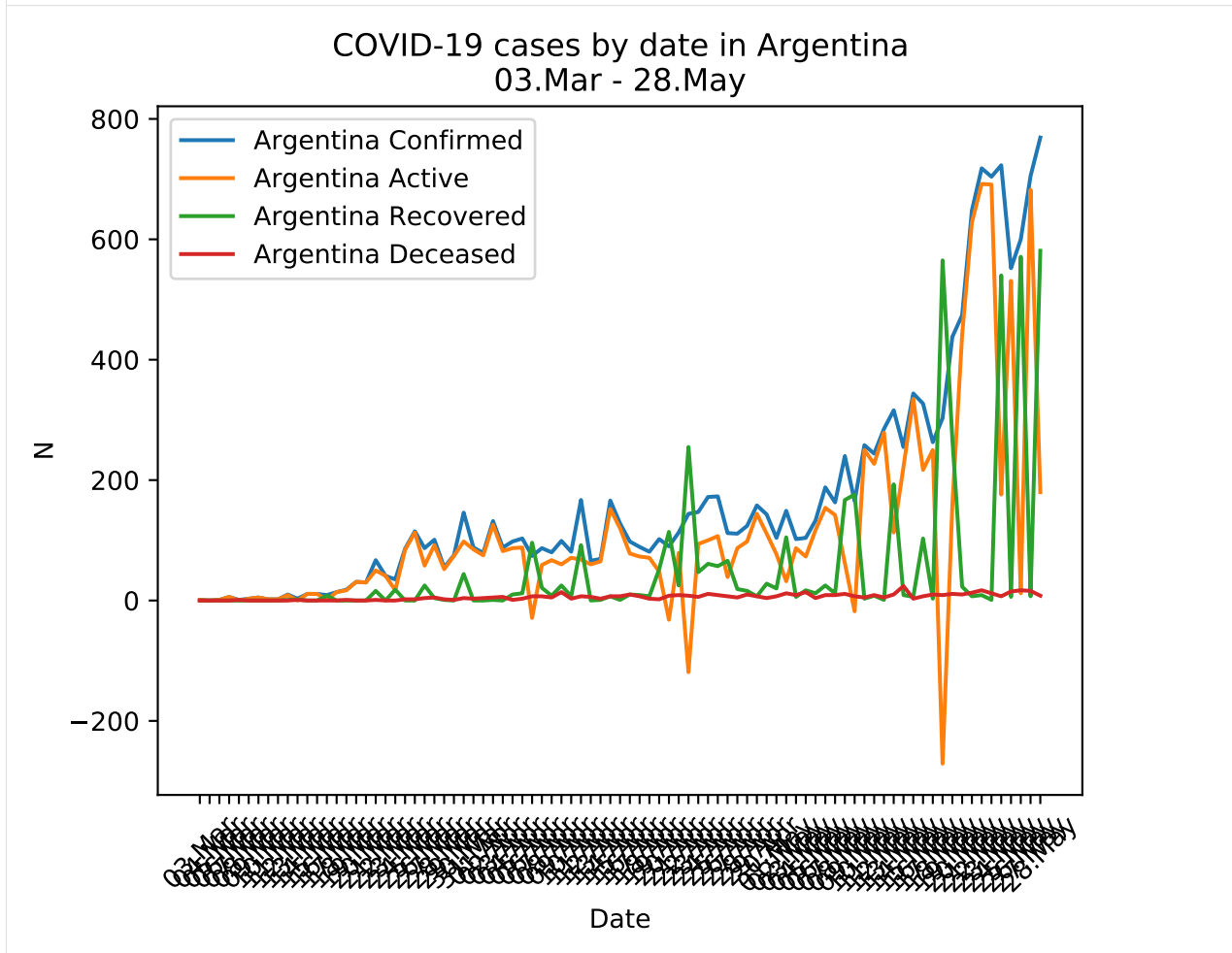
```
[6]: cases.plot.time_series_all()
```

```
[6]: <matplotlib.axes._subplots.AxesSubplot at 0x7f611f5475c0>
```



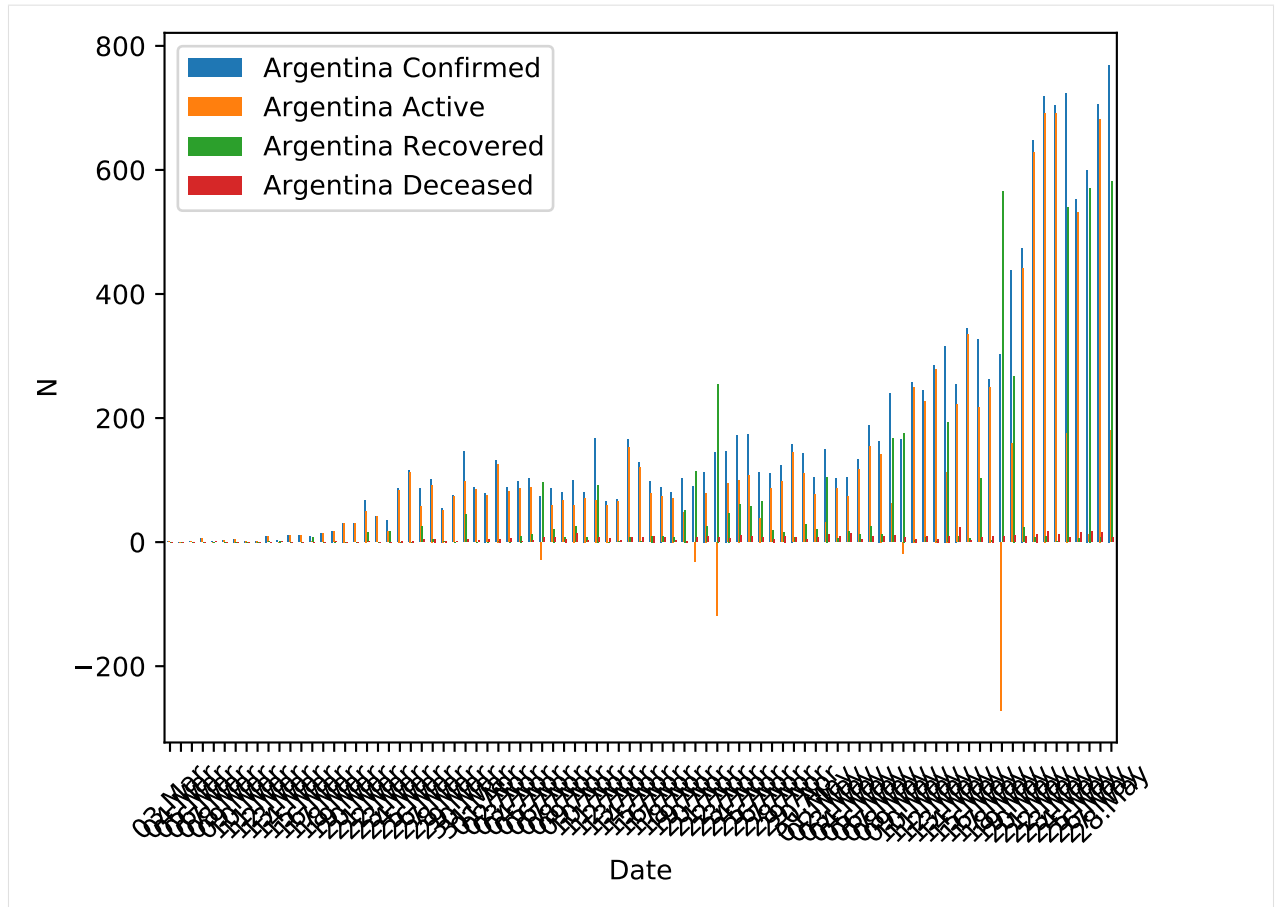

```
[7]: cases.plot.time_serie()
```

```
[7]: <matplotlib.axes._subplots.AxesSubplot at 0x7f611f725208>
```



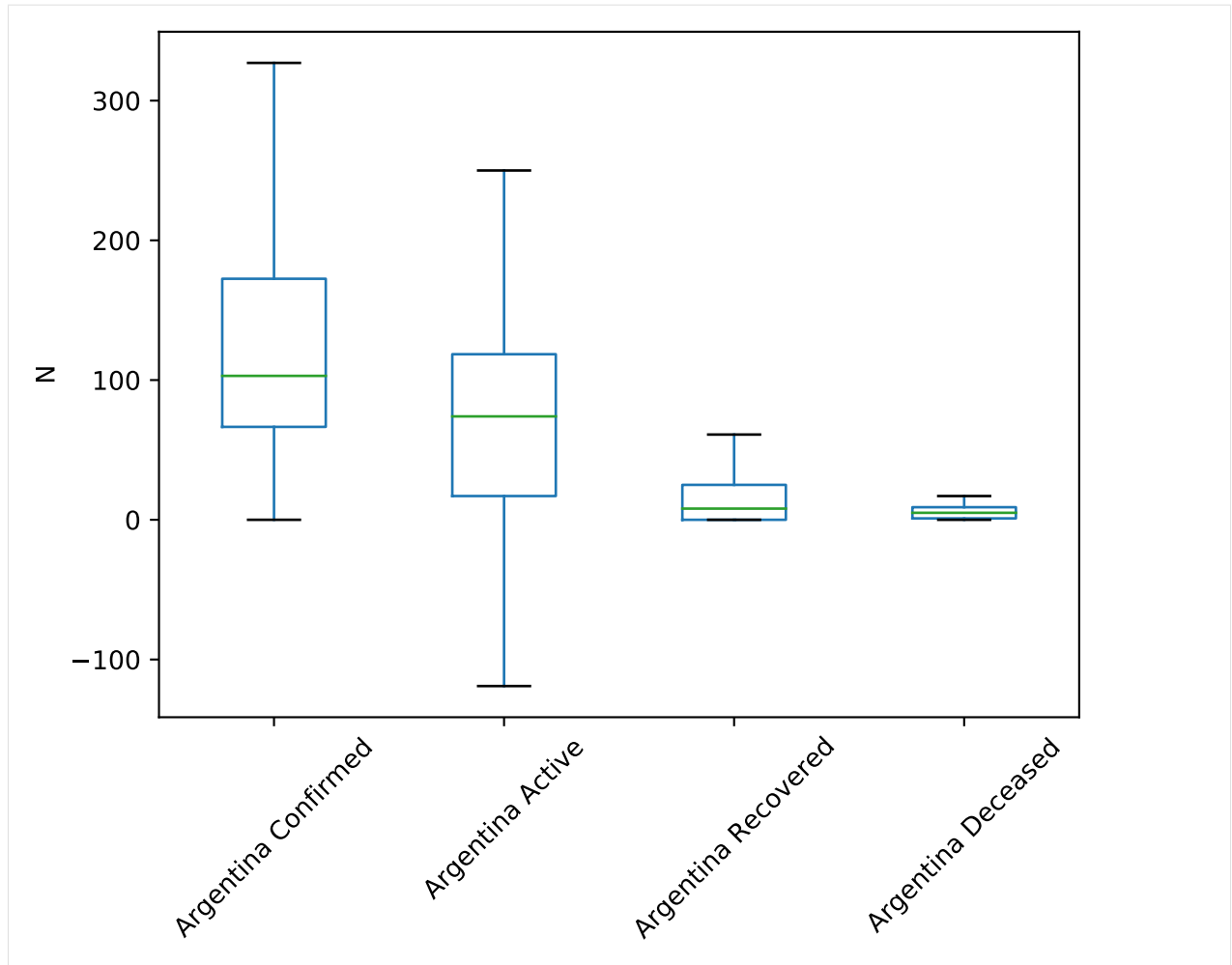
```
[8]: cases.plot.barplot()
```

```
[8]: <matplotlib.axes._subplots.AxesSubplot at 0x7f611d0152e8>
```



```
[9]: cases.plot.boxplot(showfliers=False)
```

```
[9]: <matplotlib.axes._subplots.AxesSubplot at 0x7f611cbdeb8>
```



Finally, at the following link [part 3](#) you will find an updated guide on March 28, 2020, of part 3 of the tutorial.

5.4 Web Tutorial

Arcovid19 presents a tool for the online visualization of epidemiological models.

The simplest way to execute this application is simply to run

Warning: This is a test server, and should not be used in a production environment for any reason.

```
$ arcovid19 webserver
* Serving Flask app "arcovid19.web" (lazy loading)
* Environment: production
WARNING: This is a development server. Do not use it in a production deployment.
Use a production WSGI server instead.
* Debug mode: on
* Running on http://127.0.0.1:5000/ (Press CTRL+C to quit)
* Restarting with inotify reloader
```

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```
* Debugger is active!
* Debugger PIN: XXX
```

This launches a **local** application which you can access through the url `http://localhost:5000`.

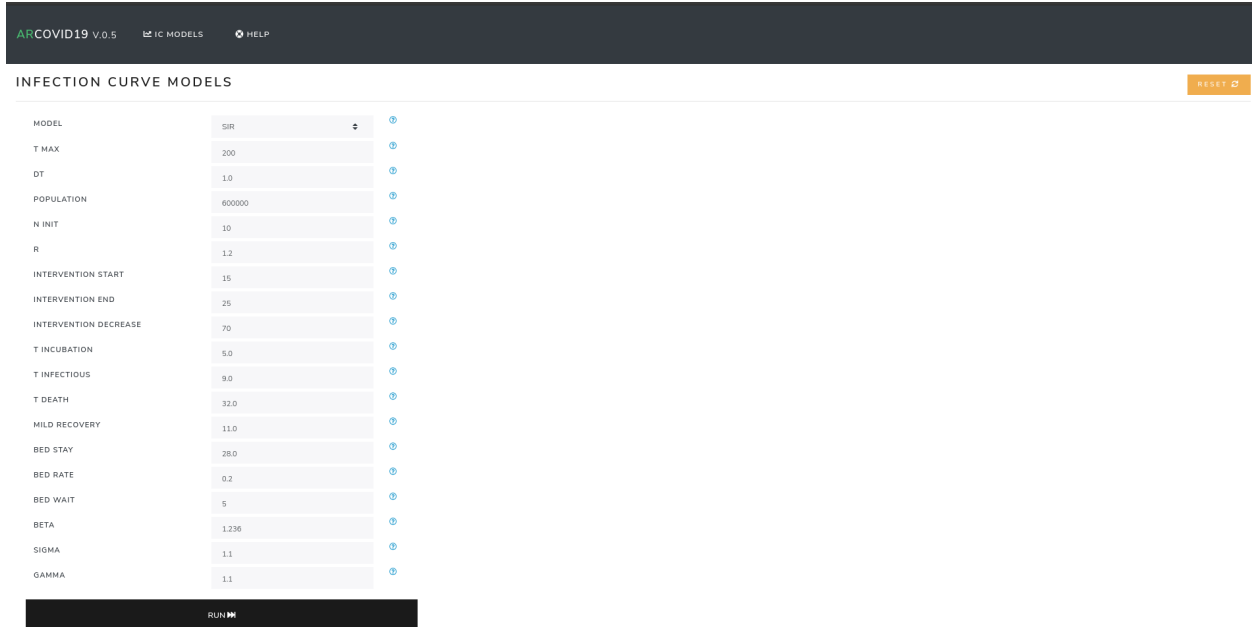


Fig. 1: Default view of arcovid19 webclient in version 0.5.

If for some reason it is necessary to launch the app in another *IP* or *port*, this can be specified with the options `--host` and `--port` respectively. For example, if you want to serve the local network on port `8000` the command would be

```
$ arcovid19 webserver --host 0.0.0.0 --port 8000
* Serving Flask app "arcovid19.web" (lazy loading)
* Environment: production
WARNING: This is a development server. Do not use it in a production deployment.
Use a production WSGI server instead.
* Debug mode: on
* Running on http://0.0.0.0:8000/ (Press CTRL+C to quit)
* Restarting with inotify reloader
* Debugger is active!
* Debugger PIN: 242-079-243
```

This would allow anyone connected to the same local network as the computer where the web application is started to access the website via the server's IP and port `8000`.

Note: For more web server options you can execute the command `arcovid19 webserver --help`.

5.4.1 Cambiando de idioma

So far arcovid19 web only has two languages implemented.

1. en - Ingles (Activated by default)

2. es - Español.

To activate the alternative language, you must assign a `environment variable`. called `ARCOVID19_DEFAULT_LOCALE`.

This is done with the command

```
$ export ARCOVID19_DEFAULT_LOCALE=es;
```

After that, it is simply a matter of launching the application with `arcovid19 webserver`.

5.4.2 Deployment

To run `arcovid19 webserver` in a production environment at least 2 environment variables must be configured:

- `ARCOVID19_DEBUG=false`
- `ARCOVID19_SECRET_KEY=a-lot-of-random-chars`

`ARCOVID19_SECRET_KEY` has to be a contiguous random string of random values, you can, for example, press keys without any sense.

If you want to configure the language to Spanish, you must export the environment variable

- `ARCOVID19_DEFAULT_LOCALE=es`

Deployment can be done with any available method to `Flask`, and the wsgi application is available with the function `arcovid19.web.create_app()`.

For example, to launch the application with `Gunicorn` can be done with the command

```
$ gunicorn 'arcovid19.web:create_app()'
```

In addition, the repository is already configured with the files `requirements.txt` and `Procfile` to work in `Heroku`.

An online version of the webapp is available in [here](#).

5.5 API

5.5.1 arcovid19 package

Module contents

Utilities to access different Argentina-Related databases of COVID-19 data from the Arcovid19 group.

Subpackages

`arcovid19.web` package

Submodules

`arcovid19.web.bp` module

Core functionalities for `arcovid19`

arcovid19.web.bp.wavid19 = <flask.blueprints.Blueprint object>
Flask blueprint with the views implemented in arcovid19.

arcovid19.web.forms module

WTF Forms for arcovid19

```
class arcovid19.web.forms.InfectionCurveForm (formdata=<object object>, **kwargs)  
    Bases: flask_wtf.form.FlaskForm  
  
    N_init = <UnboundField(IntegerField, (1'N Init',)), {'description': 1'Number of initial  
    R = <UnboundField(FloatField, (1'R',)), {'description': 1'Reproduction number.', 'default  
    bed_rate = <UnboundField(FloatField, (1'Bed Rate',)), {'description': 1'Hospitalization  
    bed_stay = <UnboundField(FloatField, (1'Bed Stay',)), {'description': 1'Length of hospi  
    bed_wait = <UnboundField(IntegerField, (1'Bed Wait',)), {'description': 1'Time from fir  
    beta = <UnboundField(FloatField, (1'Beta',)), {'description': 1'SEIR Model beta ($\beta  
    dt = <UnboundField(FloatField, (1'Dt',)), {'description': 1'Time step [days].', 'default  
    gamma = <UnboundField(FloatField, (1'Gamma',)), {'description': 1'SEIR Model gamma ($\gamma  
    intervention_decrease = <UnboundField(FloatField, (1'Intervention Decrease',)), {'descr  
    intervention_end = <UnboundField(IntegerField, (1'Intervention End',)), {'description':  
    intervention_start = <UnboundField(IntegerField, (1'Intervention Start',)), {'descripti  
    mild_recovery = <UnboundField(FloatField, (1'Mild Recovery',)), {'description': 1'Recov  
    model = <UnboundField(SelectField, (1'Model',)), {'choices': [(1'do_SIR', 1'SIR'), (1'do_  
    population = <UnboundField(IntegerField, (1'Population',)), {'description': 1'Populatio  
    sigma = <UnboundField(FloatField, (1'Sigma',)), {'description': 1'SEIR Model sigma ($\sigma  
    t_death = <UnboundField(FloatField, (1'T Death',)), {'description': 1'Time from end of  
    t_incubation = <UnboundField(FloatField, (1'T Incubation',)), {'description': 1'Length  
    t_infectious = <UnboundField(FloatField, (1'T Infectious',)), {'description': 1'None',  
    t_max = <UnboundField(IntegerField, (1'T Max',)), {'description': 1'Time range [days].'
```

Submodules

arcovid19.core module

Core functionalities for arcovid19

```
class arcovid19.core.Frame (df, extra=NOTHING)  
    Bases: object  
  
    plot_cls  
  
class arcovid19.core.Plotter (frame)  
    Bases: object  
  
    default_plot_name_method
```

arcovid19.cache module

Cache abstractions for arcovid19

```
arcovid19.cache.DEFAULT_CACHE_DIR = '/home/docs/arcovid19_data/_cache_'
    Default cache location, (default=~/.arcovid_19_data/_cache_)
```

```
arcovid19.cache.CACHE = <diskcache.core.Cache object>
    Default cache instance
```

```
arcovid19.cache.CACHE_EXPIRE = 3600
    Time to expire of every load_cases call in seconds
```

```
arcovid19.cache.from_cache (tag, function, force=False, *args, **kwargs)
    Simple cache orchestration.
```

Parameters

- **tag** (*str*) – Normally every function call the cache with their own tag. We suggest “module.function” or “module.Class.function”
- **function** (*callable*) – The function to be cached
- **force** (*bool* (default=False)) – If the vale of the cache must be ignored and re-execute the function.
- **and kwargs** (*args*) – All the parameters needed to execute the function.

arcovid19.cases module

Utilities to Utility function to parse all the actual cases of the COVID-19 in Argentina.

```
arcovid19.cases.CODE_TO_POVINCIA = {'BA': 'Bs As', 'CABA': 'CABA', 'CAT': 'Catamarca', 'CBZ'}
    List of Argentina provinces
```

```
arcovid19.cases.D0 = datetime.datetime(2020, 3, 11, 0, 0)
    Pandemia Start 2020-03-11
```

```
arcovid19.cases.Q1 = datetime.datetime(2020, 3, 20, 0, 0)
    Argentine quarantine starts 2020-03-20
```

```
class arcovid19.cases.CasesPlot (frame)
    Bases: arcovid19.core.Plotter
```

```
barplot (provincia=None, confirmed=True, active=True, recovered=True, deceased=True, ax=None,
        **kwargs)
```

```
boxplot (provincia=None, confirmed=True, active=True, recovered=True, deceased=True, ax=None,
        **kwargs)
```

```
curva_epi_pais (ax=None, argentina=True, exclude=None, log=False, norm=False, paint=None,
        count_days=None, **kwargs)
    method: full_period_normalized()
```

This function plots the time series, similar to grate_full_period_all, but including a second axis and comments about the start of quarantine

opciones para paint: pandemia, cuarentena

opciones para count_days: pandemia, cuarentena

```
curva_epi_provincia (*args, **kwargs)
```

```
default_plot_name_method = 'curva_epi_pais'
```

grate_full_period (*args, **kwargs)

grate_full_period_all (*args, **kwargs)

growth_provincia (provincia=None, confirmed=True, active=True, recovered=True, deceased=True, ax=None, log=False, norm=False, **kwargs)

time_serie (provincia=None, confirmed=True, active=True, recovered=True, deceased=True, ax=None, **kwargs)

time_serie_all (ax=None, argentina=True, exclude=None, **kwargs)

class arcovid19.cases.CasesFrame (df, extra=NOTHING)

Bases: *arcovid19.core.Frame*

Wrapper around the *load_cases()* table.

This class adds functionalities around the dataframe.

dates

Returns the dates for which we have data.

Useful to use as time column (row) list for wide (long) format.

get_provincia_name_code (provincia)

Resolve and validate the name and code of a given provincia name or code.

grate_full_period (provincia=None)

Estimates growth rate for the period where we have data

last_growth_rate (provincia=None)

Returns the last available growth rate for the whole country if provincia is None, or for only the named region.

plot_cls

alias of *CasesPlot*

restore_time_serie ()

Retrieve a new pandas.DataFrame but with observations by Date.

tot_cases

Returns latest value of total confirmed cases

arcovid19.cases.load_cases (cases_url='https://github.com/ivco19/libs/raw/master/databases/cases.xlsx', areas_pop_url='https://github.com/ivco19/libs/raw/master/databases/extra/arg_provs.dat', force=False)

Utility function to parse all the actual cases of the COVID-19 in Argentina.

Parameters

- **cases_url** (*str*) – The url for the excel table to parse. Default is ivco19 team table.
- **areas_pop_url** (*str*) – The url for the csv population table to parse. Default is ivco19 team table.
- **force** (*bool* (default=False)) – If you want to ignore the local cache and retrieve a new value.

Returns

CasesFrame – It features a pandas multi index, with the following hierarchy:

- level 0: cod_provincia - Argentina states
- level 1: cod_status - Four states of disease patients (R, C, A, D)

Return type Pandas-DataFrame like object with all the arcovid19 datatabase.

arcovid19.models module

Utilities to Utility function to parse all the actual cases of the COVID-19 in Argentina.

exception `arcovid19.models.NodeNotFoundError`

Bases: `KeyError`

If a node is not found inside a graph

class `arcovid19.models.Node` (*id, value, outgoing=NOTHING, incoming=NOTHING*)

Bases: `object`

This class is used to create and manipulated nodes.

add_neighbor (*neighbor, names, values*)

be_neighbor (*neighbor, names, values*)

get_connections ()

get_weight (*neighbor*)

class `arcovid19.models.Graph` (*vert_dict=NOTHING*)

Bases: `object`

This class is used to create and manipulated graphs.

It makes a heavy use of the node class A graph is made of nodes and edges. This class allows to store a value for each node and different “weights” for each edge. Also, edges are directed.

Example

```
>>> g = Graph()
>>> for i, inode in enumerate(['A', 'B', 'C', 'D']):
...     print(i)
...     g.add_node(inode, 0)
```

```
>>> nms = ['x', 'y']
>>> g.add_edge('A', 'B', nms, [1, 100])
>>> g.add_edge('A', 'C', nms, [2, 200])
>>> g.add_edge('B', 'D', nms, [3, 300])
>>> g.add_edge('D', 'B', nms, [4, 400])
>>> g.add_edge('D', 'C', nms, [5, 500])
>>> g.add_edge('C', 'C', nms, [6, 600])
```

A node can be connected to itself. >>> `g.add_edge('B', 'B', nms, [333, 333])` >>> `g.show()` ...

vert_dict

a dict containing the vertices

Type dict

add_edge (*frm, to, names=None, values=0*)

Link two nodes inside a graph.

Notes

Does not verify if edge already exists

Raises `NodeNotFound`: – If one of the node are not registered in the graph

add_node (*nnode, value*)

Adds a new node to a graph.

The node must have a value.

format_connections ()

format_weights ()

get_edge (*frm, to, field*)

get_node (*n*)

Retrieve a node if exists.

Parameters *n* (*str*) –

Returns *node*

Return type a node object

get_node_ids ()

get_node_value (*n*)

Returns the value of the node.

Parameters *n* (*str*) –

Returns *value*

Return type float

get_nodes_from (*nnode*)

get_nodes_to (*nnode*)

node_activation (*nnode, key*)

node_upgrade (*nnode, key*)

resume_connections ()

resume_weights ()

set_node (*n, value*)

Updates the Node value. The node must exists inside the graph.

Parameters

- **n** (*str*) – The ID or name of the node
- **value** (*float*) – The value to be assigned to the node

class arcovid19.models.**InfectionCurve** (*population: int = 600000, N_init: int = 10, R: float = 1.2, intervention_start: int = 15, intervention_end: int = 25, intervention_decrease: float = 70, t_incubation: float = 5.0, t_infectious: float = 9.0, t_death: float = 32.0, mild_recovery: float = 11.0, bed_stay: float = 28.0, bed_rate: float = 0.2, bed_wait: int = 5, beta: float = 1.236, sigma: float = 1.1, gamma: float = 1.1*)

Bases: object

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Parameters

- **population** (*int* (*default=600000*)) – Population.
- **N_init** (*int* (*default=10*)) – Number of initial infections.

- **R** (*float* (*default=1.2*)) – Reproduction number.
- **intervention_start** (*int* (*default=15*)) – Start intervention days.
- **intervention_end** (*int* (*default=25*)) – End intervention days.
- **intervention_decrease** (*float* (*default=70*)) – Decrease in transmission for intervention, percentage (0-100) 100 means total isolation.
- **t_incubation** (*float* (*default=5.*)) – Length of incubation period.
- **t_infectiou** (= (*default=9.*)) – Duration patient is infectious.
- **t_death** (*float* (*default=32.*)) – Time from end of incubation to death.
- **mild_recovery** (*float* (*default=0.2*)) – Recovery time for mild (not severnot severe) cases, days hospitalization rate (fraction).
- **bed_stay** (*float* (*default=28.*)) – Length of hospital stay, days.
- **bed_rate** (*float* (*default=0.2*)) – Hospitalization rate (fraction).
- **bed_wait** (*int* (*default=5*)) – Time from first synthoms to hospitalization (days).
- **beta** (*float* (*default=1.236*)) – SEIR Model beta (β).
- **sigma** (*float* (*default=1.1*)) – SEIR Model sigma (σ).
- **gamma** (*float* (*default=1.1*)) – SEIR Model gamma (γ).

References

“Stochastic SIR model with Python,” *epirecipes*. [Online]. Available: <https://tinyurl.com/y8zwvfk4>. [Accessed: 09-May-2020].

do_SEIR (*t_max=200, dt=1.0*)

This function implements a SEIR model without vital dynamics under the assumption of a closed population. Recovered individuals become immune for ever. ref.: <https://www.idmod.org/docs/hiv/model-seir.html>

Parameters

- **t_max** (*int* (*default=200*)) – Time range [days].
- **dt** (*float* (*default=1.*)) – Time step [days].

Returns value

Return type Time series for S, E, I and R

do_SEIRF (*t_max=200, dt=1.0*)

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Parameters

- **t_max** (*int* (*default=200*)) – Time range [days].
- **dt** (*float* (*default=1.*)) – Time step [days].

Returns value

Return type Time series for S, E, I, R and F

do_SIR (*t_max=200, dt=1.0*)

This function implements a SIR model without vital dynamics under the assumption of a closed population.

Recovered individuals become immune for ever. In this model exposed individuals become instantly infectious, i.e., there is no latency period like in the SEIR model.

Parameters

- **t_max** (*int (default=200)*) – Time range [days].
- **dt** (*float (default=1.)*) – Time step [days].

Returns value

Return type Time series for S, I and R

`arcovid19.models.load_infection_curve (**kwargs)`

Return a new instance of infection curve.

arcovid19.cli module

Command line interfaces to access different Argentina-Related databases of COVID-19 data from the Arcovid19 group.

`arcovid19.cli.main()`

Run the arcovid19 command line interface.

`arcovid19.cli.cases (*, url='https://github.com/ivco19/libs/raw/master/databases/cases.xlsx', force=False, out=None)`

Retrieve and store the cases database in CSV format.

url: str The url for the excel table to parse. Default is ivco19 team table.

out: PATH (default=stdout) The output path to the CSV file. If it's not provided the data is printed in the stdout.

force: If you want to ignore the local cache or retrieve a new value.

`arcovid19.cli.webserver (*, host=None, port=None, nodebug=False, reload=False, load_dotenv=True)`

Run a development server for arcovid19 utilities.

host: str the hostname to listen on. Set this to '0.0.0.0' to have the server available externally as well. Defaults to '127.0.0.1' or the host in the SERVER_NAME config variable if present.

port: int the port of the webserver. Defaults to 5000 or the port defined in the SERVER_NAME config variable if present.

nodebug: bool if given, disable debug mode.

reload: bool If its True any change of the code will restart the webserver.

load_dotenv: Load the nearest '.env' and '.flaskenv' files to set environment variables. Will also change the working directory to the directory containing the first file found.

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