

## **Wikiprint Book**

**Title: FAQ**

**Subject: TracMeteo - WRF4GFAQ**

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## FAQ

Through this page we will try to answer the most common questions about how to use WRF4G.

If you execute `wrf4g_submit` and see this error. You probably has GridWay in **Stopped status**:

```
[user@mycomputer~]$ wrf4g_submit -e test
connect(): Connection refused
FAILED: failed connection to gwd
```

In order to solve the problem, you have to first check out the `wrf4g_framework` status.

```
[user@mycomputer~]$ wrf4g_framework status
GridWay Stopped
MySQL Stopped
```

Then you have to execute:

```
[user@mycomputer~]$ wrf4g_framework start
Starting GridWay..... OK
Starting MySQL..... OK
```

- You want to modify the features of your experiment that you created it before. The new experiment will have the same name and different features that the old experiment.

```
[user@mycomputer~]$ wrf4g_prepare
Warning: You are using resources.wrf4g located in the /home/user/WRF4G/etc/ directory.
Preparing namelist...
WRFV3/run/namelist.input
WRF Check Warning: CAM radiation selected but paerlev/levsiz/cam_abs_dim1/cam_abs_dim2 was not set. Fixing...
WRF Check Warning: radt is shorter than dx (0.500000)

---> Single params run
---> Continuous run
    ---> cycle_chunks: test 2011-08-28_12:00:00 2011-09-01_00:00:00
        ---> chunks 1: test 2011-08-28_12:00:00 2011-08-29_00:00:00
        ---> chunks 2: test 2011-08-29_00:00:00 2011-08-29_12:00:00
        ---> chunks 3: test 2011-08-29_12:00:00 2011-08-30_00:00:00

[user@mycomputer~]$ wrf4g_prepare
Warning: You are using resources.wrf4g located in the /home/carlos/WRF4G/etc/ directory.
Experiment already exists
```

You need to execute `wrf4g_prepare --reconfigure`.

```
[user@mycomputer~]$ cat experiment.wrf4g | grep "end_date=\""
end_date="2011-08-30_00:00:00"

[user@mycomputer~]$ cat experiment.wrf4g | grep "end_date=\""
end_date="2011-09-01_00:00:00"

[user@mycomputer~]$ wrf4g_prepare --reconfigure
Warning: You are using resources.wrf4g located in the /home/carlos/WRF4G/etc/ directory.
Preparing namelist...
WRFV3/run/namelist.input
WRF Check Warning: CAM radiation selected but paerlev/levsiz/cam_abs_dim1/cam_abs_dim2 was not set. Fixing...
WRF Check Warning: radt is shorter than dx (0.500000)
```

```

----> Single params run
----> Continuous run
----> cycle_chunks: test 2011-08-28_12:00:00 2011-09-01_00:00:00
----> chunks 1: test 2011-08-28_12:00:00 2011-08-29_00:00:00
----> chunks 2: test 2011-08-29_00:00:00 2011-08-29_12:00:00
----> chunks 3: test 2011-08-29_12:00:00 2011-08-30_00:00:00
----> chunks 4: test 2011-08-30_00:00:00 2011-08-30_12:00:00
----> chunks 5: test 2011-08-30_12:00:00 2011-08-31_00:00:00
----> chunks 6: test 2011-08-31_00:00:00 2011-08-31_12:00:00
----> chunks 7: test 2011-08-31_12:00:00 2011-09-01_00:00:00

```

The experiment **test1** has finished with an error and you want to run the experiment again. What should you execute on shell?

```

[user@mycomputer~]$ wrf4g_submit --rerun -f -e test1
Submitting realization: "test1"
Submitting Chunk 1:    2011-08-28_12:00:00    2011-08-29_00:00:00
Submitting Chunk 2:    2011-08-29_00:00:00    2011-08-29_12:00:00
Submitting Chunk 3:    2011-08-29_12:00:00    2011-08-30_00:00:00

```

Imagine, you want to resubmit the chunk number **1** of the realization **test2**. In this case, it is highly recommended that you use the option `--dry-run` of `wrf4g_submit` command before you submit your chunk. If you want to make sure you are submitting that chunk.

```

[user@mycomputer~]$ wrf4g_submit --dry-run --rerun -c 1 -f -r test2
Submitting realization: "test2"
Submitting Chunk 1:    2011-08-28_12:00:00    2011-08-29_00:00:00

[user@mycomputer~]$ wrf4g_submit --rerun -c 1 -f -r test2
Submitting realization: "test2"
Submitting Chunk 1:    2011-08-28_12:00:00    2011-08-29_00:00:00

```