Building multiple alignments and exploratory trees with MUSCLE and RAxML

Example command line for stand-alone installation of MUSCLE and RAxML

#MUSCLE commands for 2 iterations

```
muscle -in tufA.fas -seqtype auto -maxiters 2 -out tufA_aln.fasta
```

#RAxML for 10 restarts with 4 threads

```
raxml -s tufA_aln.fasta -n tufA -m GTRGAMMA -p 5463789 -T 4 -N 10
```