

CROCO Webinars #1

Python pre-processing tools for CROCO – 2.0.3

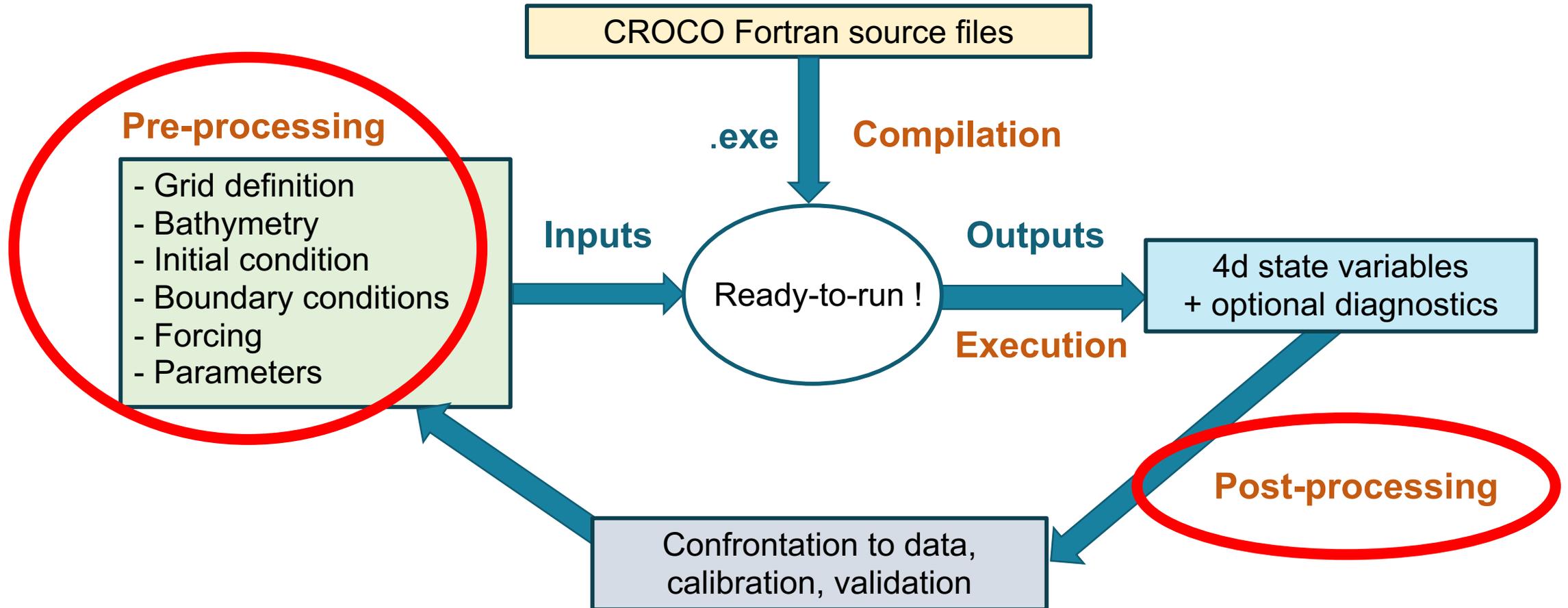
Friday, February 6th, 2026 at 1:30 PM (France time)

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On behalf of the CROCO pytools development team



Communauté
d'universités
et établissements
de Toulouse

Why do we need tools ?



What is CROCO_PYTOOLS ?

It's a python toolbox for pre-processing, post-processing, and visualization for CROCO

It includes :

- **prepro**: preparation of input files required to run realistic CROCO simulations
- **xcroco**: analysis of CROCO outputs using xarray and xgcm



Code repository:

https://gitlab.inria.fr/croco-ocean/croco_pytools



Documentation:

https://croco-ocean.gitlabpages.inria.fr/croco_pytools

Forum:

<https://forum.croco-ocean.org/>

Installation

env.yml → python environment → f2py compilation → Ready to use



Python environment :

Pre-configured environment provided: env.yml

Includes all Python dependencies (e.g., NumPy, matplotlib, f2py)

Create and activate, example with conda:

```
conda env create -f env.yml
conda activate croco_pyenv
```

Fortran tools, interfaced through f2py : must be compiled

Some routines are written in Fortran and must be compiled

Compilation example:

```
cd prepro/Modules/tools_fort_routines
make
```

Reference and tutorial :

https://croco-ocean.gitlabpages.inria.fr/croco_pytools/prepro/tuto.env.html



Improve on
Apple chip

Preprocessing pipeline

Splitting the preparation of CROCO inputs into independent steps:

Grid > Initial conditions > Boundary conditions > (Tides) > (Rivers)

CROCO_FILES
READY!



What CROCO_PYTOOLS/prepro provides

- Python scripts & notebooks (`make_*.py`, `nb_*.ipynb`)
- Configurable via `.ini` files (see Examples)
- Data readers (variable mapping dictionary `readers.jsonc`)
- Download tools included (`download_*.py`)



Use separate `.ini` files per step, or a single merged configuration file for the whole pipeline

Downloading data

Scripts available for : **ERA5**, **GLOFAS**, **HYCOM** and **Mercator** products

Use `-h` or `--help` to have usage informations

Configuration file with parameters :

- to request an extent from a CROCO grid netcdf file or a specified area,
- to choose a product and the variables needed
- to request a specific period of time ...

Download scripts can be long to run if you request a large amount of data



Accounts needed :

- ERA5: define your `~/.cdsapirc` with key & url `https://cds.climate.copernicus.eu/api`
- GLOFAS: define your `~/.cdsapirc` with key & url `https://ewds.climate.copernicus.eu/api`
- MERCATOR: provide login and password directly with command options or provide credentials file `~/.copernicusmarine/.copernicusmarine-credentials`



Follow up
copernicus API

Make grid

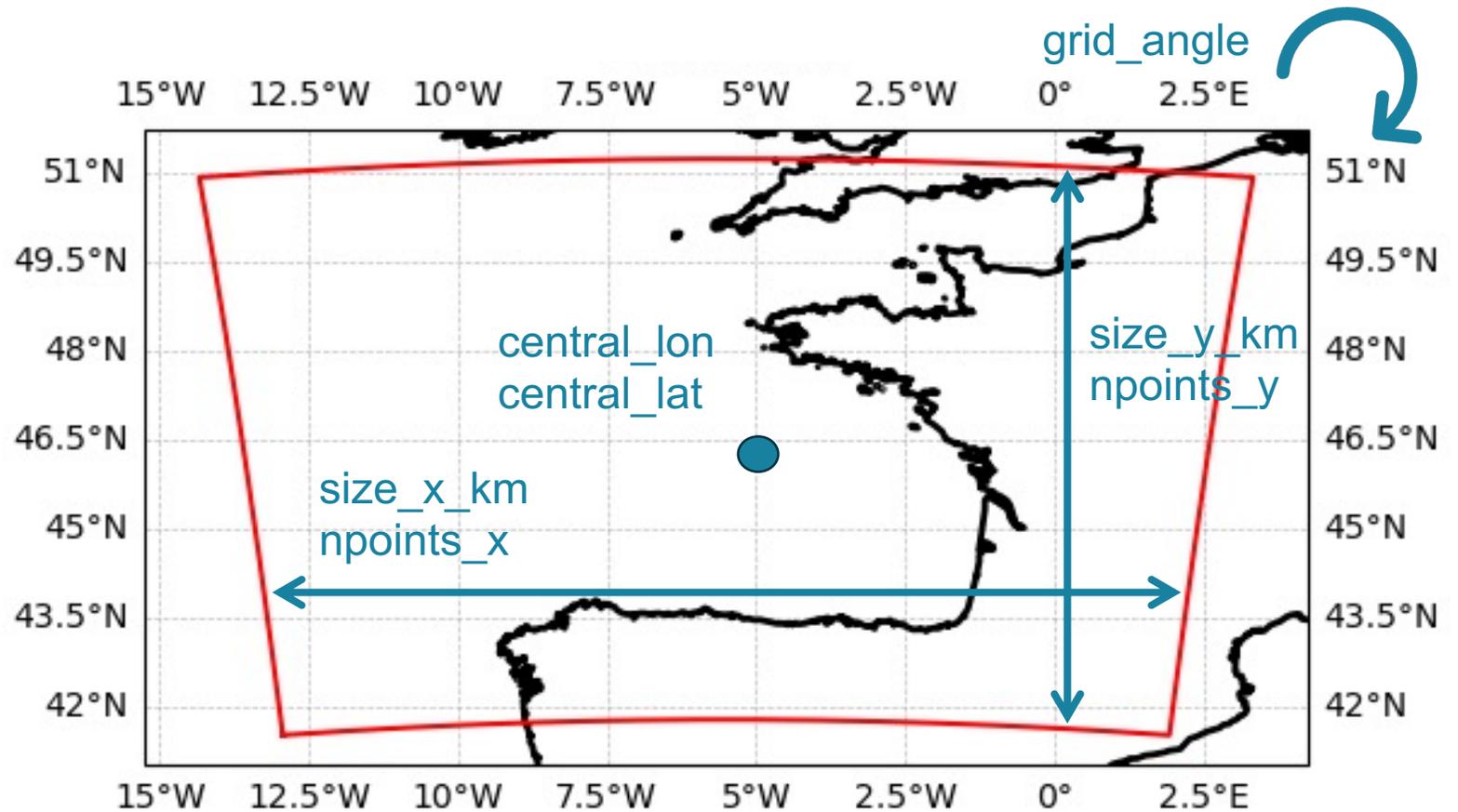
How to make a grid ? `./make_grid.py my_grid_configuration.ini`

Tutorial in the documentation

To remember:

- Input data must be gridded (raster only, no vector data)
- Grid uses Oblique Mercator projection
- Several smoothing methods are available
- Isolated waterbodies can be masked

 croco_grd.nc

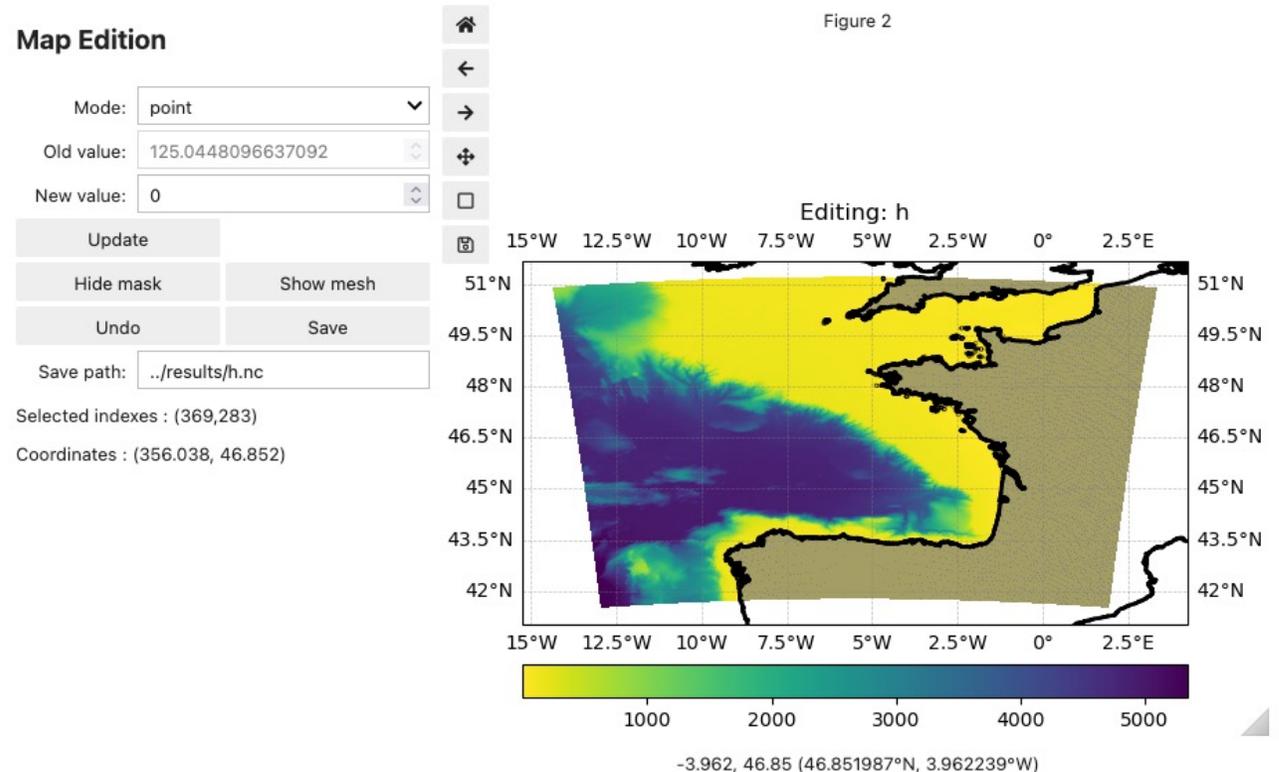


Edit the result with grid widget

How to update bathymetry « h » or mask « maskr » ? [Editor demonstration](#)

To remember :

- Point or rectangle selection
- Leave zoom mode to select a point
- Save button does NOT update croco_grd.nc directly
- After editing, bathymetry smoothing should usually be applied again



Use jupyter-forward if you are on remote computer

Make_ini and make_bry

How to generate the initial and boundary conditions ?

```
./make_ini.py my_ibc_configuration.ini  
./make_bry.py my_ibc_configuration.ini
```

Interpolation :

Z-coordinate > S-coordinate : general case

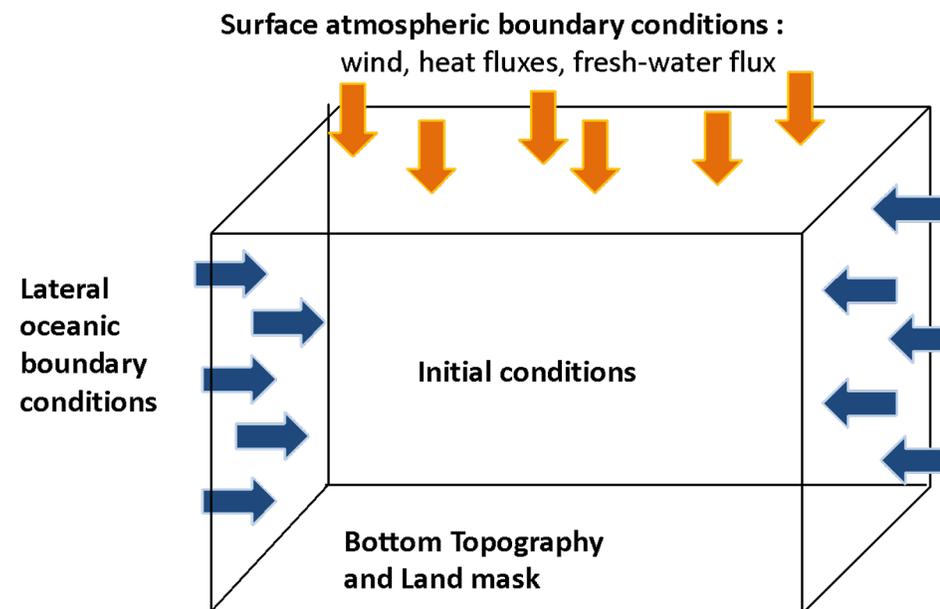
S-coordinate > S-coordinate : zoom offline case

To remember :

- S-coordinate parameters must be coherent with CROCO parameters (N, hc, theta_s, theta_b)



```
croco_ini_xxx_YxxxxMxx.nc  
croco_bry_xxx_YxxxxMxx.nc
```



Example with
BGC tracers
available

Make_tides

How to make tidal forcing file ?

```
./make_tides.py my_tides_configuration.ini
```

Careful with CROCO param coherency :

- Ntides
- Precompilation keys : #TIDE_MAS #USE_CALENDAR #SSH_TIDES #UV_TIDES #POT_TIDES

Check the tide atlas use : Re_Im or Amp_phase format

Check readers.jsonc file

Unless using TIDEMAS, correction at a fixed date, update to do at every restart file for long run



`croco_frc_xxx.nc`



All FES tide waves now available with correction

Make_rivers

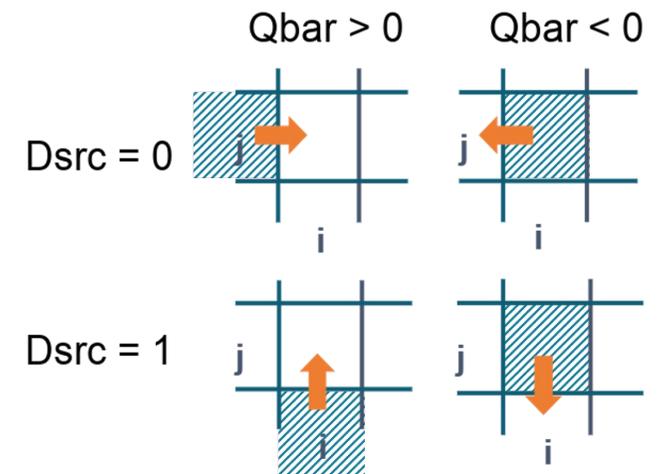
How to make runoff forcing file ?

```
./make_rivers.py my_rivers_configuration.ini
```

Rivers data can have different format :

- 2D netcdf file, like GLOFAS product
- 1D netcdf file (temporal series of one river)
- 1D netcdf file (temporal series of several rivers)
- Text file (one for each river)

| # input_file | Lon | Lat/Qmin |
|--------------|-------|----------|
| river.nc | -2.05 | 47.30 |
| loire.dat | -2.05 | 47.30 |
| Rhone.txt | 4.82 | 43.37 |
| 2ddata.nc | X | 10 |



Make_rivers output

- croco_runoff.nc
- text file to complete croco.in psource section

Rivers widget

How to update rivers ? Editor demonstration

To remember :

- Leave zoom mode to select a river
- On save, misplaced rivers are moved
- This widget does NOT support PSOURCE_MASS option

Rivers Edition

Hide summary

Number of rivers: 5 Number of points: 7 Number of splitted rivers: 2

| River | i | j | lon | lat | dsrc | qbardir |
|--------------------------|-----|-----|--------|--------|------|---------|
| cems_glofas_ALL_0 | 171 | 17 | -8.880 | 42.026 | 0 | -1 |
| cems_glofas_ALL_1 | 474 | 107 | -1.454 | 43.620 | 0 | -1 |
| cems_glofas_ALL_2_split0 | 498 | 198 | -0.711 | 45.220 | 1 | 1 |
| cems_glofas_ALL_2_split1 | 497 | 198 | -0.736 | 45.221 | 1 | 1 |
| cems_glofas_ALL_3 | 437 | 305 | -2.163 | 47.208 | 0 | -1 |
| cems_glofas_ALL_4_split0 | 517 | 434 | 0.195 | 49.440 | 0 | -1 |
| cems_glofas_ALL_4_split1 | 517 | 433 | 0.193 | 49.423 | 0 | -1 |

River: None selected

Left click to select and move, right click to unselect

dsrc:

qbardir:

Delete selection Add point to selecti...

Show mask Show mesh

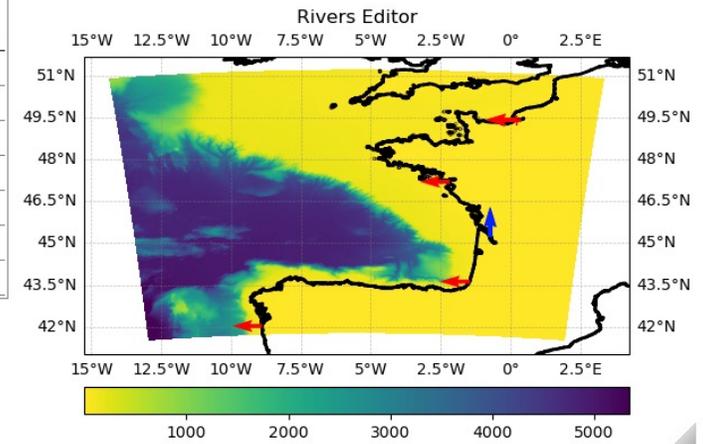
Undo

Save files

Rivers IN:

Rivers NC:

Figure 5



Use jupyter-forward if you are on remote computer

Last release 2.0.3 (early February 2026)

- Improve installation on Apple chip
- Update python environment to follow copernicus API change
- Add example for handling of other tracers than T,S in `make_ini`, `make_bry`
- Improve FES tide atlas handling
- Bugs fixes

For now ...

Still no widget to create grid, no multi-files handling to interpolate grid and no vector data (soundings) handling, no creation of river widget (just modification), no tracers in rivers input, only ONLINE atmospheric forcing available... ..

... **Work in progress !**



Feedback welcome ! <https://forum.croco-ocean.org/>

To conclude ...



Code download

https://gitlab.inria.fr/croco-ocean/croco_pytools/-/releases



Help and details. Try Tutorials !

https://croco-ocean.gitlabpages.inria.fr/croco_pytools



Feedback welcome !

<https://forum.croco-ocean.org/>

Questions

Jupyter-forward example

Install jupyter-forward locally, see details in :

<https://jupyter-forward.readthedocs.io/en/stable/>

Example of command line :

```
jupyter-forward
  --port 8888
  --conda-env "/path_env_xxx"
  --notebook-dir "/path_nb_xxx/"
  --port-forwarding
  --shell "/bin/bash"
  --launch-command "qsub -N jlab -j oe -l select=1:ncpus=1:mem=8g -l walltime=9:00:00"
  xxx@xxx
```

Adding BGC tracers

If all tracers are in the same product, just add the tracers in the list and check the readers.jsonc file.

If the tracers are in a different product than classical fields, make 2 downloads, see Examples/benguela_multifiles

Then in ibc.ini use prefix to indicate the right product where the variables are available

```
[IBC_Input_Files]
ibc_dir = ../data/DATA_IBC/GLORYS/
ibc_prefix = GLORYS_PHY

[Mercator_Download]
dataset = cmems_mod_glo_phy-all_my_0.25deg_P1
variables = ["vo_glor", "vo_glor", "thetao_glor"]
depths = [0, 5000]
```

```
[IBC_Input_Files]
ibc_dir = ../data/DATA_IBC/GLORYS/
ibc_prefix = GLORYS_BIO

[Mercator_Download]
dataset = cmems_mod_glo_bgc_my_0.25deg_P1M-m
variables = ["chl", "o2", "no3", "po4", "si"]
depths = [0, 5000]
```

```
[IBC_Options]
obc_dict = {'south':1, 'north':1, 'west':1, 'east':1}
tracers = ["temp", "salt", "chl", "o2", "no3", "po4", "si"]
uv_conserv = 1
min_nb_valid_data = 4

[IBC_Input_Files]
ibc_reader = mercator_glorys_0.25deg
ibc_dir = ../data/DATA_IBC/GLORYS
ibc_prefix = GLORYS
ibc_extension = .nc
ibc_freq = 1M
ibc_multi_files = True
ibc_file_ssh = PHY
ibc_files_tracers = ["PHY", "PHY", "BIO", "BIO", "BIO", "BIO", "BIO"]
ibc_file_u = PHY
ibc_file_v = PHY
ini_filedate = Y2013M01
ini_idx = 0
```

Ini format

An INI file is a plain text file commonly used to store configuration settings.

It is a simple and widely supported format that organizes information into sections and key-value pairs.

Example :

```
[Croco_Files]
croco_files_dir = ../results/CROCO_FILES
croco_grd_prefix = croco_grd
```

```
[Zoom_Options]
is_zoom = False
is_agrif = False
agrif_level = 0
```

```
[Grid_Position]
central_lon = 15.0
central_lat = -32.0
size_x_km = 1556
size_y_km = 1334
npoints_x = 41
npoints_y = 42
grid_angle = 0.0
```

Rivers input format

4 types :

- 2D netcdf file :
 - .nc extension
 - Qbar = variable with unit attribute in ['m³ s⁻¹', 'm^{**3} s^{** -1}', 'm³/s']
 - lon/lat : search by name and unit
- 1D netcdf file with one river :
 - .nc extension
 - Qbar = variable with unit attribute in ['m³ s⁻¹', 'm^{**3} s^{** -1}', 'm³/s']
- 1D netcdf file with several rivers :
 - .nc extension
 - Qbar = variable with unit attribute in ['m³ s⁻¹', 'm^{**3} s^{** -1}', 'm³/s']
 - lon/lat : search by name and unit
- text file : other extension than .nc

```
1950/01/01 12:00:00 730.000000
1950/01/02 12:00:00 695.000000
1950/01/03 12:00:00 675.000000
1950/01/04 12:00:00 645.000000
1950/01/05 12:00:00 610.000000
```

Readers.jsonc

```
{
  "Topo": {
    "etopo2022": {
      "lon": "lon",
      "lat": "lat",
      "topo": "z",
      "zaxis": "up"
    },
  },
  ...
  "Tides": {
    "tpxo9": {
      "lonr": "lon_z",
      "lonu": "lon_u",
      "lonv": "lon_v",
      "latr": "lat_z",
      "latu": "lat_u",
      "latv": "lat_v",
      "H_r": "hRe",
      "H_i": "hIm",
      "U_r": "uRe",
      "U_i": "uIm",
      "V_r": "vRe",
      "V_i": "vIm",
      "topor": "h",
      "topou": "hu",
      "topov": "hv",
      "invert_phase": true
    },
  },
  ...
  "IBC": {
    "mercator": {
      "depth": "depth",
      "lonr": "longitude",
      "lonu": "longitude",
      "lonv": "longitude",
      "latr": "latitude",
      "latu": "latitude",
      "latv": "latitude",
      "ssh": "zos",
      "temp": "thetao",
      "salt": "so",
      "u": "uo",
      "v": "vo",
      "time": "time",
      "time_dim": "time"
    },
  },
}
```