



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.