

# Optimising R code for BlueBEAR

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# Optimising R code for BlueBEAR

## Topics

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- R packages on BlueBEAR
- R parallel workflows


# What is BlueBEAR?

- UoB's "tier 3" HPC (high-performance computing) system
  - Also HTC workflows: high-throughput computing
- It's heterogeneous:
  - What does this mean?
  - Why is this important for R package installs?
- [BEAR Portal](#)  = graphical frontend to BlueBEAR (sort of).

# R Package Installs: part 01

## Do you need to install?




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- Check if we've already installed package via a module.
  - <https://bear-apps.bham.ac.uk>  [demo]
  - N.B. changes 2023a and above.
- RStudio (with “meta” modules).
- JupyterLab – R kernels with modules tool. [demo]

# R Package Installs: part 02

## R package install: process overview

Don't use RStudio!



- Start an [interactive job](#) 
  - Where to run? (See “Heterogeneous issues” below.)
  - How long for?
  - Alternatively, use a [batch job](#)  to run the install. (Reproducible!)
- Follow the documentation here:  
[https://docs.bear.bham.ac.uk/bluebear/software/self\\_installs\\_r/](https://docs.bear.bham.ac.uk/bluebear/software/self_installs_r/) 

### Heterogeneous issues


- Pay close attention to the node type on which you're running the session.
- You may need to install the R packages multiple times.

# R Package Installs: part 03

Other options...

- R Conda environments
  - <https://astrobiomike.github.io/R/managing-r-and-rstudio-with-conda> 
  - The same heterogeneous issues apply!
- Python dependency stack
  - *reticulate*  package.
  - The same heterogeneous issues apply!

# R Parallel Jobs: part 01






- What speeds-up a job?
  - More resource...
    - ... if the script is written to parallelise.
    - ... if the workflow is not storage-bound.
  - Putting the data in the correct location.
    - RDS (network storage) is relatively slow.
    - [Local \(/scratch\) storage](#)  is much faster.
    - ... but memory is also fast, although has less capacity.

# R Parallel Jobs: part 02

- Note that array jobs may be a good solution for some types of workflow.
  - [https://docs.bear.bham.ac.uk/bluebear/array\\_jobs/](https://docs.bear.bham.ac.uk/bluebear/array_jobs/) ↗
- Our recommendation: single-node parallelisation is best for R.
- Accurate detection of cores: <https://docs.bear.bham.ac.uk/bluebear/software/guides/R/#accurate-detection-of-cores> ↗

```
library(parallelly)
library(parallel)
# Detect how many cores are available for R to use
n <- availableCores()
cat("Using", n, "cores\n")
# Create a PSOCK cluster using all available cores
cl <- makeClusterPSOCK(n)
# Function each worker will run
hello_fun <- function(i) {
paste("Hello from worker", i, "on", Sys.info()[["nodename"]])
}
# Run in parallel
results <- parLapply(cl, seq_len(n), hello_fun)
print(results)
# Shut down the cluster
stopCluster(cl)
```

# R Parallel Jobs: part 03 – existing tools

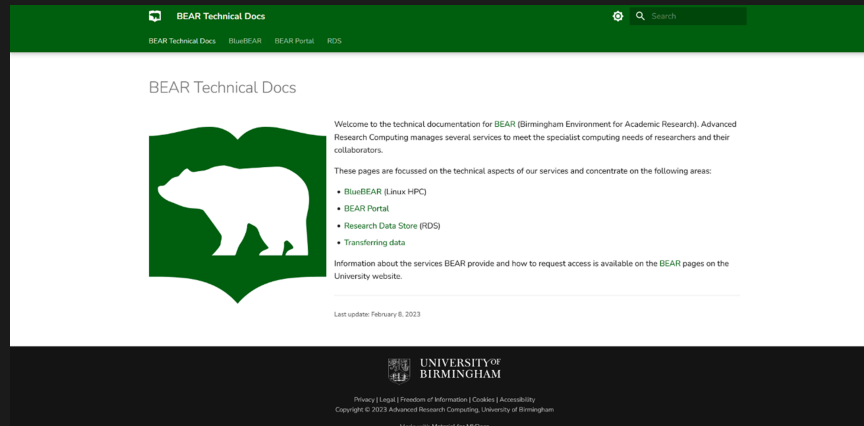
- <https://mirai.r-lib.org/> 
  - Cluster config: [https://mirai.r-lib.org/reference/cluster\\_config.html](https://mirai.r-lib.org/reference/cluster_config.html) 
- [future](#) 
  - Slurm example: <https://future.batchtools.futureverse.org/> 
- BiocParallel
  - Available on BlueBEAR: <https://apps.bear.bham.ac.uk/applications/R-bundle-Bioconductor/> 

# Miscellaneous

- Batch jobs over interactive RStudio / JupyterLab.
- Why has my R Studio session died?!
  - OOM-kills.
    - Accumulation of variables.
    - Cores and memory.
- GPU jobs:
  - Batch job or interactive job only. No RStudio.
  - We don't have any R modules built with GPU capability.
    - Historically, much of the GPU functionality was provided via Python packages. (See again *reticulate*).

# Support

# BEAR Technical Docs



BlueBEAR Technical Documentation Site: <https://docs.bear.bham.ac.uk>

- The BEAR Technical Docs website should be your first resource for BEAR questions
- Continually updated...
  - ... and please let us know if anything is unclear or needs further explanation

## IT Service Desk

- <https://www.birmingham.ac.uk/bear/sd/other-bear-request>
  - Questions and issues running jobs
  - Advice
- <https://www.birmingham.ac.uk/bear/sd/new-bear-software>
  - Use to request new software on BlueBEAR with a link to the software and preferred version

 Visibility for Masters students

Masters students cannot see some research tickets so should submit a general ticket, clearly stating that it's for BEAR.

Any Questions??