

Supplementary Figures: S1A-F

AcetoBase Version 2: A database update and re-analysis of formyltetrahydrofolate synthetase amplicon sequencing data from anaerobic digesters

Abhijeet Singh^{1*}, Anna Schnürer^{1*}

¹Anaerobic Microbiology and Biotechnology Group, Department of Molecular Sciences, Swedish University of Agricultural Sciences, Almas Allé 5, Uppsala, SE-750 07, Uppsala, Sweden

*For correspondence. E-mail: anna.schnurer@slu.se; abhijeet.singh@slu.se, Tel. +46 18671000; Fax +46 18672000; Address: Department of Molecular Sciences, Box 7025, 75007 Uppsala, Sweden

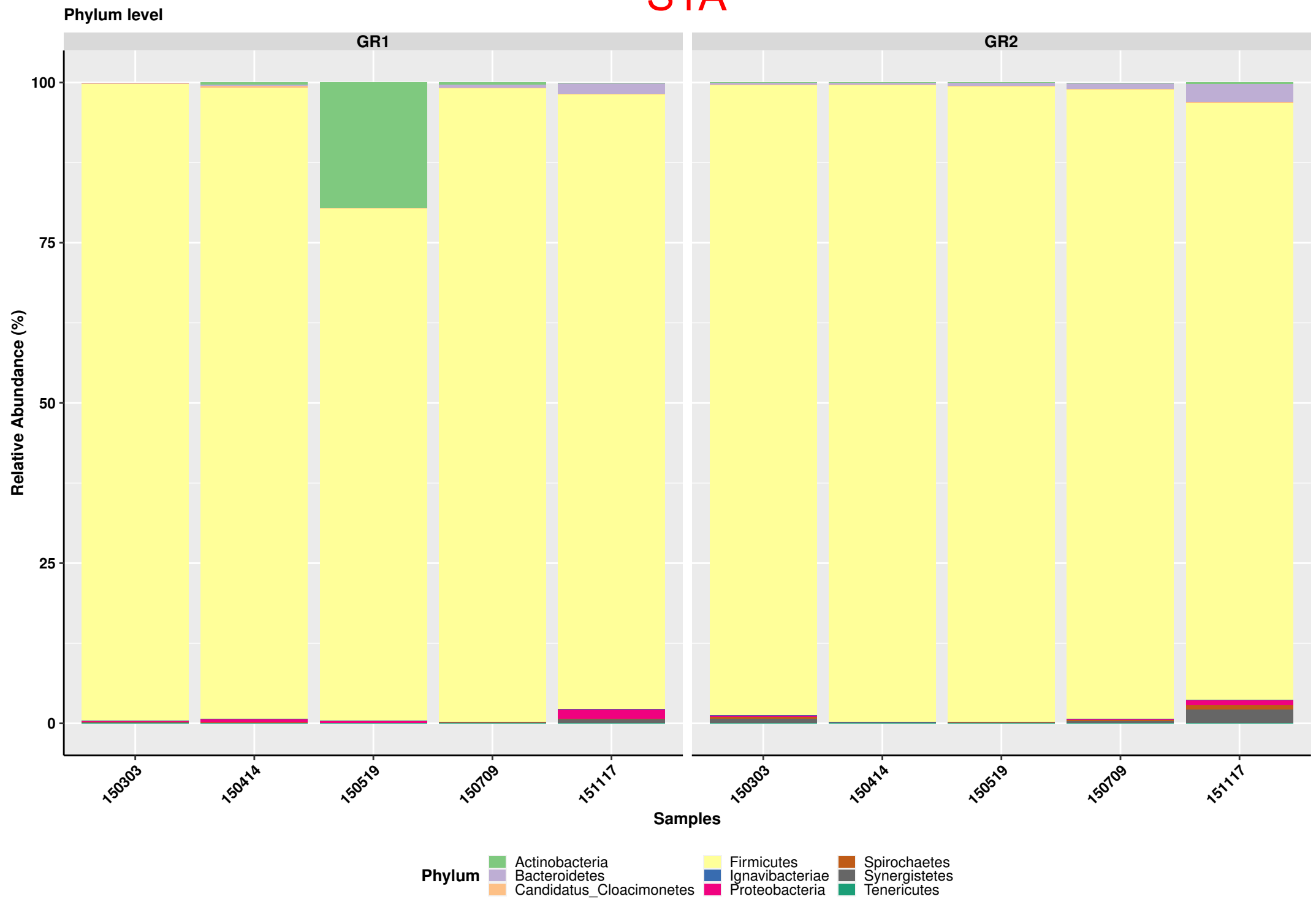
Description:

The supplementary figures S1A-F (phylum-species level) are generated in present study by the reanalysis of the FTHFS amplicon sequence data (reactor GR2) from the study by Singh *et al.* (2020) and additional data mentioned in the present study (reactor GR1).

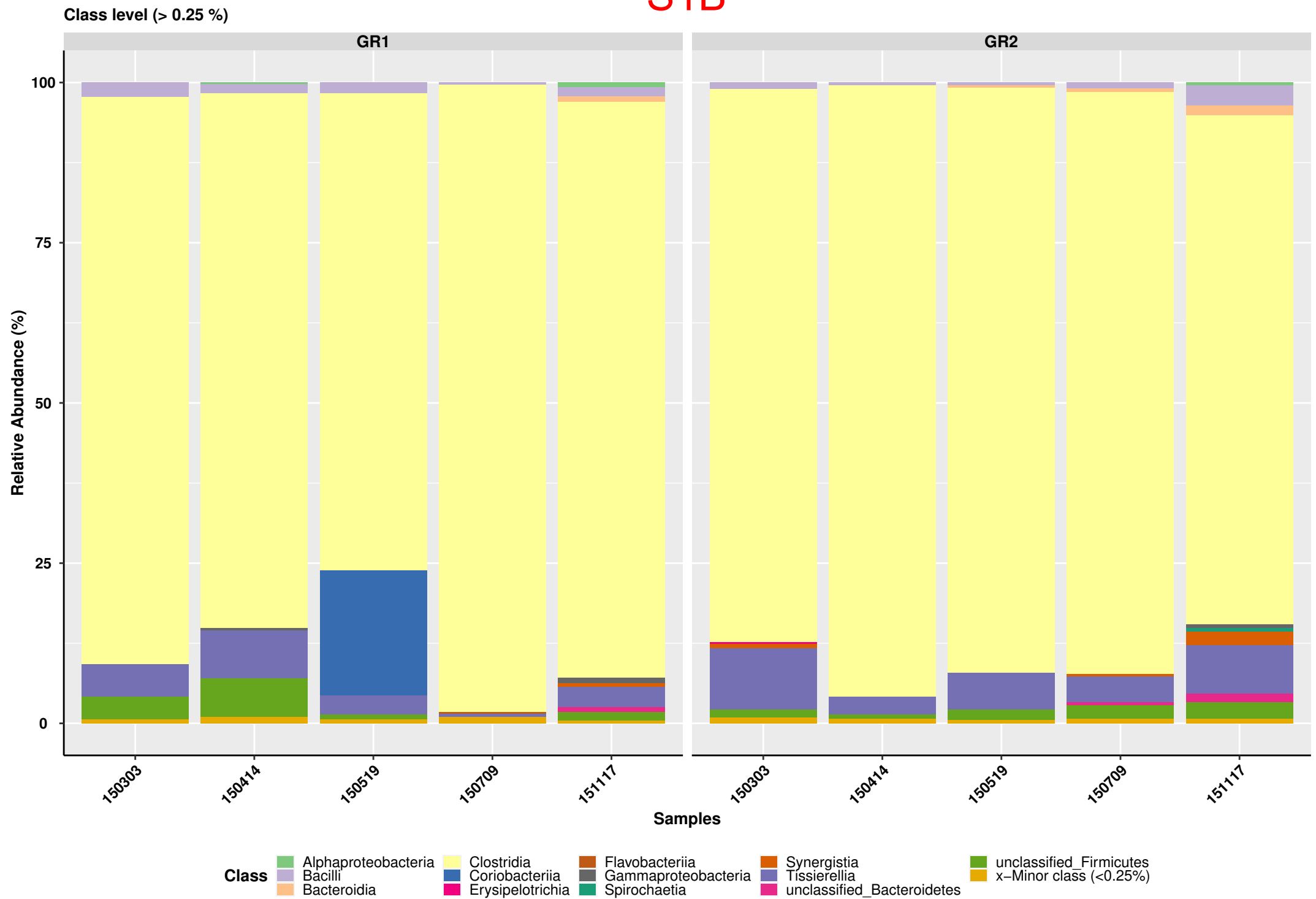
Reference:

Singh, A., Nylander, J. A. A., Schnürer, A., Bongcam-Rudloff, E., and Müller, B. (2020). High-Throughput Sequencing and Unsupervised Analysis of Formyltetrahydrofolate Synthetase (FTHFS) Gene Amplicons to Estimate Acetogenic Community Structure. *Front. Microbiol.* 11, 1–13. doi:10.3389/fmicb.2020.02066.

S1A

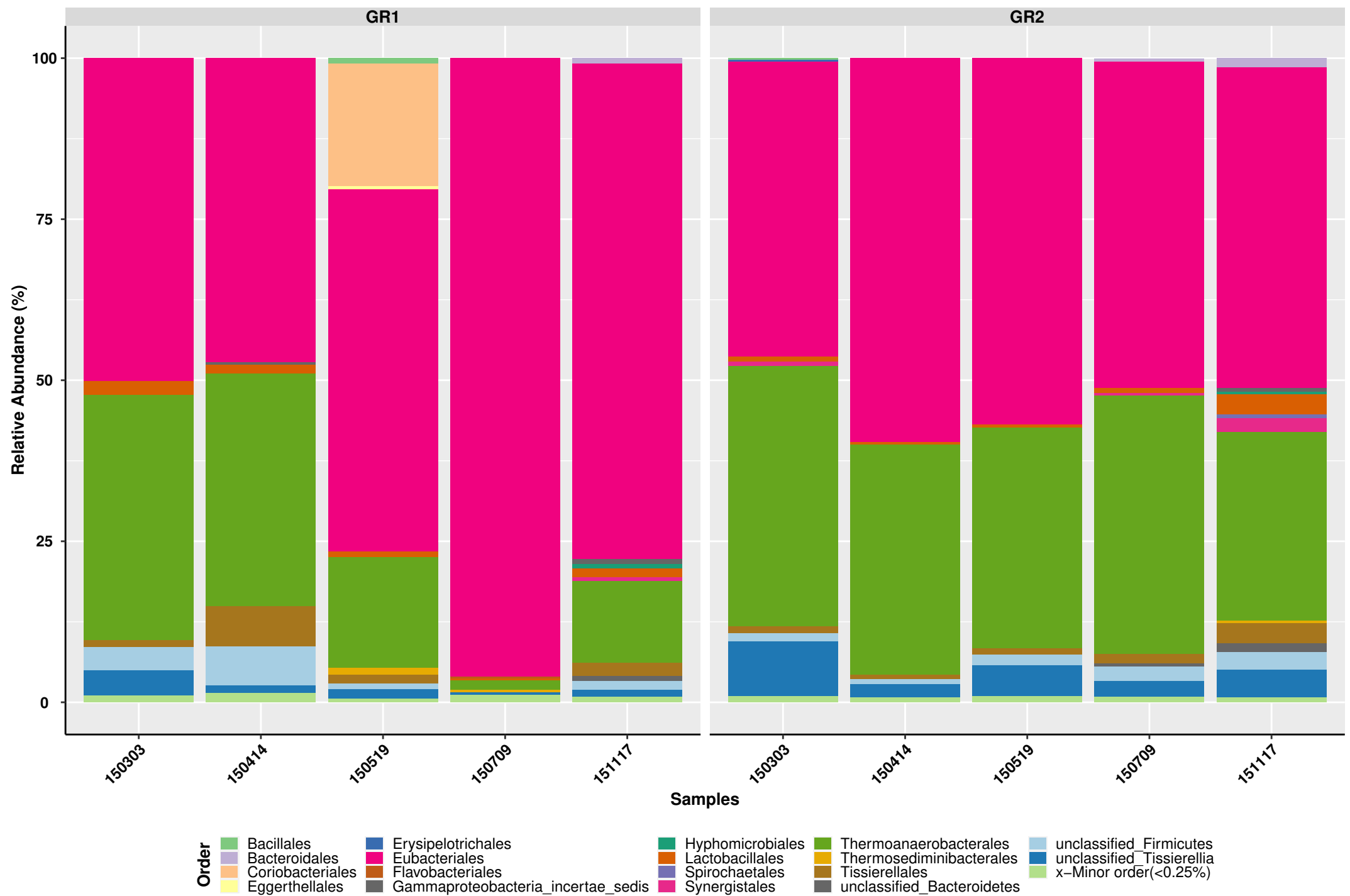


S1B

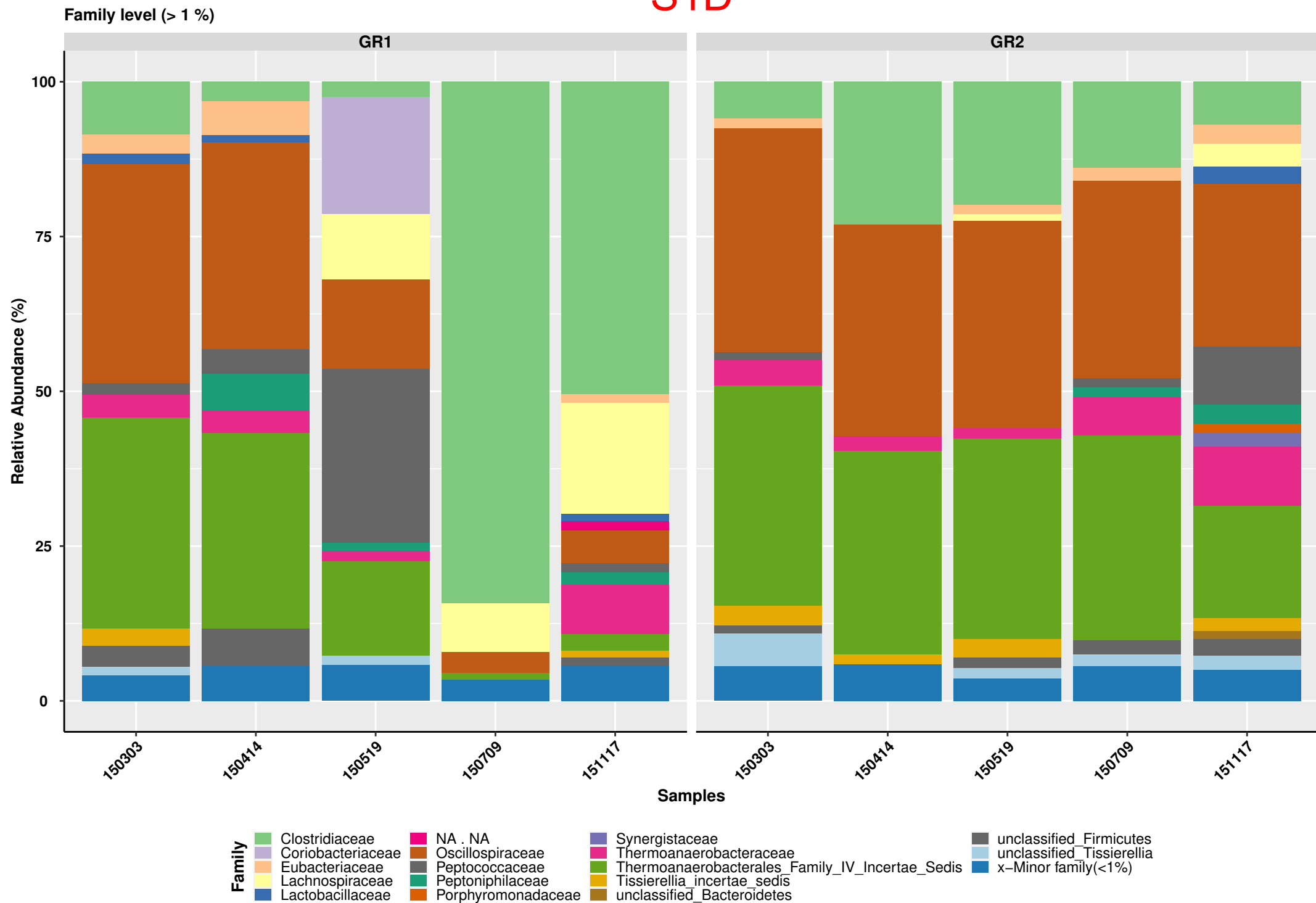


S1C

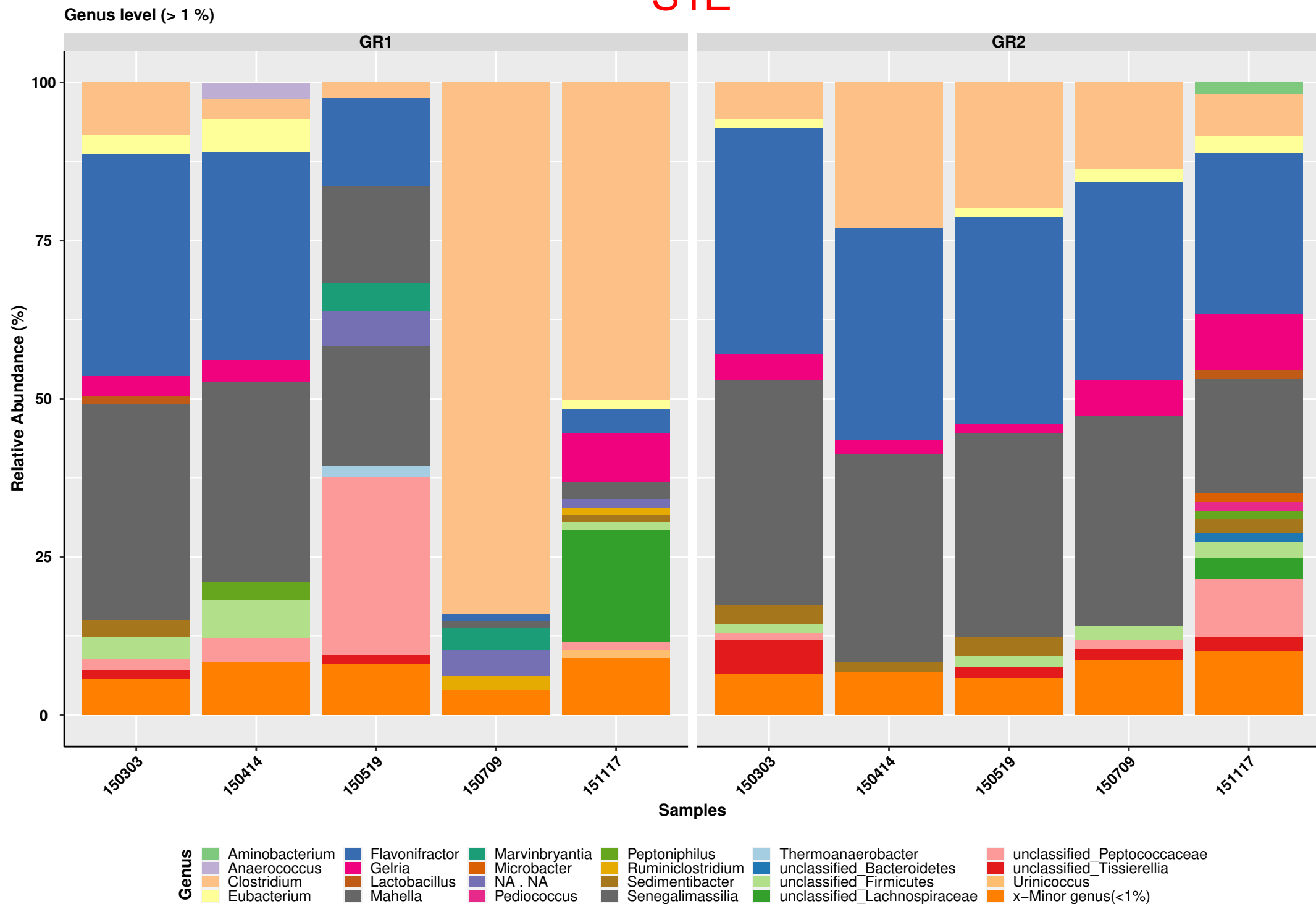
Order level (> 0.25 %)



S1D



S1E



S1F

