

Figure B1 – Phylum level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %

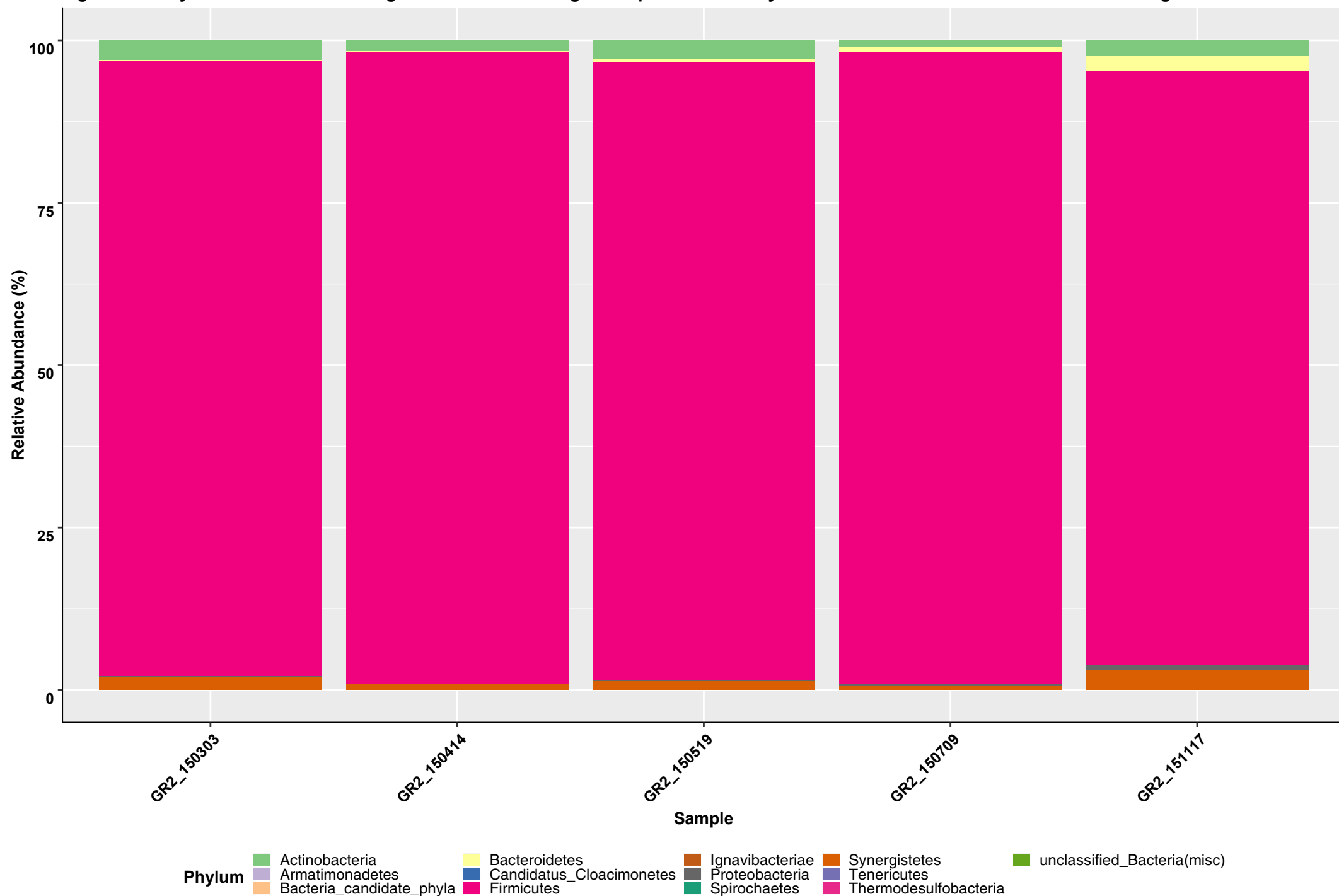


Figure B2 - Class level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %(> 0.25 %)

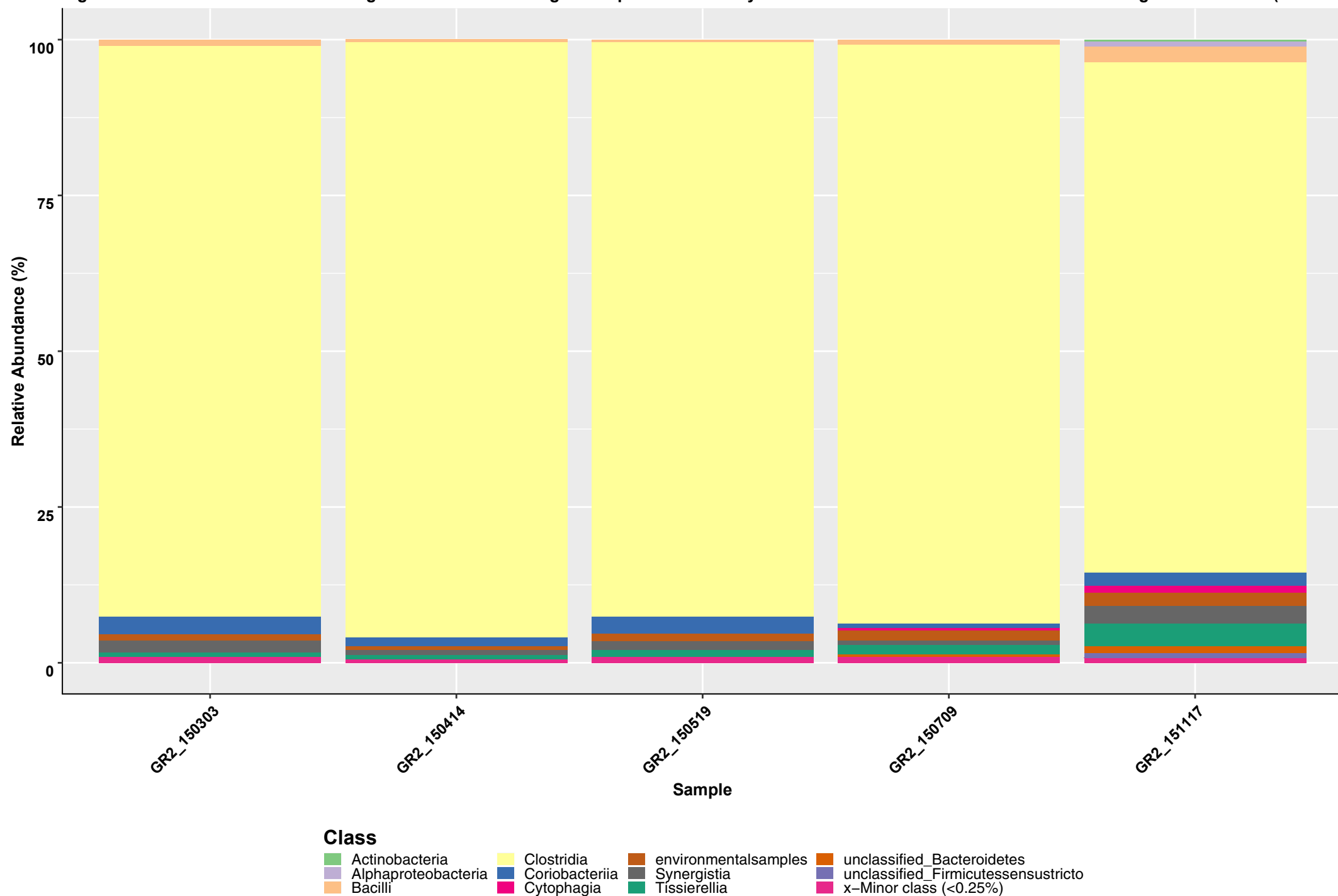


Figure B3 - Order level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %(> 0.25 %)

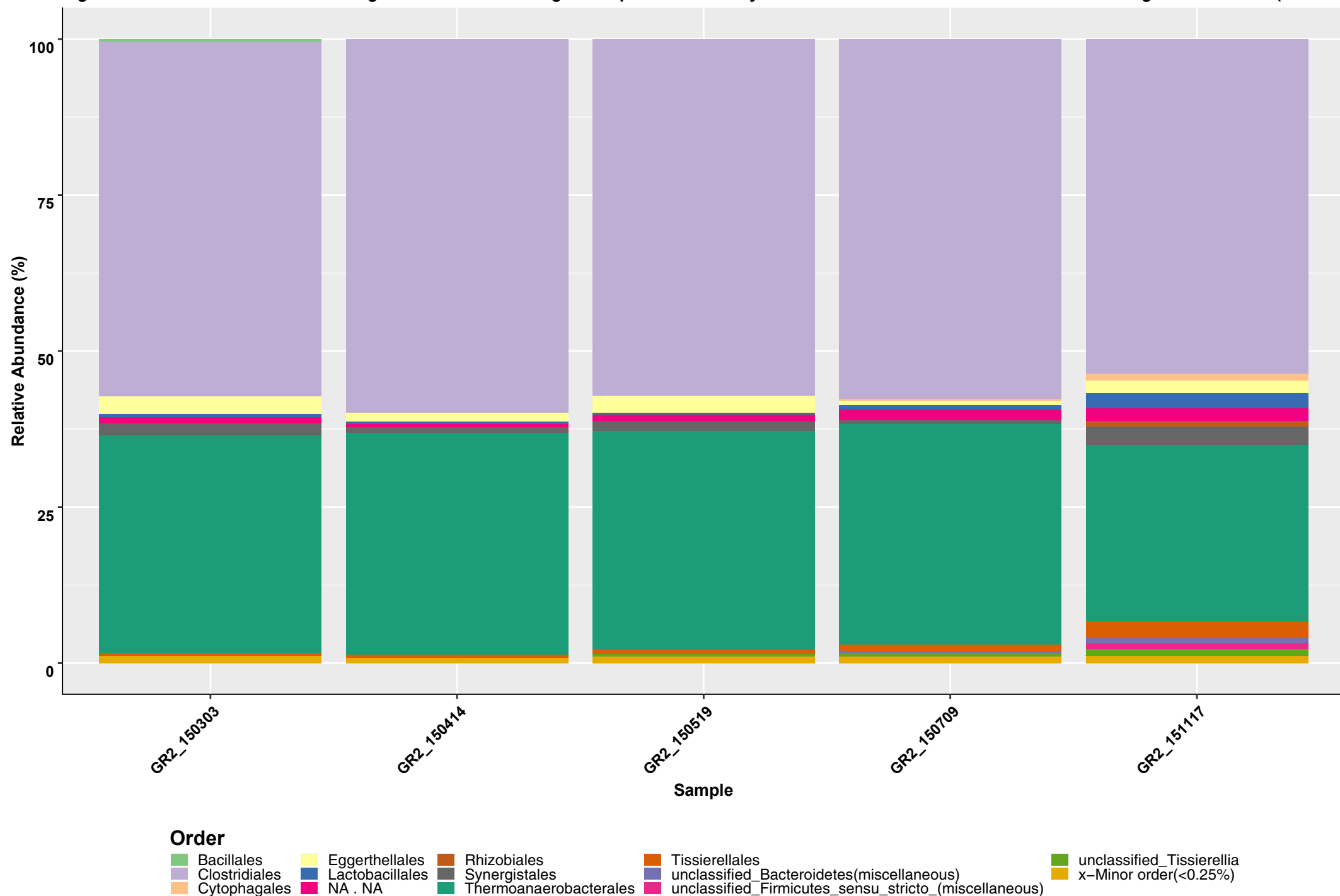


Figure B4 - Family level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %(> 1 %)

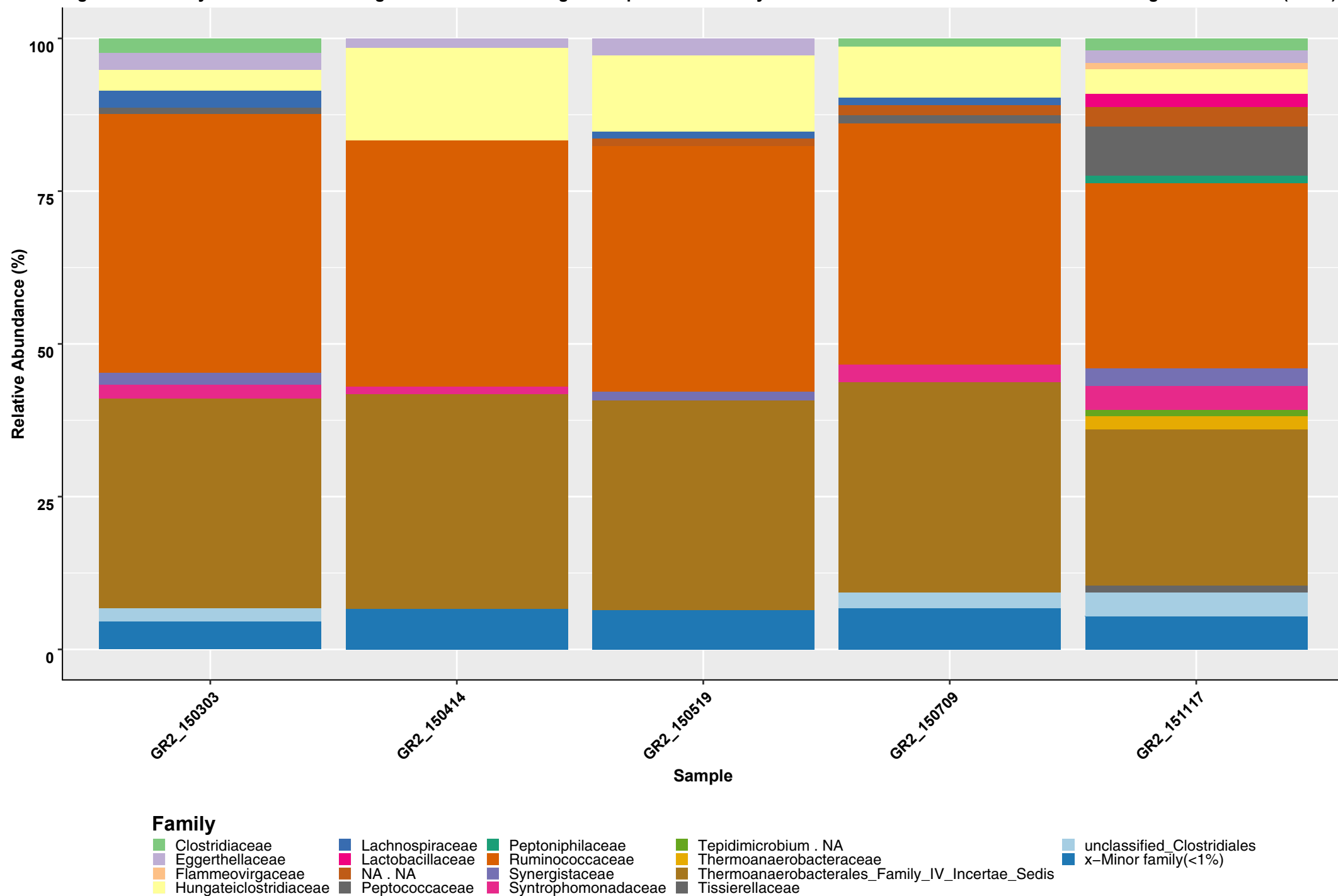


Figure B5 - Genus level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %(> 1 %)

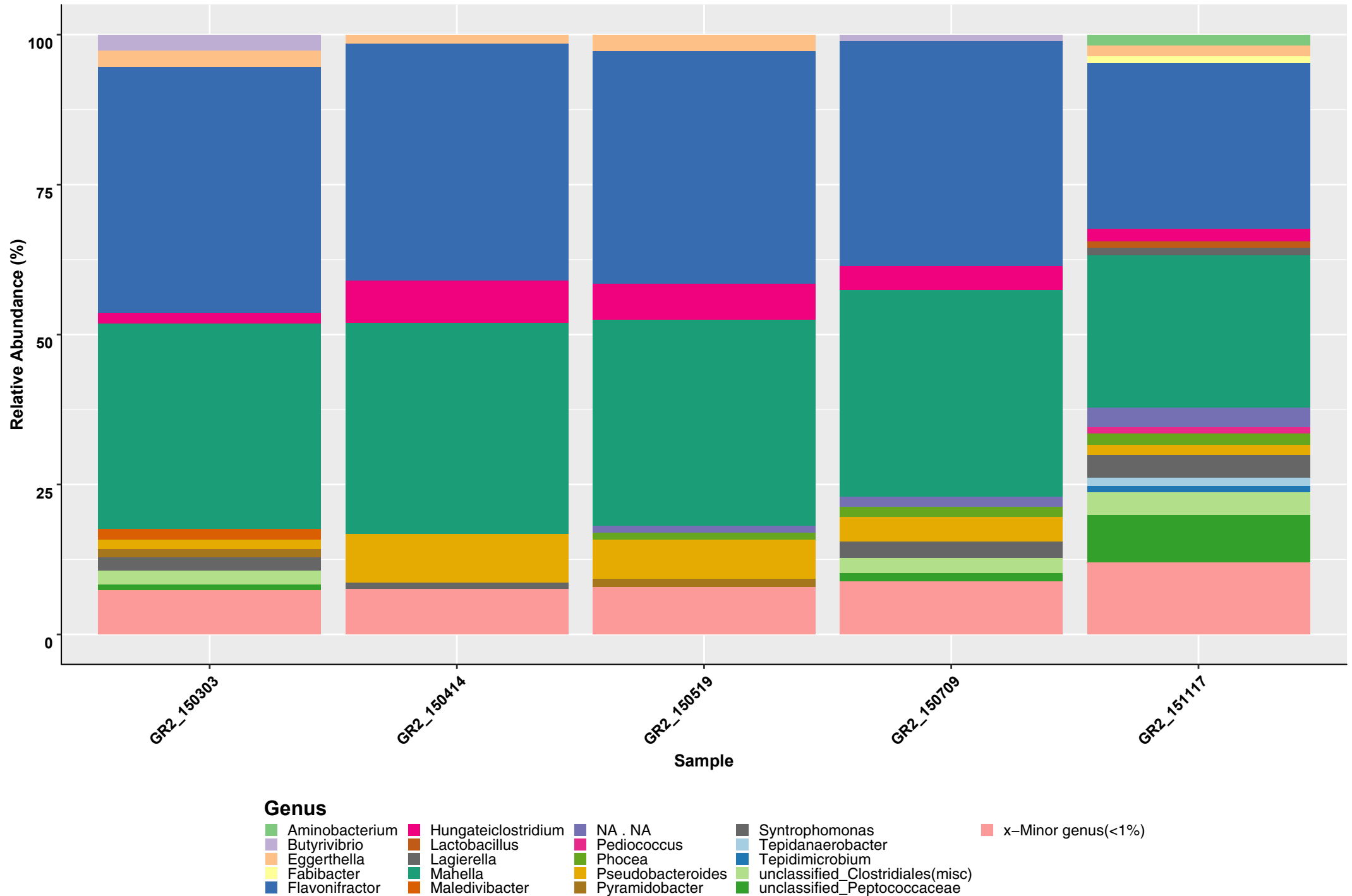


Figure B6 – Species level taxonomic assignments of the FTHFS gene sequence data analysis of forward reads with AcetoScan at clustering threshold 80 %(> 5 %)

