



Figure S5. Relative abundance (%) of all nitrifying groups obtained by *amoA* and *nxB* gene sequencing in plots with single and interactive effects of elevated temperature (eT) and atmospheric CO₂ (eCO₂) concentration – (A); and plots with single and interactive effects of future climate conditions (eT x eCO₂) and drought (D) – (B). The color code represents different genera/OTUs/clades/lineages. Multiple horizontal lines within the same color represent individual OTUs within a group. Taxa that comprised less than 0.1% of all reads per treatment are grouped as ‘Other’.