

Figure SI1. Rank abundance curves for the 12 April 2010 DNA-based libraries (three replicate libraries from each treatment summed together) derived from field plots under the four field treatments (A=Ambient CO_2 , AF=Ambient CO_2 -N-fertilized, E=Elevated CO_2 , EF=Elevated CO_2 -N-fertilized; black bars). The summed abundance of OTUs from the three corresponding cDNA-based libraries from each of the field treatments are also plotted (red bars).