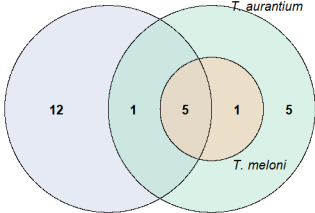
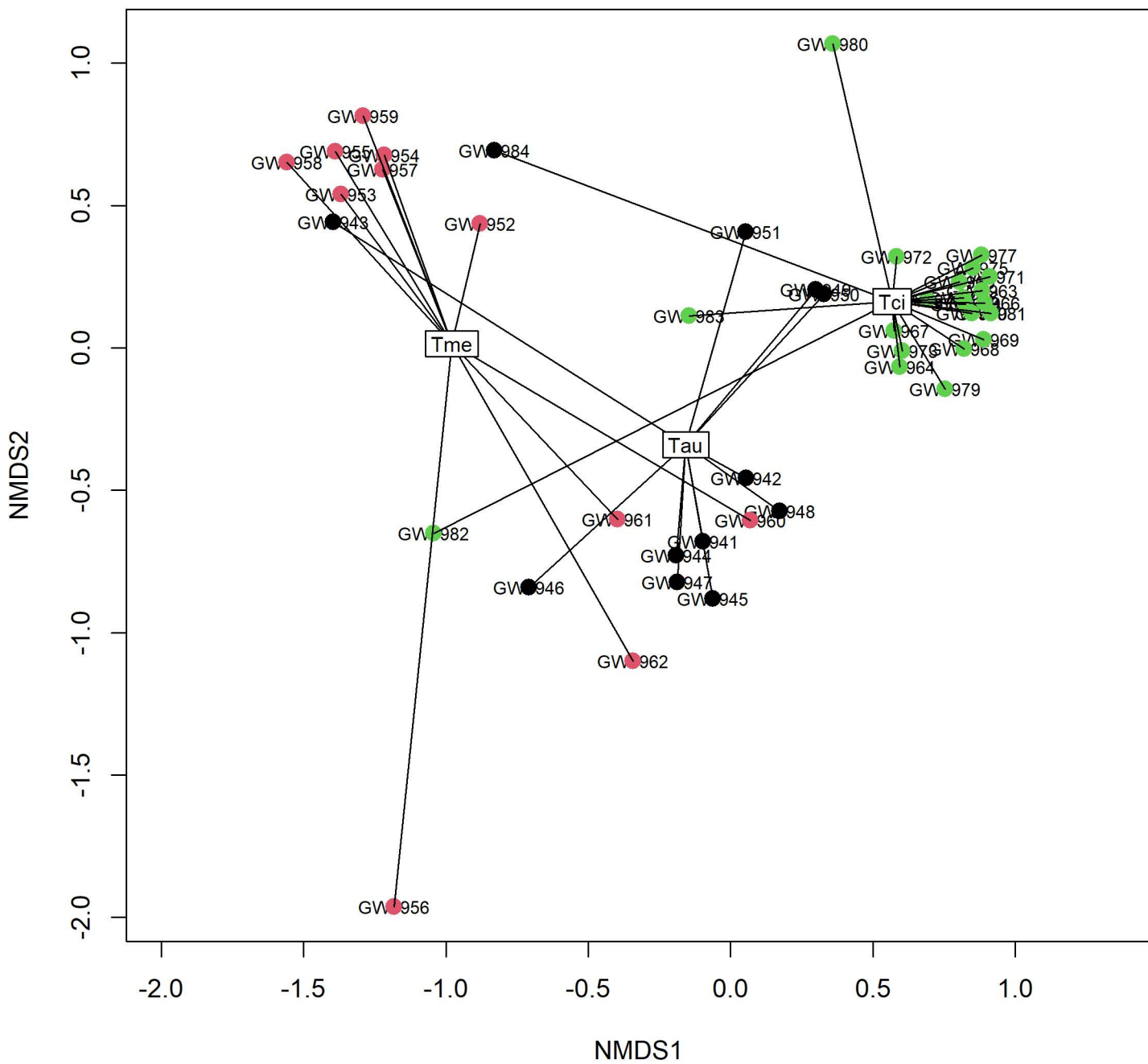


# Core bacterial communities across and between species

*T. citrina*

*T. aurantium*





NMDS2

NMDS1

