* Handouts?
  + Command line cheat sheet ready for Monday
  + Peter almost done with AMR++ sheet for Tuesday
* During the R portions, should we try to show examples with both data types (microbiome, resistome)
* Does AMR++ include fastqc or multiqc? Both- fastqc on each sample and then multiqc to create a report with all samples.
* Will Lee go into detail about QC issues during 16S content? - Very briefly, Noelle to expand in lecture
* Learning objectives? Do we need them? When/where do we need them?
* Tuesday start == waldo lecture??
* What’s up with Marit participant? (see slack message)