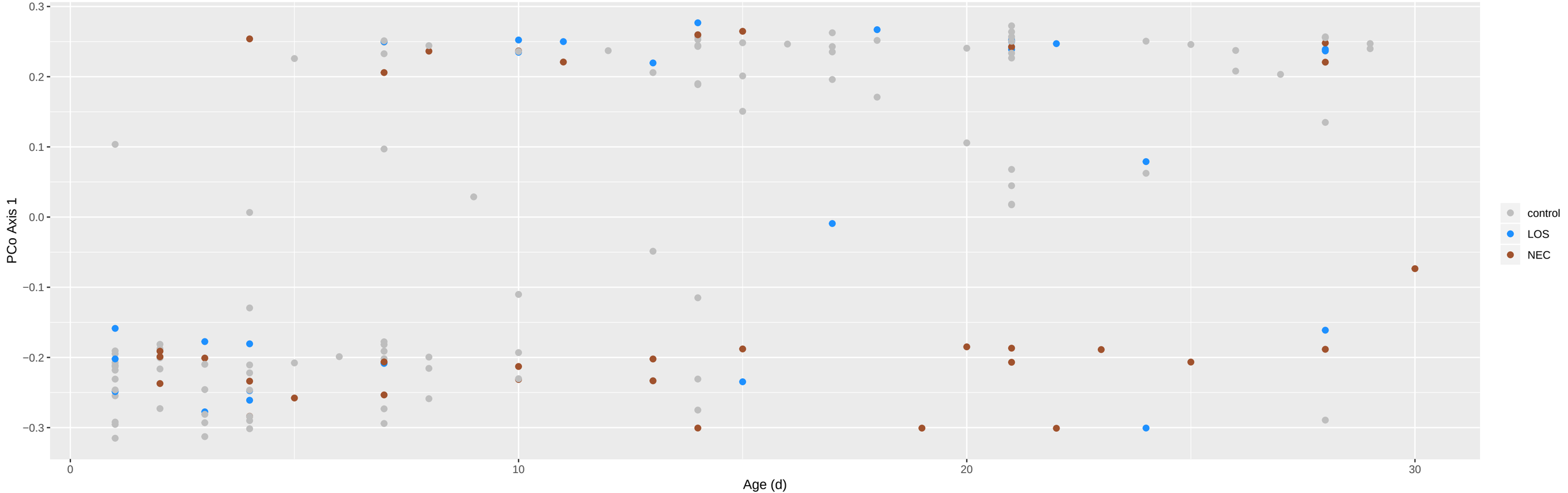PCoA based on weighted UniFrac distances between bacterial communities plotted over time, genus level