



- 1 First we have to find out the size of a piece of our pathway pairs that runs in the cluster without memory problems. This will serve as the size of each piece later on in this pipeline
- 2 An R script that splits the total pathways into runnable pieces (based on the above)
- 3 A bash script that fires a number of `improved_PCxN_estimates01.R` scripts simultaneously, each time using a different piece from step 2.
- 4 The basic `improved_PCxN_estimates01.R` script with some changes that deal with the different piece inputs.