

## INC1609499 - Update - Help with BEAR compute

From Service Desk <servicedesk@servicedesk.bham.ac.uk>

Date Wed 18/06/2025 16:39

To Anubrata das (Cancer and Genomic Sciences) <a.das.1@bham.ac.uk>



UNIVERSITYOF BIRMINGHAM

IT SERVICES

Reference Number: <u>INC1609499</u>

Short Description: Help with BEAR compute

Dear Anubrata,

There's been an update to your call. Here's what's been added:

## 18-06-2025 16:38:36 BST - Chris Greasley (Advanced Research Computing)

User Notes

At some point in the pipeline, Matplotlib should be told how to cache files. How is this being configured at present?

I don't know what your config file looks like, but you could workaround this issue with the pipeleine and expose your home directory in the nexflow config file:

Something like:

```
apptainer {
runOptions = "-B ${HOME}"
}
```

Or if you wanted to leave your home directory from being exposed, you could define a new location for the matplotlib cache and bind that in the nextflow config. (This would need to already exist on your host system)

```
e.g.
env {
MPLCONFIGDIR = "${PWD}/.apptainer_cache/matplotlib"
}
apptainer {
runOptions = "-B ${PWD}/.apptainer_cache"
}
```

If you want to see a default config for working on BlueBEAR, have a look at https://github.com/nf-core/configs/blob/master/conf/bluebear.config.

Best regards, chris

**Attachments** (requires login): matplotlib\_directory.png

## **Thank You**

IT Services University of Birmingham Want to give us feedback?





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