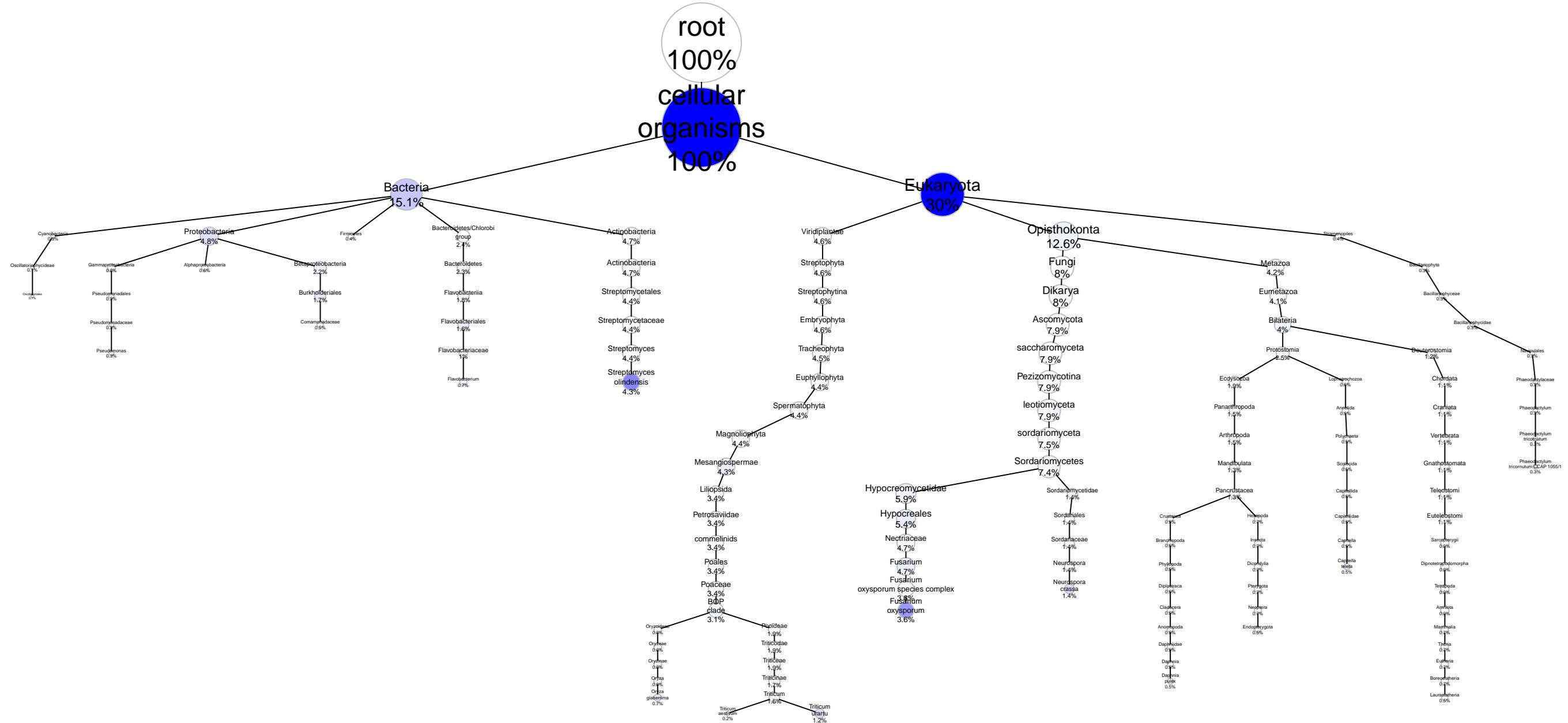


sample\_SM11\_norm1\_sequence\_fastq (total reads: 1441333)

















sample\_SM8\_uc3\_sequence\_fastq (total reads: 967825)

