



Figure S4: Matrix showing the amount of sequences for all possible primer combinations. Combinations which were used for tagging samples are highlighted in green (with 6 samples marked with asterisks were excluded from the dataset, as they belong to another project). Other combinations with matching tags were highlighted in blue based on relative sequence abundance. Combinations in Orange and Red are highlighting mismatching tags likely due to sequencing errors or in the case of NA+NA PhiX reference sequences that were spiked into the Illumina run to increase sequence diversity.