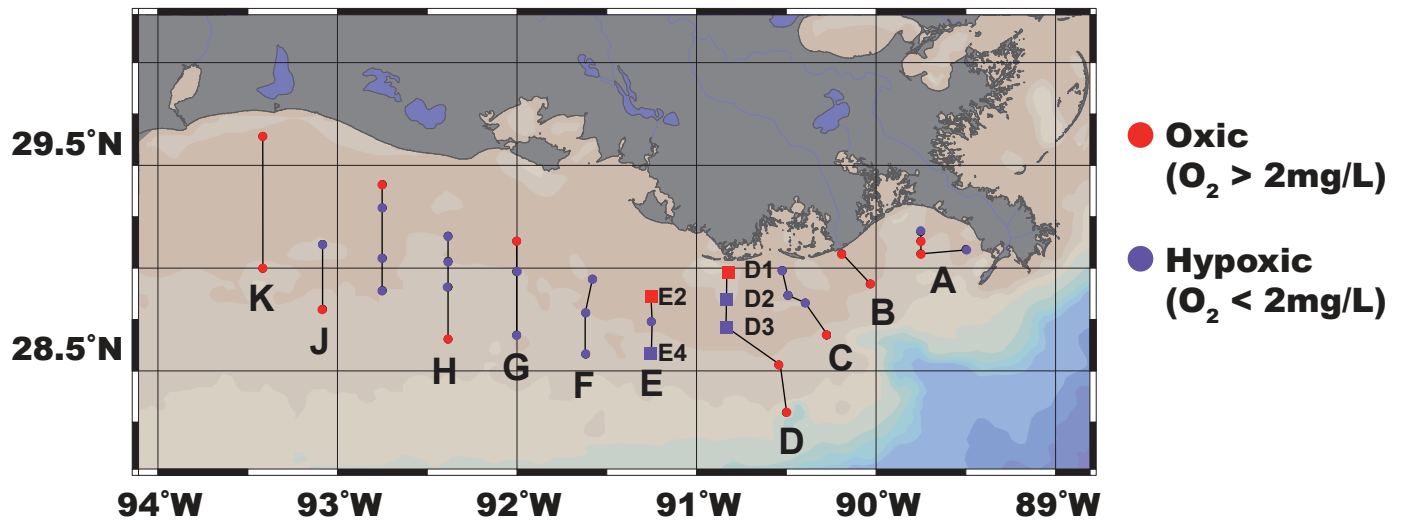




Supplemental Figure 1. Map from Gillies et al. (2015) of all samples collected in the 2013 nGOM dead zone. Samples selected for metagenomic and metatranscriptomic sequencing, O_D1, H_D2, H_D3, O_E2, and H_E4, are labelled.