

```
library(phyloseq)
data(GlobalPatterns)
```

```
source('embed_krona.R')
```

```
# if an up to date Krona is already installed and ktImportText is available in path:
#plot_krona(GlobalPatterns)
```

```
# otherwise, manually install KronaTools in the embed_krona directory and supply its path
plot_krona(GlobalPatterns, install_dir='.')
```

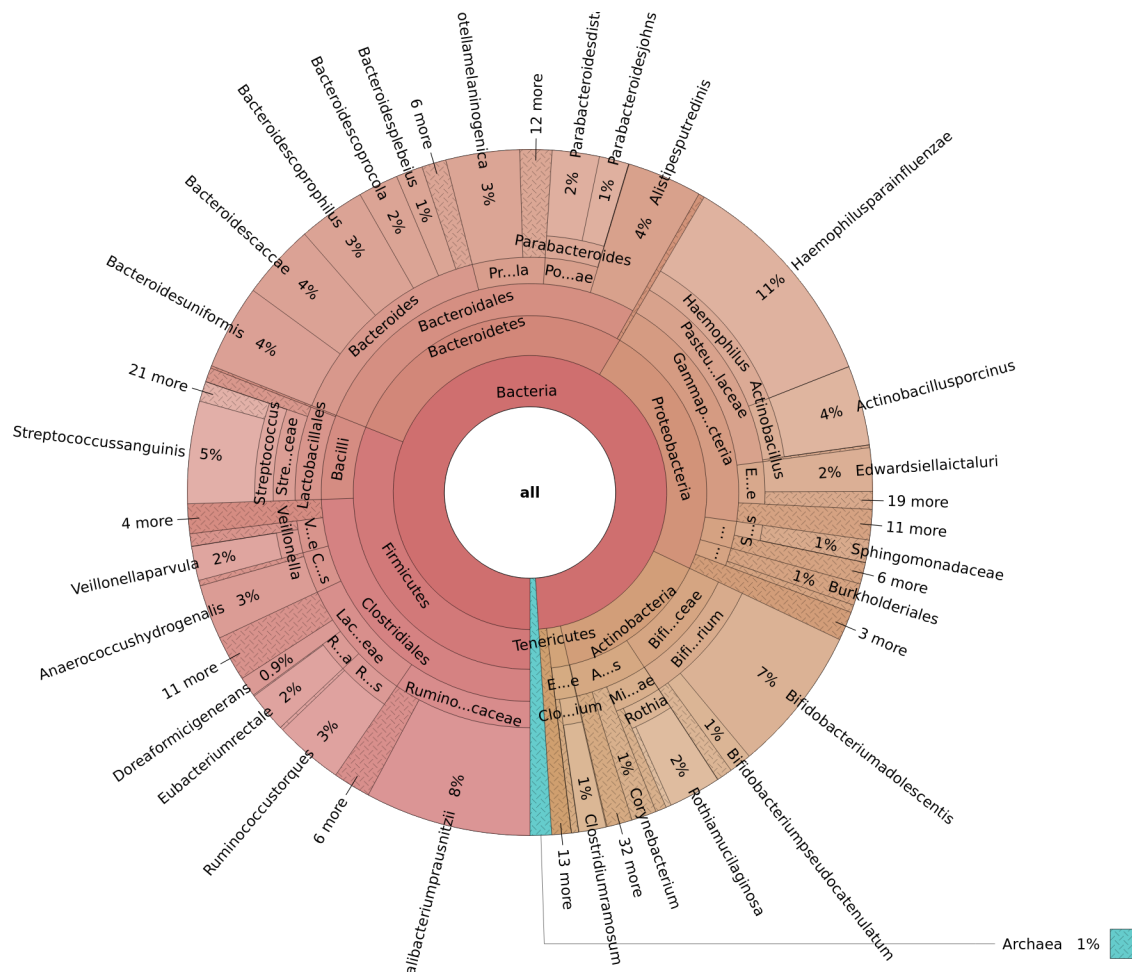


Figure 1: Overview of the GlobalPatterns dataset

```
# Note: the size of snapshots is best set using the snapshot_dim argument, but it
# is also possible to set the chunk option out.width='100%' for similar behavior
# across file formats.
```