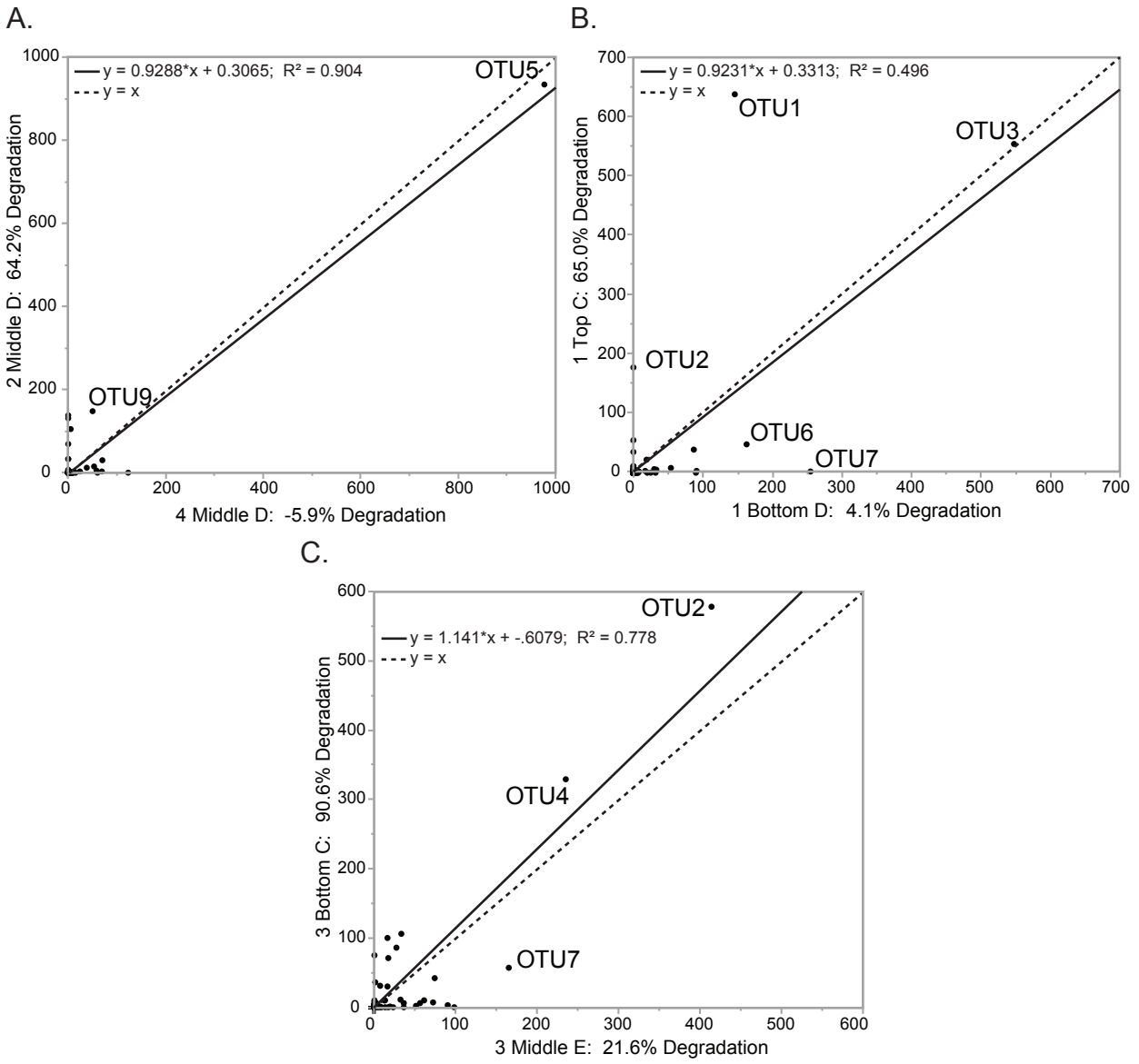
S4 Fig



S4 Fig. Pairwise comparisons of OTU abundance between samples with similar diversity. Samples were selected with high similarity in diversity but large differences in cellulolytic ability according to the cladogram in Fig 4. The abundance of reads for each OTU is plotted for the indicated samples. We show the best fit line for each plot to indicate the correlation between read abundances of each sample and the y = x line. Abundant OTUs are indicated. See Table 2 or S2 Table for taxonomic classification of each OTU.