Figure S2. Whole community diversity comparisons. Correspondence analysis calculated from the percent taxonomic distributions of the total OTUs. Dots are colored according to Ice Extent: <2007 in blue and >2007 in red. (A) Bacteria 97% phylum-level OTUs. (B) Eukarya 98% major-group-level OTUs. (C) Archaea 97% genus-level OTUs.