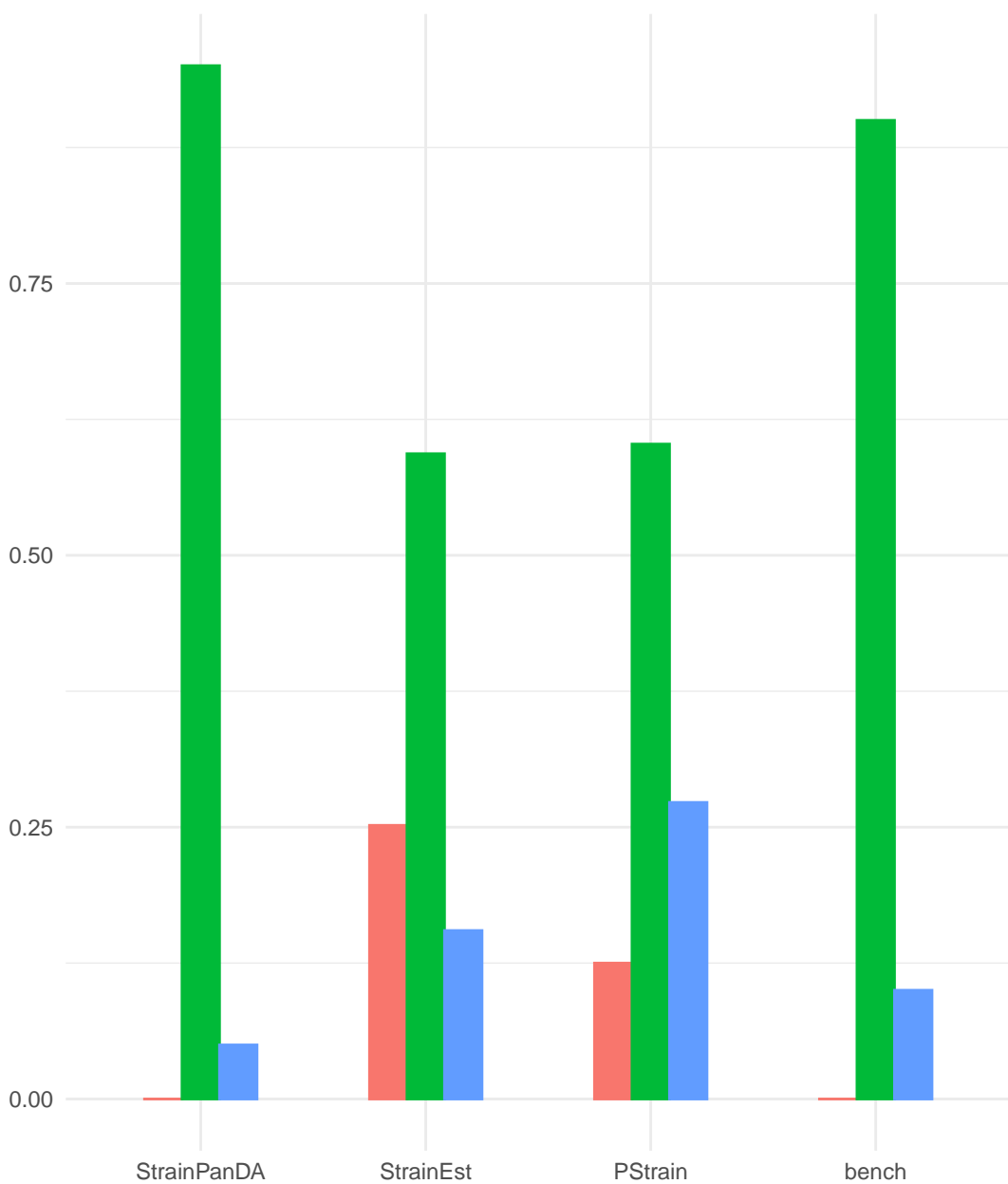


Relative_Abundance



Strains

all_FPs
GCA_str1
GCA_str2

Rank

all_FPs
GCA_str1
GCA_str2