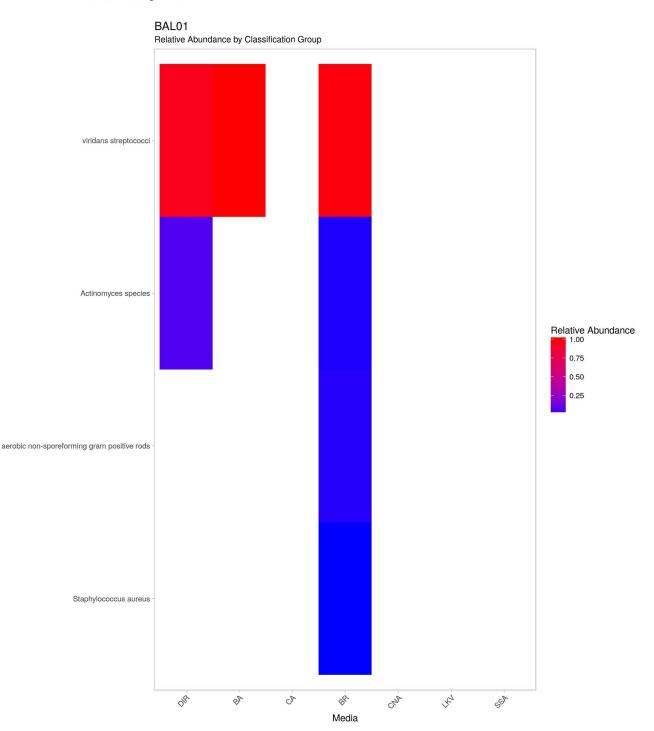
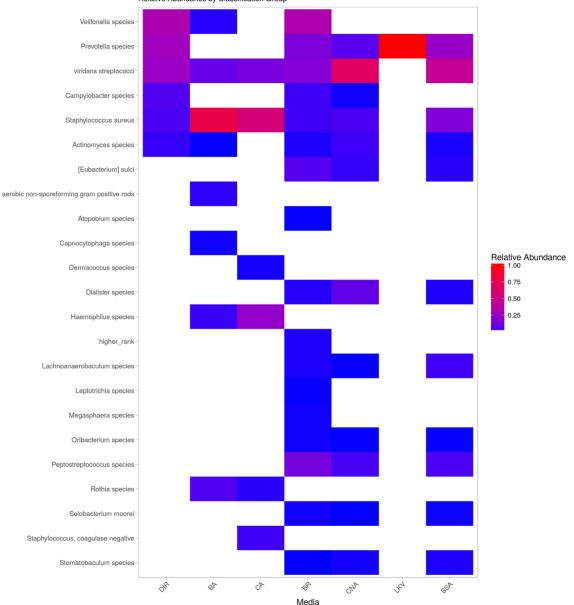
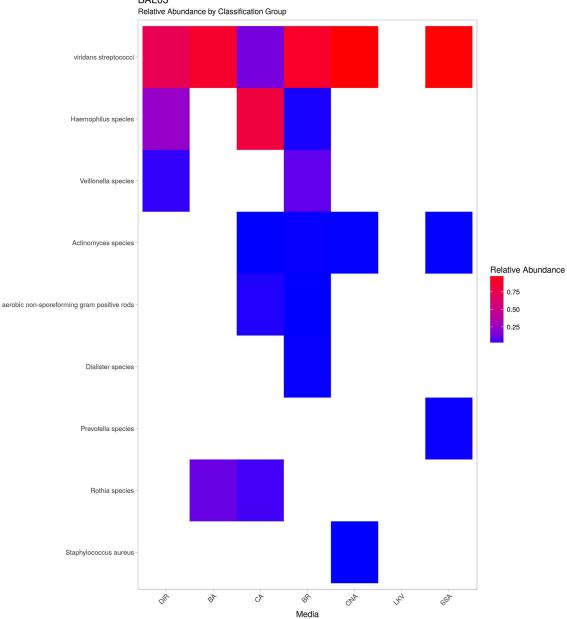
**Supplementary Figure S1**. Heat maps showing the CRCs identified by NGS analysis for each specimen. Individual heat maps showing the CRCs identivied by NGS16S for each specimen are shown (19 total; no heatmap was generated for BAL18 because no organisms were identified for that specimen). Relative abundance of each classification group is shown by media type; relative abundance is indicated by color. DIR = NGS16S sequencing of DNA extracted directly from the sample; plate washes from media as described in Figure 1.



BAL02

Relative Abundance by Classification Group

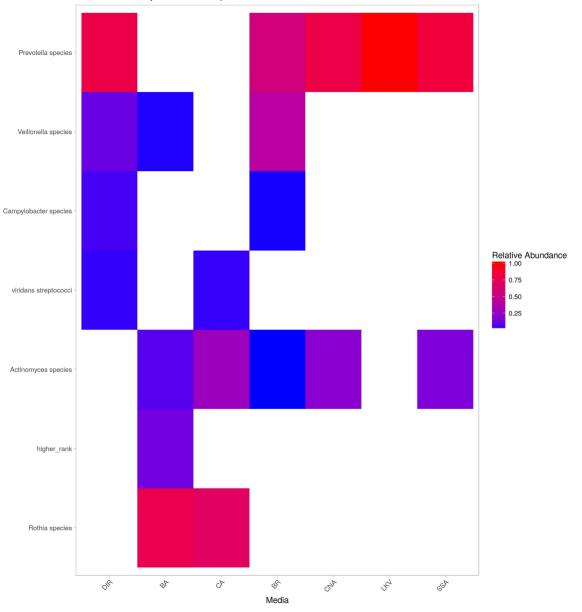


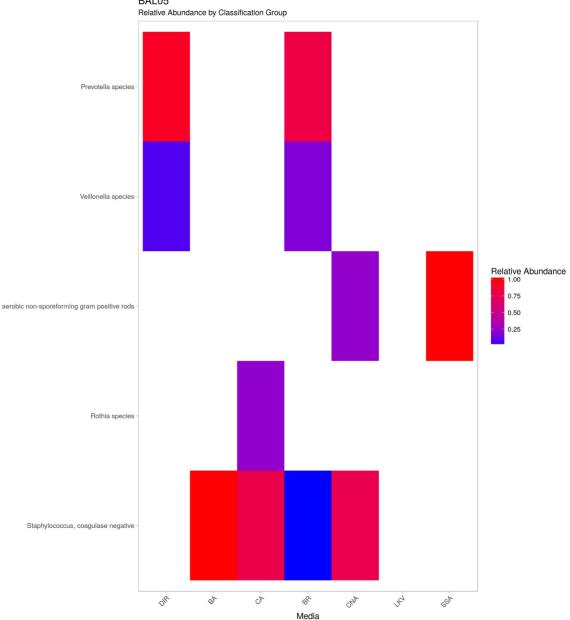


BAL03

BAL04

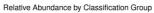
Relative Abundance by Classification Group

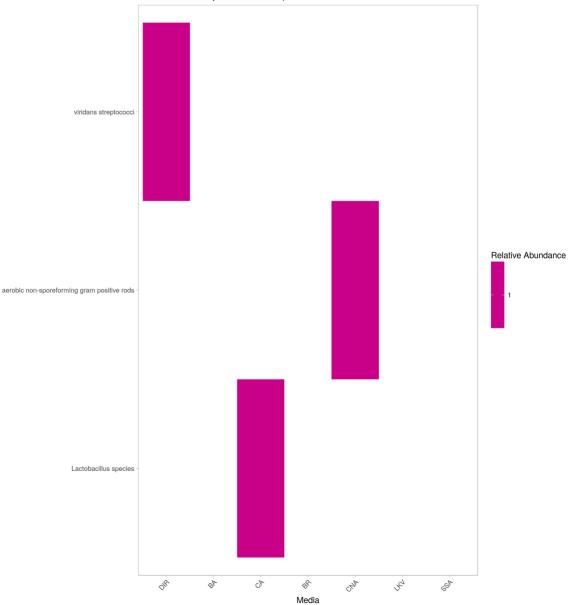


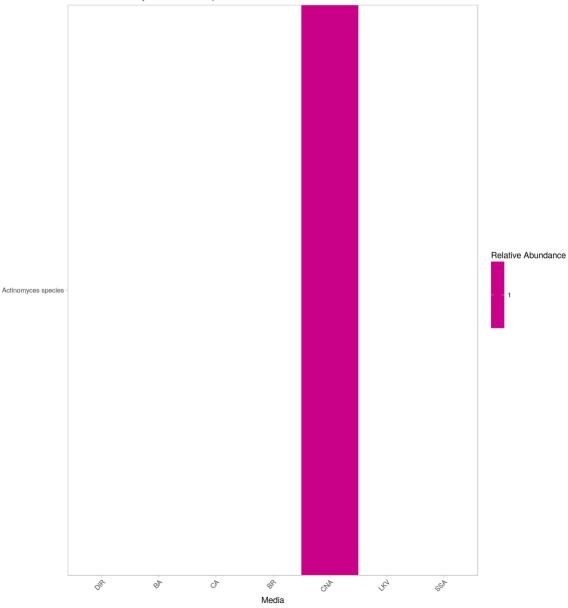


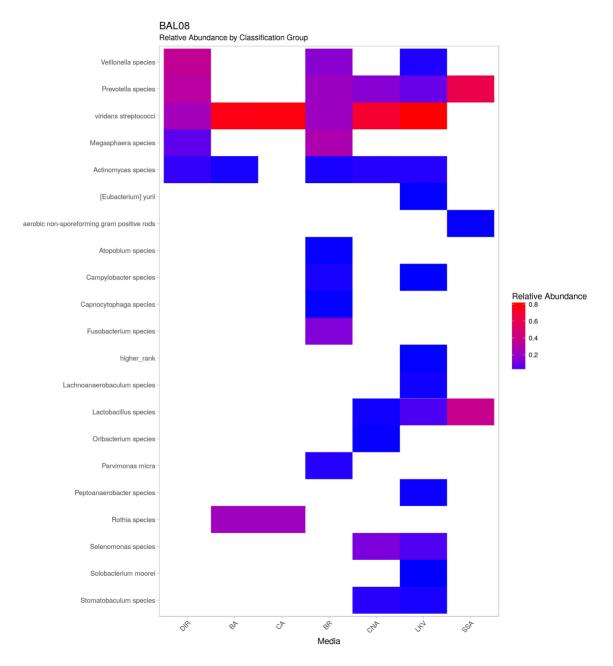
## BAL05

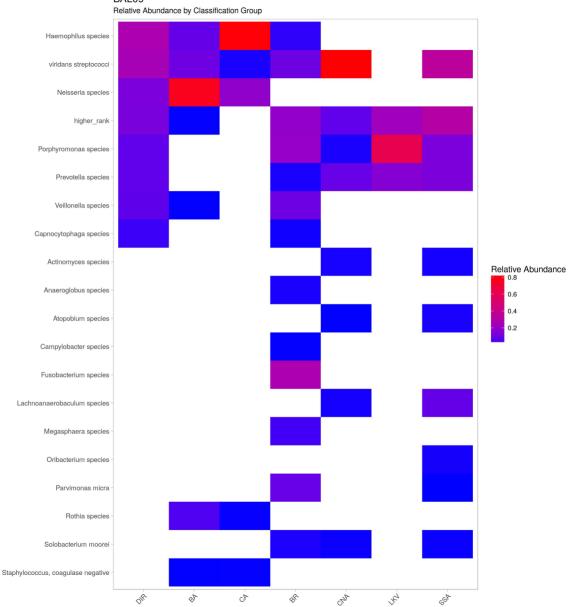










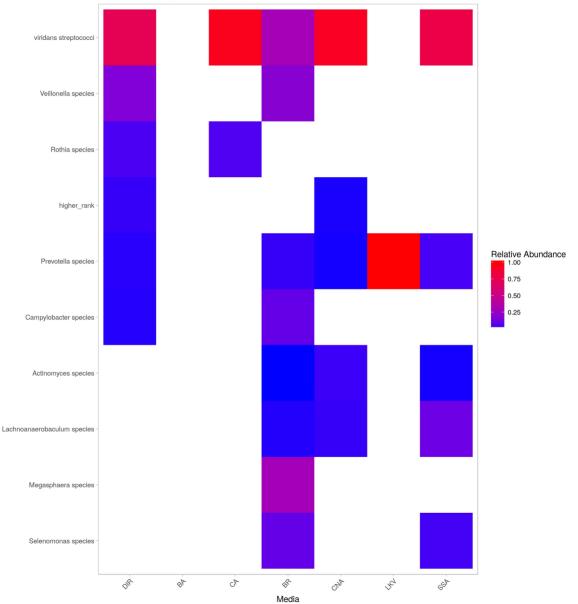


BAL09

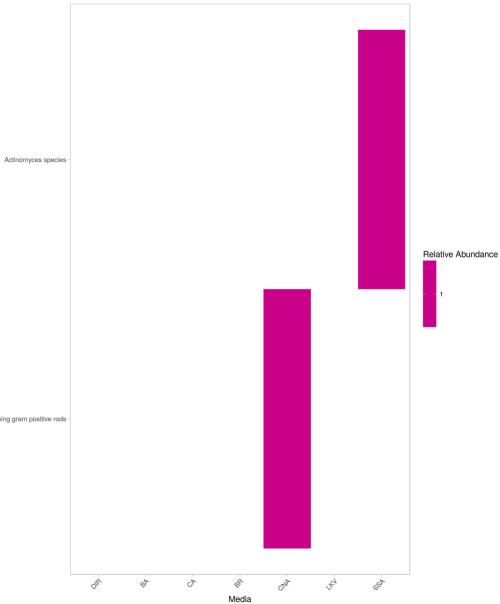
Media

BAL10

Relative Abundance by Classification Group

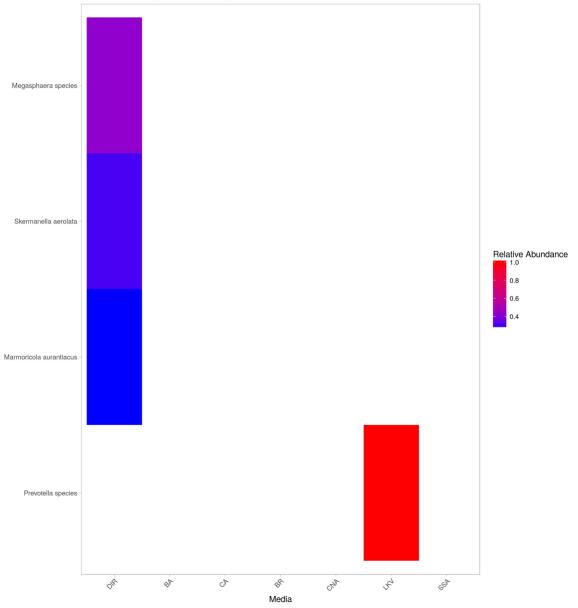


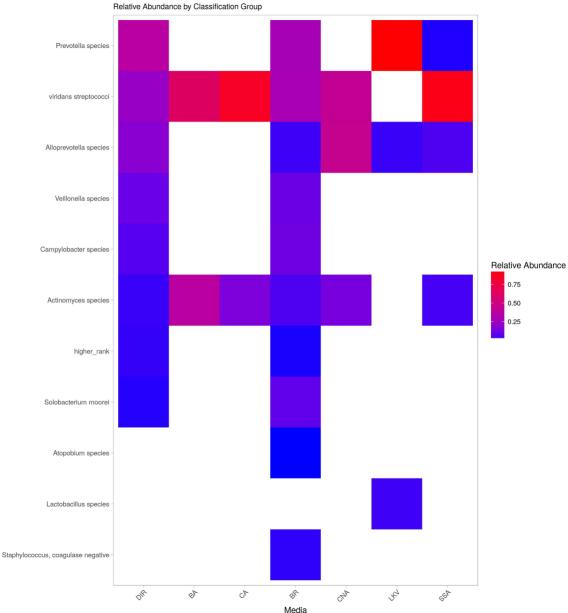
BAL11 Relative Abundance by Classification Group



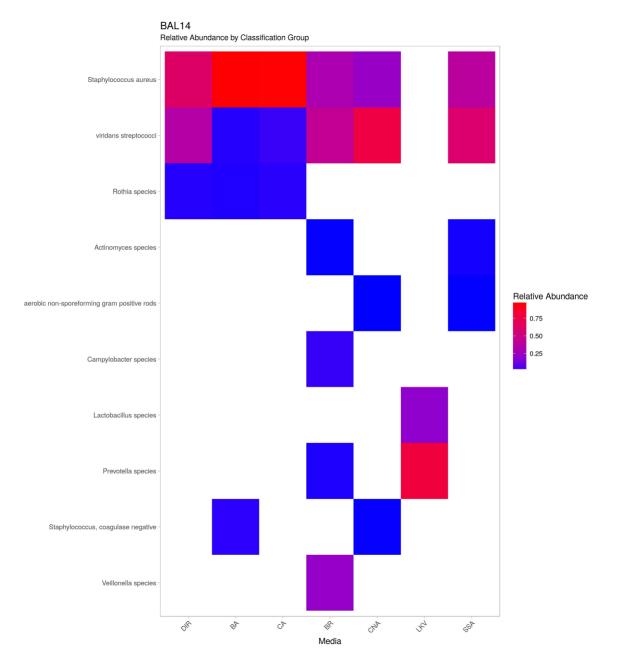
aerobic non-sporeforming gram positive rods

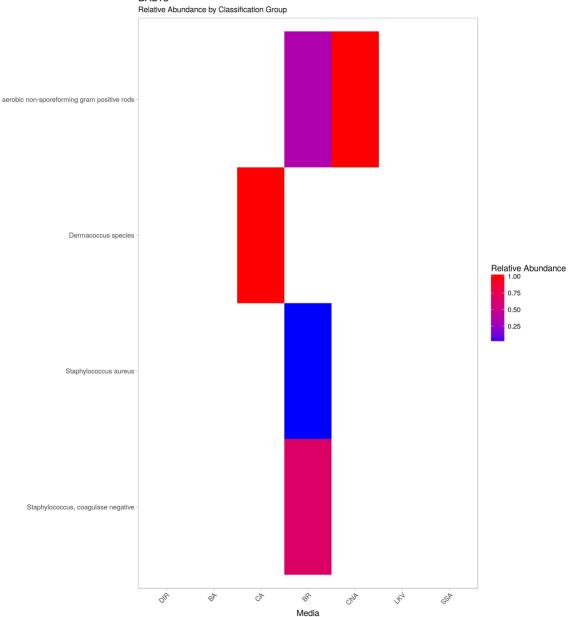
BAL12 Relative Abundance by Classification Group



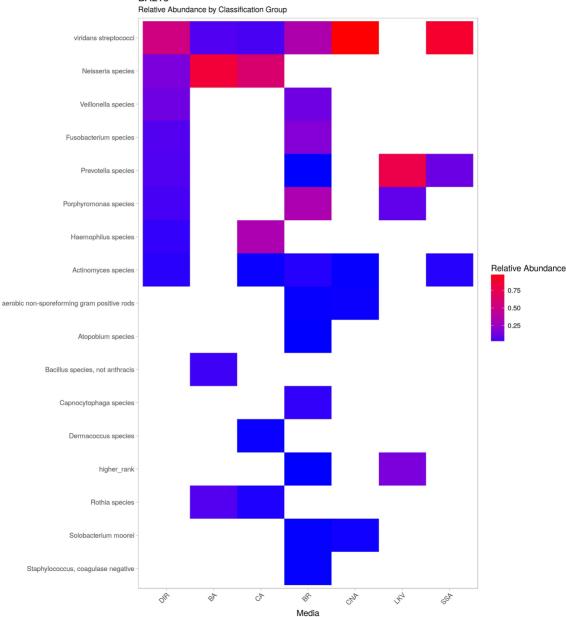


BAL13 Belative Abundance by Classification Grou

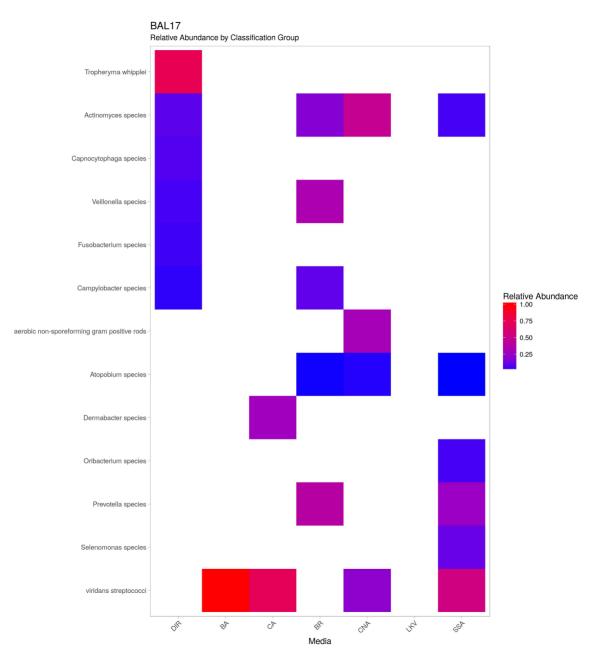




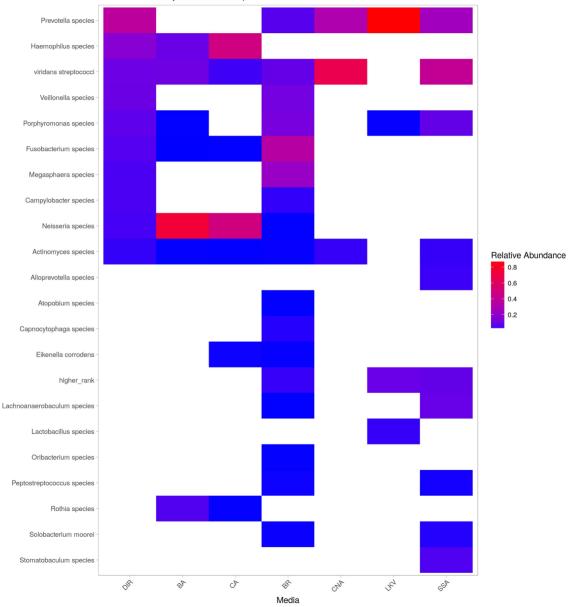
BAL15

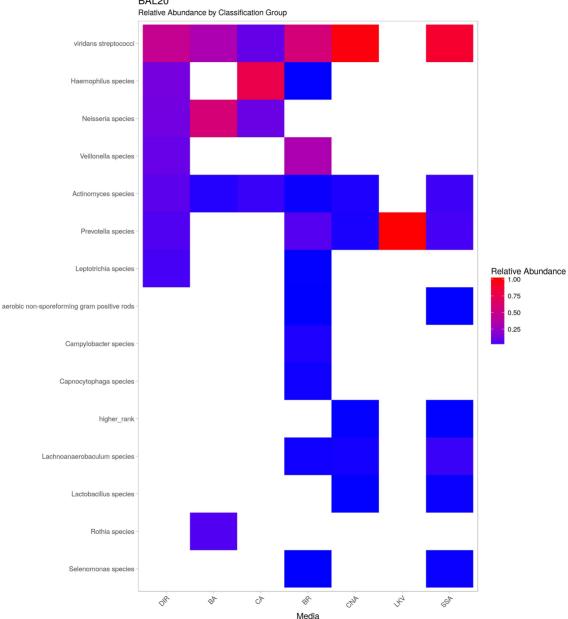


BAL16



BAL19 Relative Abundance by Classification Group





BAL20