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Quickstart

Please prepare your job_script which should be capable of storing and loading configuration snapshots and of returning the relevant metadata for the particular sampling method (at least reaction coordinate and calculation steps/time).

Server

Open a terminal and start the server (you may edit the configuration file first):

```
cd freshs/server
python main_server.py -c server-sample.conf
```

Client(s)

Open another terminal and start the client, e.g.:

```
cd freshs/client
python main_client.py -c client-sample.conf
```

Thereby, the executable of your simulation and the harness script are read from the configuration file. According to your resources, you can connect multiple clients to the server.

MPI

If you'd like to use a prefix for your executable, e.g. mpirun, then use

```
python main_client.py -c client-sample.conf -e "mpirun -np 8"
```

SSH tunnel

If you need a secure ssh tunnel to your local machine, the ssh tunnel can be enabled and the tunnelcommand can be specified in the configuration file of the client:

```
# ssh tunnel
ssh_tunnel = 1
```

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ssh tunnelcommand

ssh_tunnelcommand = **ssh** -N -L 10000:localhost:10000 tunneluser@tunnelhost

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