Read me

* In the submission box, as well as the link, put some accompanying text on another line something like "tidy data as per the ReadMe that can be read into R with read.table(header=TRUE) {listing any settings you have changed from the default}" This is just to make it really easy for your reviewer.
* In the readMe in explaining what the script does put "and then generates a tidy data text file that meets the principles of ...etc"
* the truly cunning may want to put in a citation to this discussion and/or Hadley's paper

The codebook still has the specific description of the tidy data file contents (and you mention that it exists and it's role in the ReadMe)

the assignment says \_The code should have a file run\_analysis.R in the main directory that can be run as long as the Samsung data is in your working directory.\_ so it is fine to start with the decompressed UCI HAR folder in the working directory. The clever thing to do is to document the starting conditions in the readme.

2 use whatever packages you want to. You are the data scientist and it is up to your judgemen