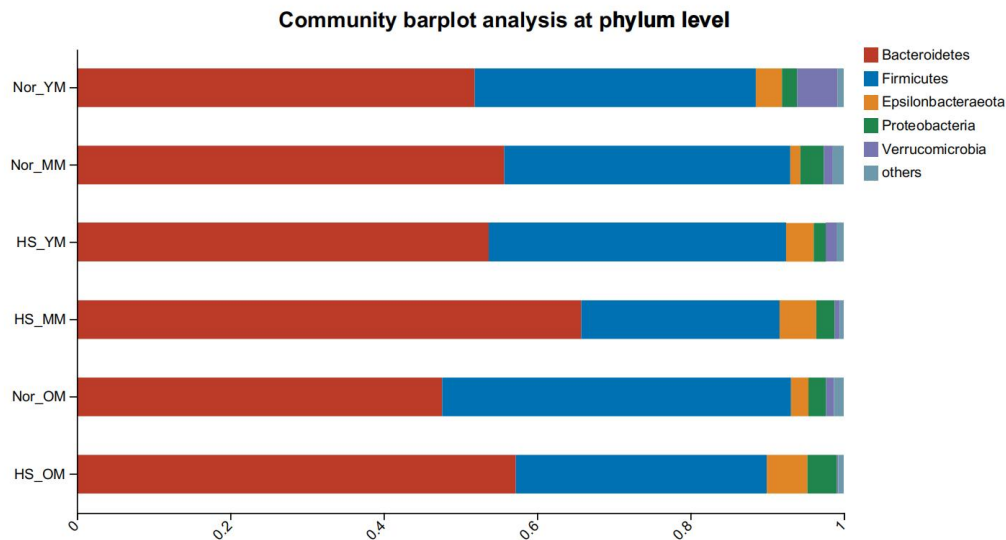
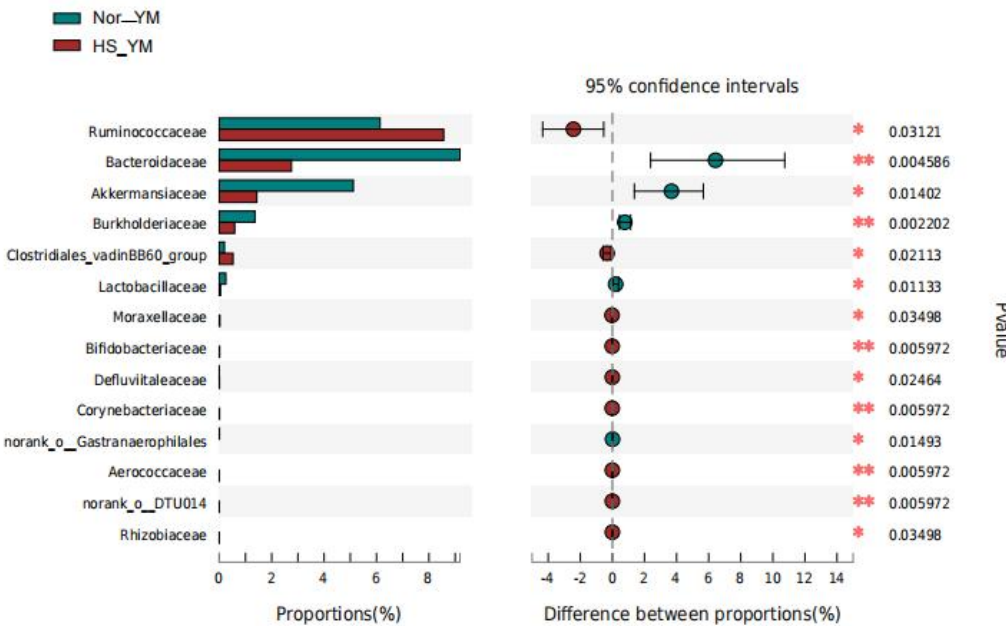
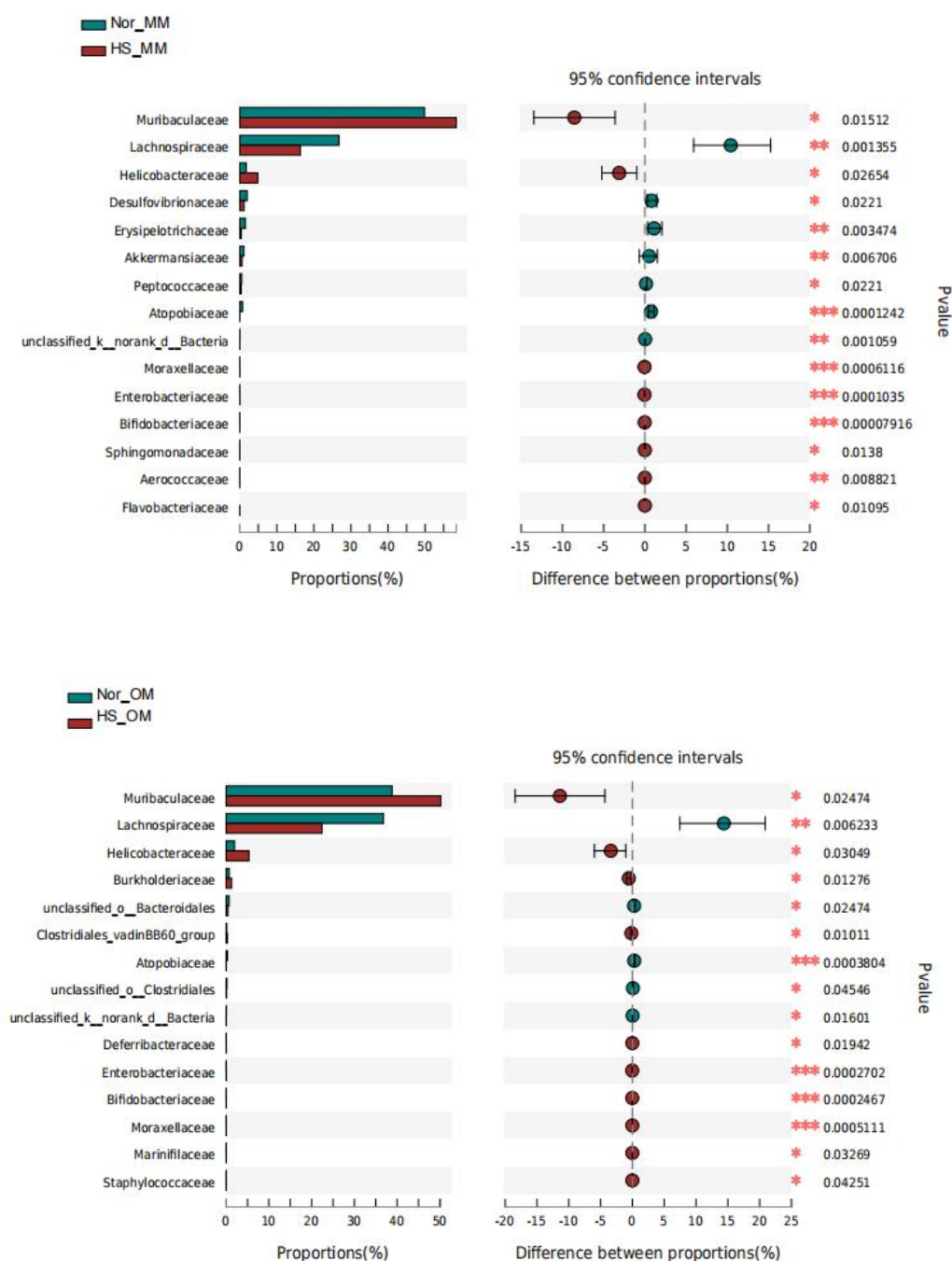


FigS1
A



B





FigS1. Long-time high sucrose intake effected the composition of gut microbiota. A: community barplot analysis at phylum level. B: different relative abundance of gut microbes between Nor and HS groups. YM: young mice group; MM: middle age mice group; OM: old age mice group.