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## **1 Provenance of the image data files**

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Where downloaded, what done for downsampling.

## **2 Session Info**

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- R version 4.1.1 (2021-08-10), x86\_64-pc-linux-gnu
- Locale: LC\_CTYPE=en\_US.UTF-8, LC\_NUMERIC=C, LC\_TIME=en\_GB, LC\_COLLATE=C, LC\_MONETARY=en\_US.UTF-8, LC\_MESSAGES=en\_US.UTF-8, LC\_PAPER=en\_US.UTF-8, LC\_NAME=C, LC\_ADDRESS=C, LC\_TELEPHONE=C, LC\_MEASUREMENT=en\_US.UTF-8, LC\_IDENTIFICATION=C
- Running under: Ubuntu 20.04.3 LTS
- Matrix products: default
- BLAS: /home/biocbuild/bbs-3.14-bioc/R/lib/libRblas.so
- LAPACK: /home/biocbuild/bbs-3.14-bioc/R/lib/libRlapack.so
- Base packages: base, datasets, grDevices, graphics, methods, stats, utils
- Other packages: MSMB 1.12.0, knitr 1.36
- Loaded via a namespace (and not attached): BiocManager 1.30.16, BiocStyle 2.22.0, compiler 4.1.1, digest 0.6.28, evaluate 0.14, fastmap 1.1.0, htmltools 0.5.2, magrittr 2.0.1, rlang 0.4.12, rmarkdown 2.11, stringi 1.7.5, stringr 1.4.0, tools 4.1.1, xfun 0.27, yaml 2.2.1