

## Generate a run script

### First step

There are several pre defined experiments available in the run-directory, listed as `exp.<experiment-name>`. To generate a run script for one of these experiments do

```
./make_runscripts <experiment-name>
```

Be careful not to forget `experiment-name`. `make_runscripts` will generate **all** possible runscripts otherwise. `make_runscripts` only works after configuring the model, as it needs the proper information of the compiler used.

The resulting runscript `exp.<experiment-name>.run`, for some experiments also a postprocessing script `post.<experiment-name>.run`, will be stored in the run-directory and might need additional editing.

### Use your experimenter ID

In general and to simplify collaboration it is an established way to copy a given script according to our **personalized naming convention**, see [List of IDs](#).

### Adaptations for accounting and parallelization

#### Example: Run the script as developer abc on Levante:

Copy the `exp.amip` experiment-template to your personal experiment-id:

```
cp run/exp.aes_amip run/exp.abc0001
./make_runscripts abc0001
cd run
```

Edit the file `exp.abc0001.run` and change the model settings (account e.g. `mh0287`):

```
#SBATCH --account=mpiaes
to
#SBATCH --account=mh0287
```

```
#SBATCH --nodes=1
to
#SBATCH --nodes=100
```

This number of nodes fit for the default setting of the script, which is R2B8 resolution.

Submit the job:

```
sbatch exp.abc0001.run
```

You can check the status of your jobs with:

```
squeue -u <user>
```

Results will be written to `icon-mpim/experiments/abc0001`

**In case you want to run one month** of this experiment in R2B8 you have to adapt the run time in the run script: Set the run time of the job to at least 5 hours, also depending on your amount of output:

```
#SBATCH --time=05:00:00
```

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— *Monika Esch* 2024/08/19 12:19

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