

### Ejercicio 1

```
$ head -n 1 file.vcf > merged.file  
$ cat file2.vcf >> merged.file
```

### Ejercicio 2

```
$ head -n 1 prokaryotes.txt > akkermansia.txt  
$ grep "Akkermansia" prokaryotes.txt >> akkermansia.txt  
$ head -n 10 akkermansia.txt  
$ grep -c "Akkermansia muciniphila" akkermansia.txt  
1205  
$ grep -c "Akkermansia biwaensis" akkermansia.txt  
1
```