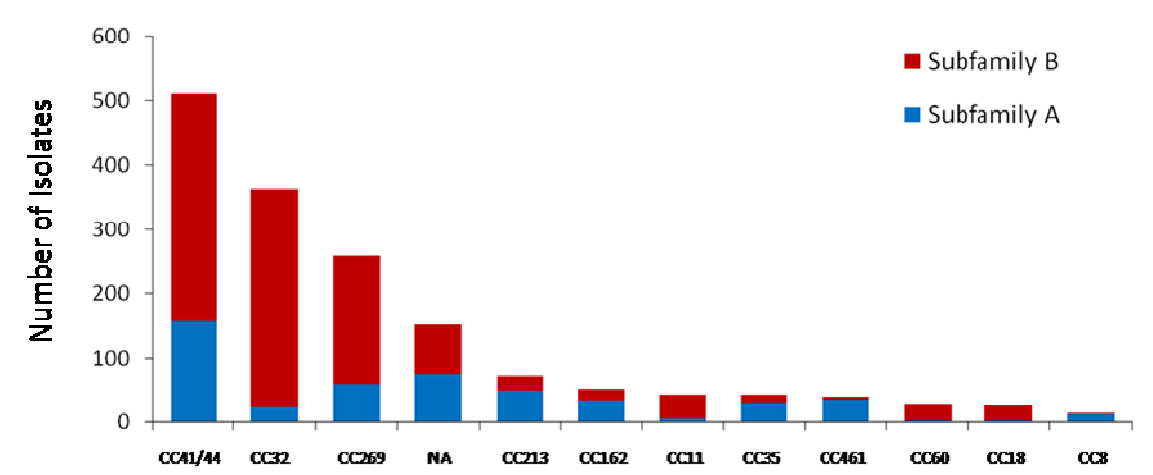


SUPPLEMENTAL FIGURE S4. Distribution of factor H binding protein (fHBP) subfamily (A) and variants (B) within clonal complexes (CCs). Data are shown for complexes with >7 strains. NA indicates sequence type (ST) not assigned to a complex. USA isolates are weighted for Oregon. Numbers on top of bars indicate the number of different fHBP variants found in each CC. The number in parenthesis represents the SDI.

A.



B.

