

```
#####  
#NAME: Erin Nishimura  
#  
#DATE: 8/25/2016  
#  
#ASSIGNMENT: 1  
#####  
1) cd Dropbox/COURSES/2016_LINUX_computational_platform/DEVELOPMENT/Assignment1/Assignment1/  
   --> Many answers accepted  
  
2A) pwd  
2B) ls  
  
3) less genome_releases.README  
   --> more, nano also accepted  
  
4) cd S288C_reference_genome_R64-1-1_20110203  
  
5) ls *.fasta  
  
6) ls -Sl *.fasta  
   --> on some systems, only -s works; also took that  
  
7) mkdir 01_genome 02_fastfiles 03_annotations 04_other  
  
8A) mv *fsa 01_genome/  
8B) mv *.fasta 02_fastfiles/  
8C) mv *.gff 03_annotations/  
8D) mv *.sgd 04_other/  
   --> other versions accepted as long as they work  
  
9) mv ../genome_releases.README 00_README/  
   --> other versions accepted as long as it was clear that genome_releases.README had been in  
   the parent directory  
  
10)  
DATE: 160825  
AUTHOR: Erin Nishimura  
PROJECT: Download yeast genome  
  
Purpose:  
  I downloaded the yeast genome for Assignment #1 of the Linux class.  
  
URL:  
  http://downloads.yeastgenome.org/sequence/S288C\_reference/genome\_releases/  
  
Files downloaded:  
  genome_releases.README  
  S288C_reference_genome_R64-1-1_20110203.tgz  
  
Organization:  
  I reorganized the downloaded files into the directories:  
  00_README      01_genome      02_fastfiles  03_annotations 04_other  
  
Problems:  
  No problems encountered  
  
--> lots of answers accepted
```