

Trifolium Pallescens genome annotation

Vikas Gupta and Stig U. Andersen

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1. Data

```
p <- path.expand("/Volumes/GenomeDK/LotusGenome/01_vgupta/11_clover/20140709_clover_palleszens/RogerMoraga")
setwd(p)
p

## [1] "/Volumes/GenomeDK/LotusGenome/01_vgupta/11_clover/20140709_clover_palleszens/RogerMoraga"

## File size
a <- system("ls -lh | awk '{print $9,$5}'", intern = TRUE)

for (i in a) {
  print(i)
}

## [1] " "
## [1] "F1_consensus.fasta.gz 54M"
## [1] "L4_consensus.fasta.gz 34M"
## [1] "T_pal_stolon_1_1_9_1.fastq.gz 3.8G"
## [1] "T_pal_stolon_1_1_9_2.fastq.gz 3.9G"
```