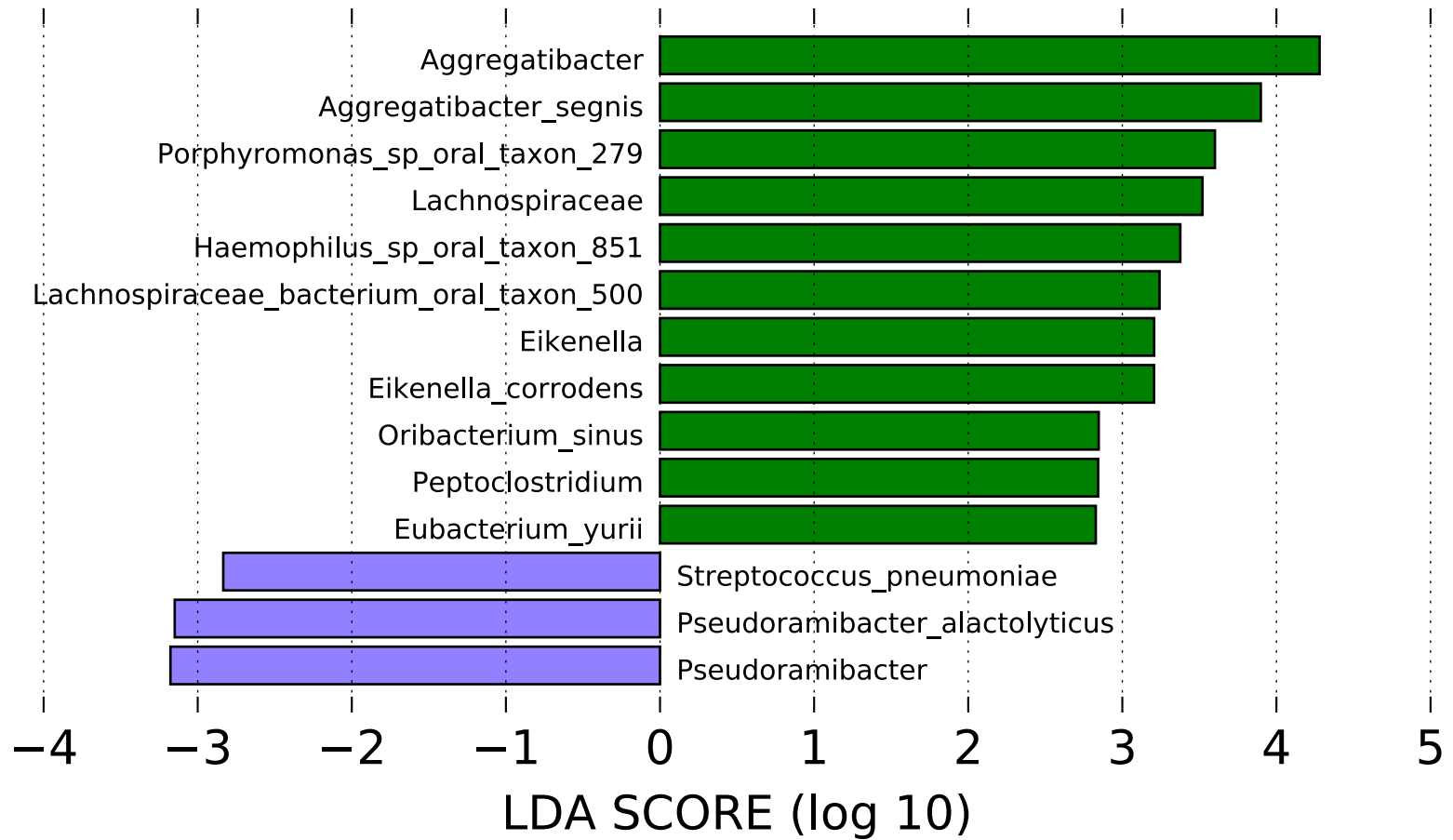


- Healthy control matching tumor site [control tumor - green]
- Buccal site from healthy controls [purple]



Supplementary Figure 1. Statistical differences in the phylogenetic composition of active communities.

Metatranscriptome hit counts were obtained using Kraken against an oral microbiome database.

Counts were then analyzed using LEfSe to identify significant differences at the species level between the microbial communities compared.

Healthy control matching tumor site [control tumor - green] vs. Buccal site from healthy controls [purple].