

We will be running a workshop on single cell RNA-Seq analysis.

**Before being able to start the workshop**, you will need to install R and some dependencies and download the data. Details of how to do this are here: <https://github.com/IMB-Computational-Genomics-Lab/SingleCells2018Workshop>

Please let me <j.powell@garvan.org.au> know if you encounter issues with installing the dependencies.

The workshop vignette is available here: <https://github.com/IMB-Computational-Genomics-Lab/SingleCells2018Workshop/blob/master/SingleCellsWorkshop.md>