

Supplemental Digital Content 3. Relative abundances of OTUs assigned to genera containing potential pneumonia pathogens: a) *Streptococcus*, b) *Haemophilus*, c) *Moraxella*, d) *Staphylococcus*. The abundance of each OTU is shown relative to the total number of 16S ribosomal RNA gene sequences assigned to the genus in nasopharyngeal samples from the study population. OTU, operational taxonomic unit

