S5 Fig. Statistical significance of regressions of taxa agglomerated at a range of taxonomic levels, against each antibody found to relate to community structure in Tables 1 and S1. We report both A $p$-values and B False discovery rates for each. False discovery rates are calculated from $p$-values at each antibody-taxonomic level combination. Horizontal lines indicate significance thresholds of 0.2 (gray), 0.05 (blue) and 0.01 (green).