

Figure C1 – Phylum level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 %

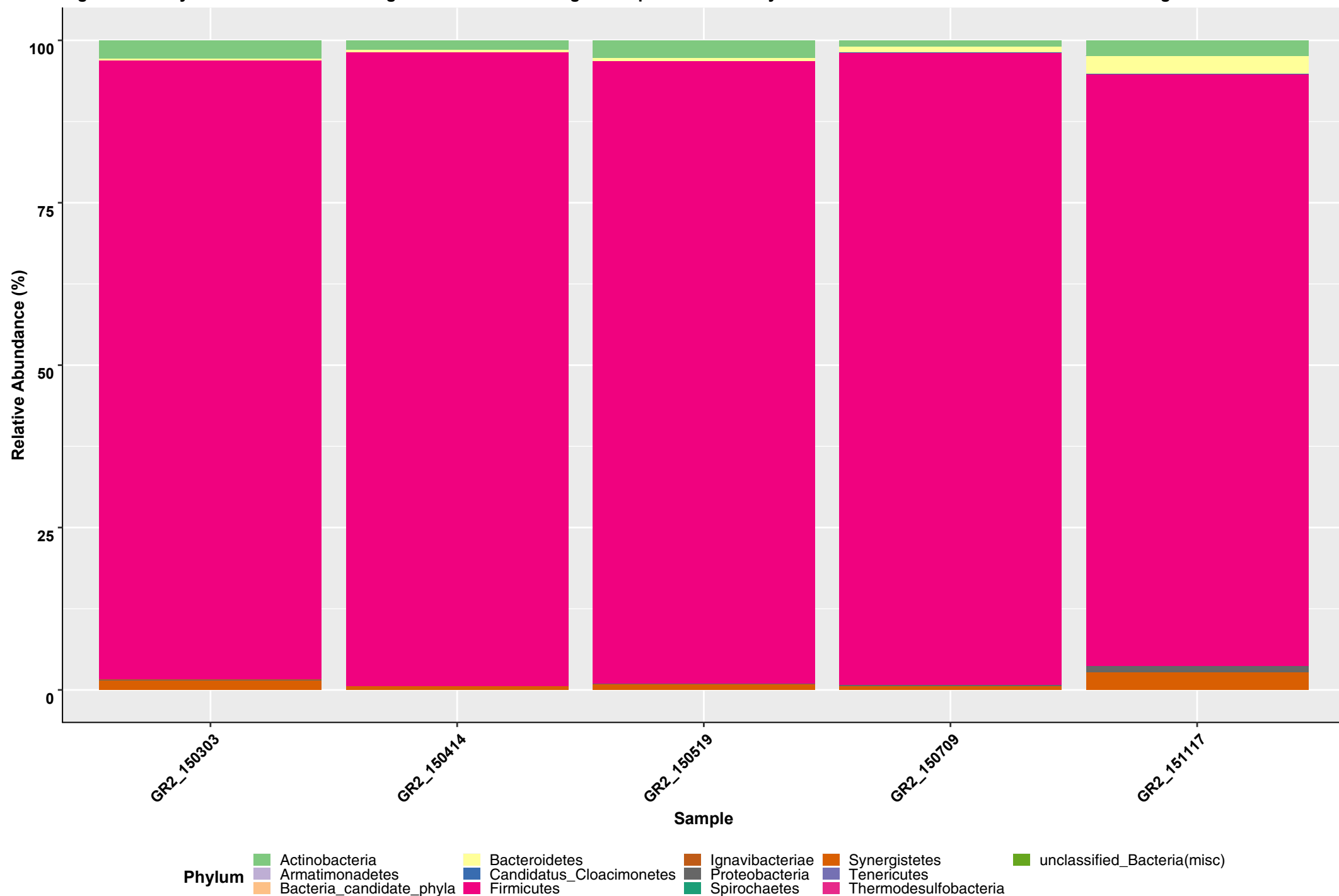


Figure C2 - Class level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 %(> 0.25 %)

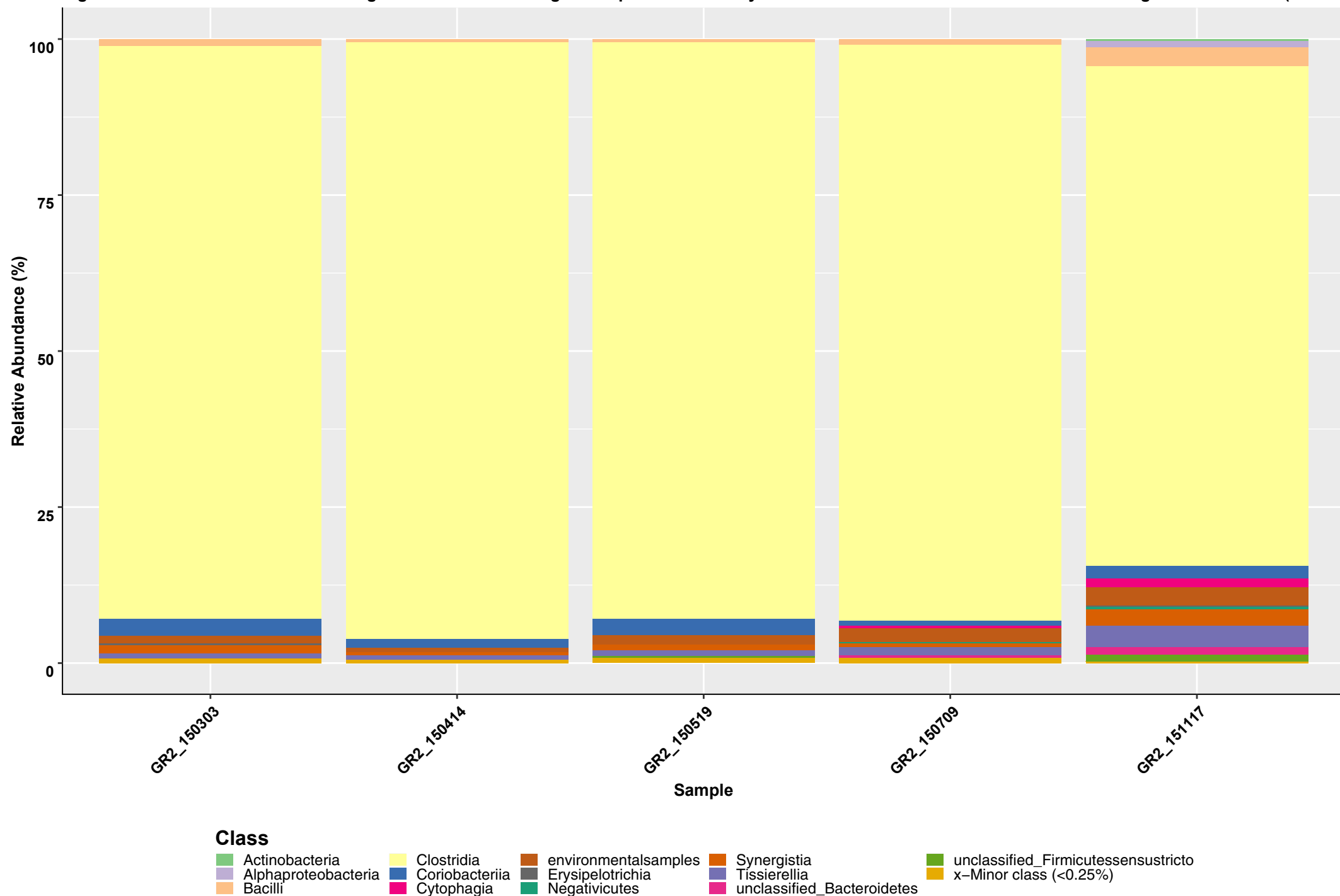


Figure C3 - Order level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 %(> 0.25 %)

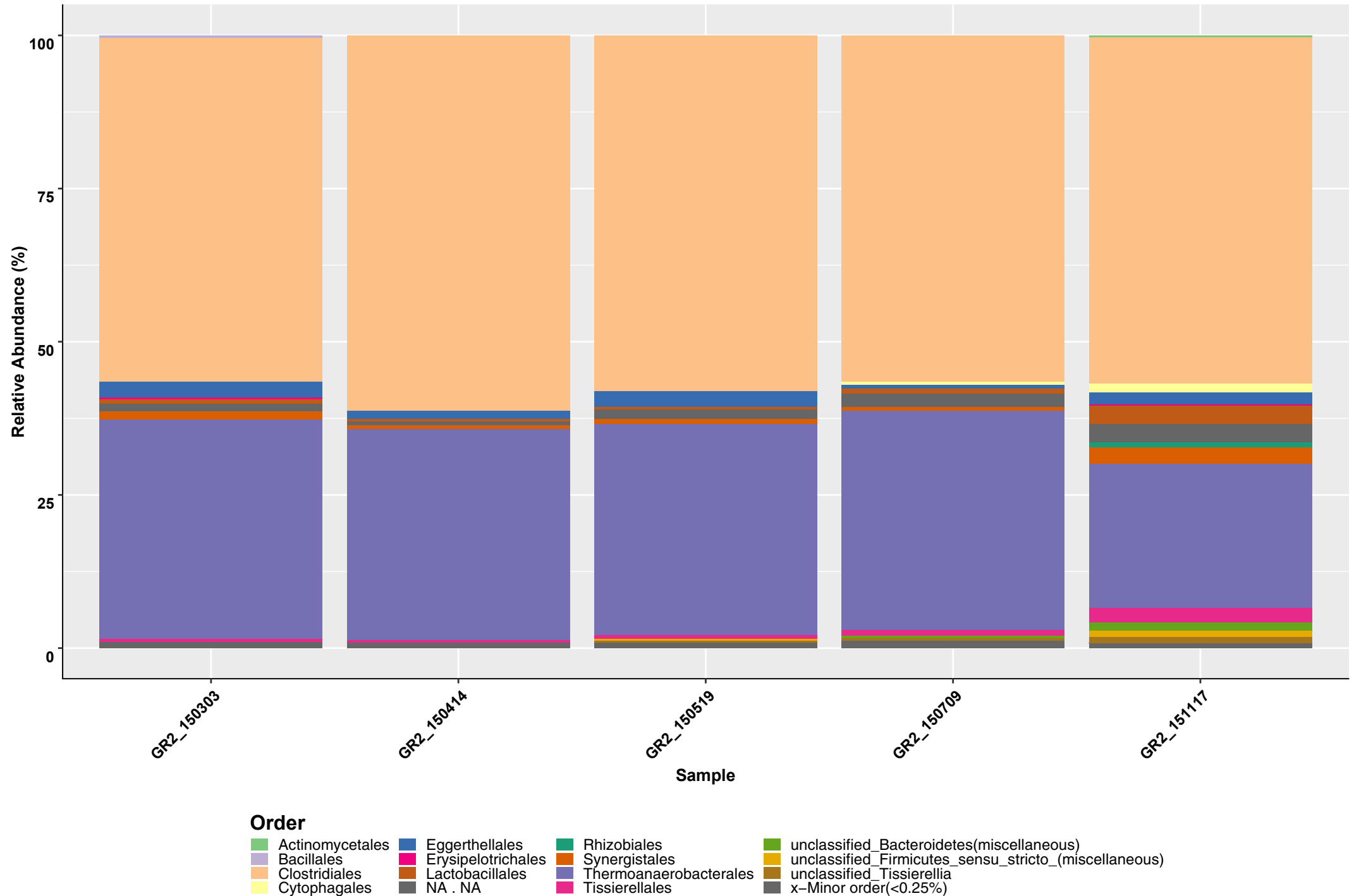


Figure C4 - Family level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 %(> 1 %)

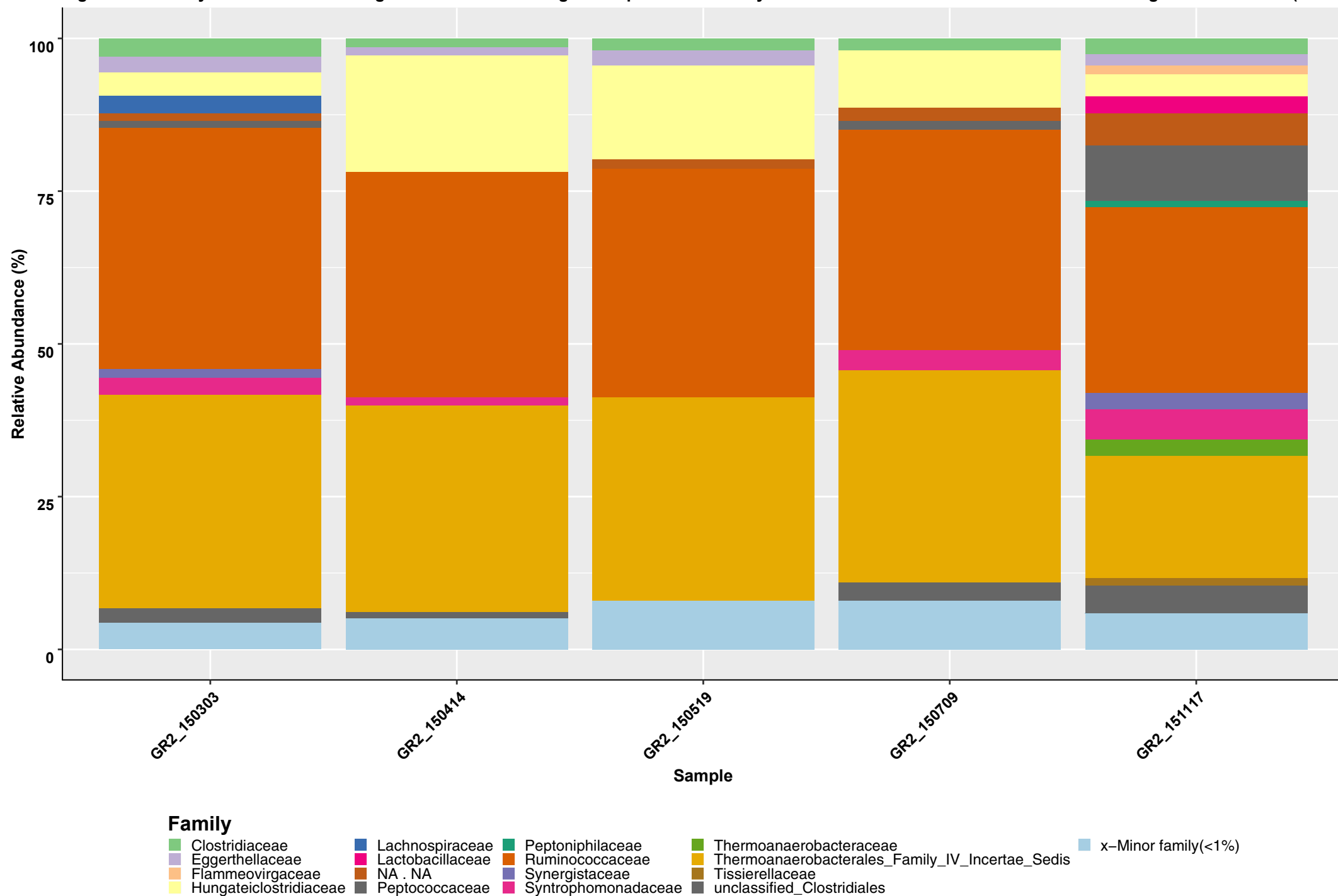


Figure C5 - Genus level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 % (> 1 %)

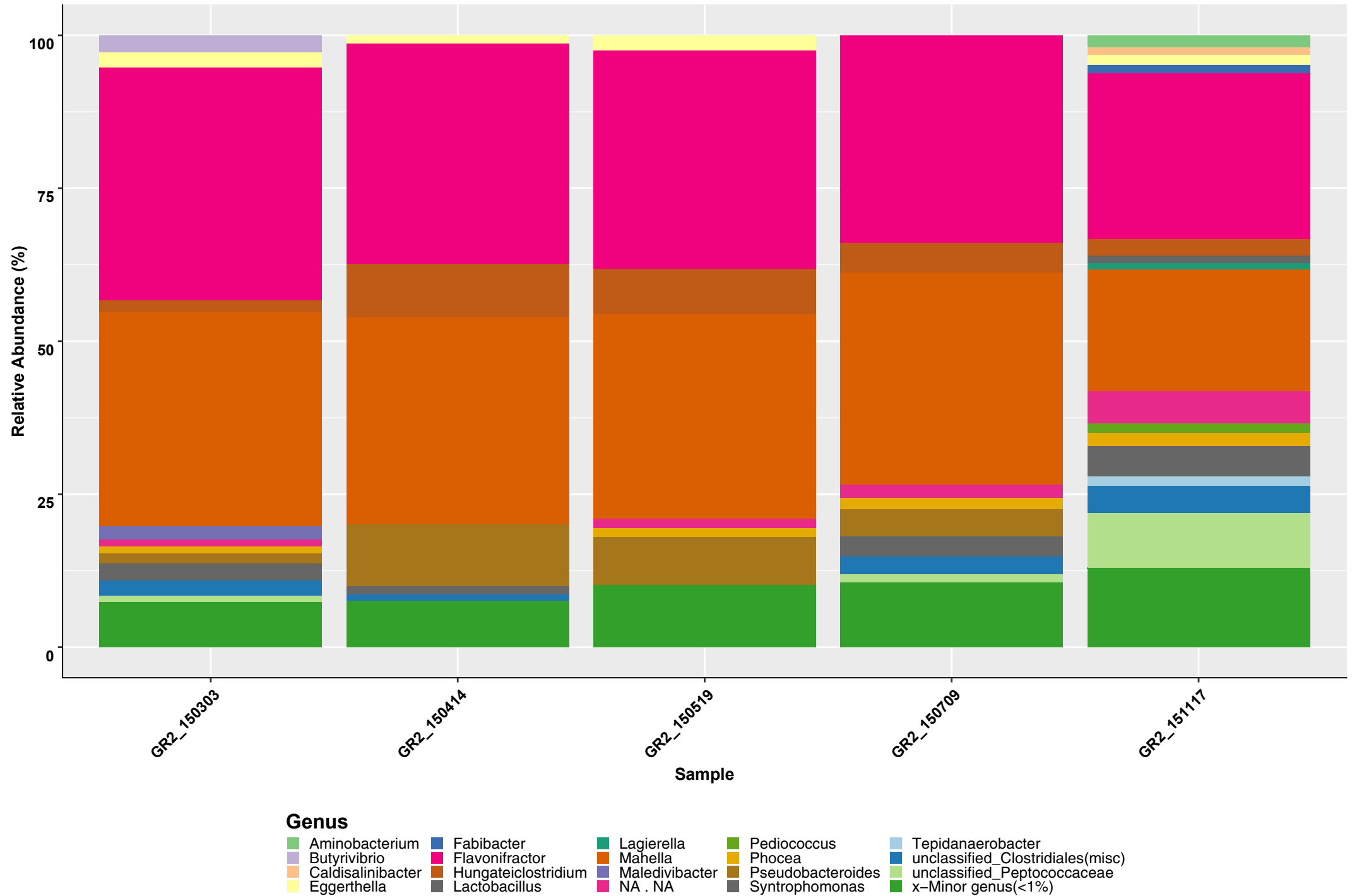


Figure C6 – Species level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 100 %(> 5 %)

