R_Assignment

list.files()

```
## [1] "fang_et_al_genotypes.txt" "maize_data"
## [3] "R_Assignment.html" "R_Assignment.pdf"
## [5] "R_Assignment.Rmd" "R_Assignment.Rproj"
## [7] "README.md" "snp_position.txt"
## [9] "teosinte_data"
```