Normalized (by genome size) average relative abundances: TCGA bacteria vs. fungi (Primary Tumor | RNA only) Relative Abundance - 0.0 **Domain** k\_\_Bacteria Normalized Mean F 0.0 - 0.2 - 0.1 - 0.0 k\_\_Eukaryota 0.083 0.065 0.058 0.046 0.049 0.057 0.038 0.035 0.034 0.028 0.034 0.033 0.029 0.009 0.008 900.0 0.007 0.003 0.002 0.04 0.003 TCGA-KICH TCGA-READ -MESO GBM KIRC SKCM TCGA Cancer Type