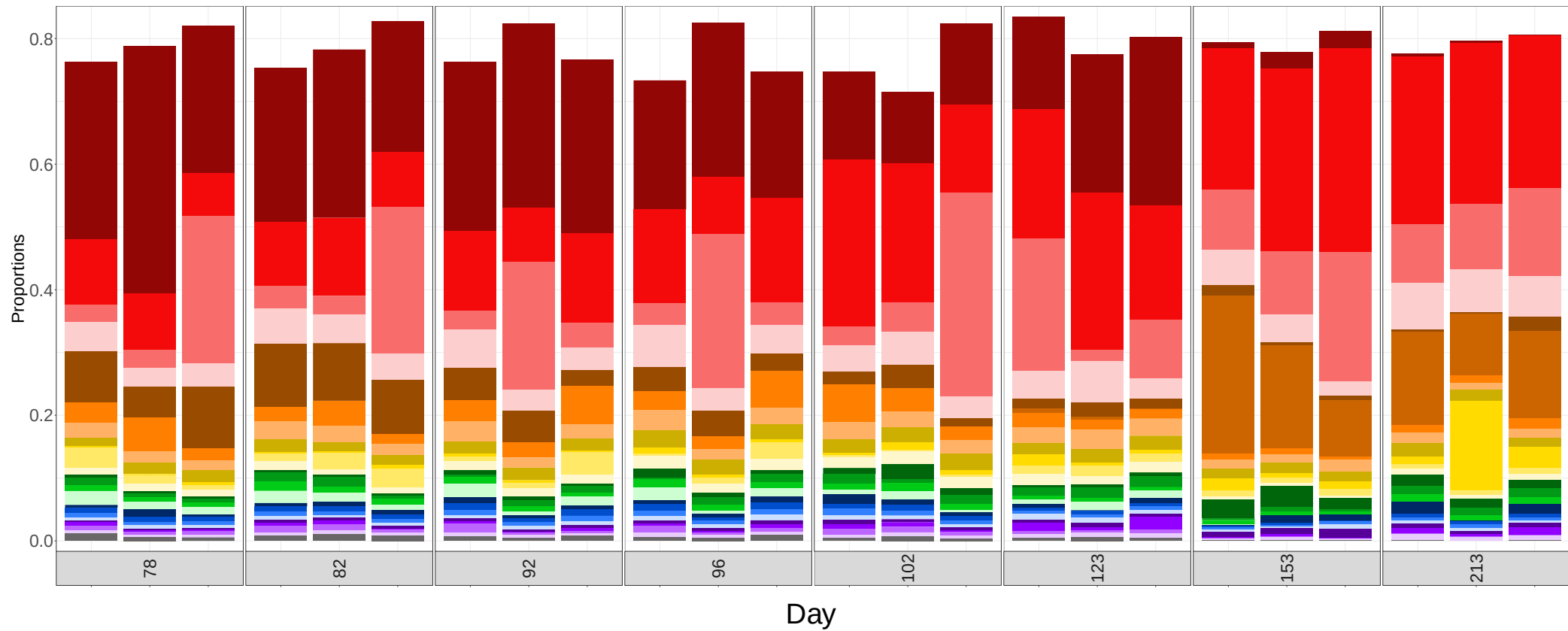




Taxa

- bin.333:d Bacteria;p Cloacimonadota;c Cloacimonadia;o Cloacimonadales;f Cloacimonadaceae;g Syntrophosphaera;s__Syntrophosphaera sp012516855
- bin.227:d Bacteria;p Bacteroidota;c Bacteroidia;o Bacteroidales;f UBA932;g DMER64;s DMER64 sp002427525
- bin.21:d Bacteria;p Firmicutes;c Bacilli;o RFN20;f CAG-826;g JAAYSA01;s JAAYSA01 sp012518455
- bin.218:d Bacteria;p Firmicutes_G;c Limnochordia;o DTU010;f DTU010;g UBA5420;s UBA5420 sp012838305
- bin.275:d Bacteria;p Firmicutes;c Bacilli;o Acholeplasmatales;f Acholeplasmataceae;g DUQW01;s
- bin.211:d Bacteria;p Cloacimonadota;c Cloacimonadia;o Cloacimonadales;f Cloacimonadaceae;g UBA3900;g UBA3900 sp002391675
- bin.184:d Bacteria;p Firmicutes_B;c Syntrophomonadales;f Syntrophomonadales;g Syntrophomonas_B;s__Syntrophomonas_B sp012729495
- bin.99:d Bacteria;p Firmicutes_G;c DTU065;o DTU065;f DTU065;g DTU065;s DTU065 sp001512545
- bin.106:d Bacteria;p Firmicutes_A;c Clostridia;o Acetivibrionales;f Acetivibrionaceae;g DTU013;s DTU013 sp002385815
- bin.50:d Bacteria;p Firmicutes_B;c Syntrophomonadia;o Syntrophomonadales;f Syntrophomonadaceae;g DTU018;s DTU018 sp002305535
- bin.170:d Bacteria;p Firmicutes_A;c Clostridia;o Tissierellales;f Sedimentibacteraceae;g Sedimentibacter;s Sedimentibacter sp012520455
- bin.276:d Bacteria;p Firmicutes_A;c Clostridia;o Acetivibrionales;f Acetivibrionaceae;g DTU013;s DTU013 sp012799845
- bin.204:d Archaea;p Halobacteriota;c Methanomicrobia;o Methanomicrobiales;f Methanospirillaceae;g Methanospirillum;s Methanospirillum sp012520015
- bin.292:d Bacteria;p Firmicutes_A;c Clostridia;o Oscillospirales;f Acutalibacteraceae;g DTU053;s DTU053 sp012520115
- bin.58:d Bacteria;p Firmicutes_A;c Clostridia;o Acetivibrionales;f DSM-8532;g DTU059;s DTU059 sp012523705
- bin.125:d Bacteria;p Bacteroidota;c Bacteroidia;o Bacteroidales;f 4484-276;g RZYY01;s RZYY01 sp012516825
- bin.217:d Bacteria;p Firmicutes_A;c Clostridia_A;o Christensenellales;f CAG-74;g DTU024;s
- bin.53:d Bacteria;p Firmicutes_A;c Clostridia;o Peptostreptococcales;f Anaerovoracaceae;g UBA1426;s
- bin.206:d Bacteria;p Bacteroidota;c Bacteroidia;o Bacteroidales;f Dysgonomonadaceae;g Petrimonas;s Petrimonas mucosa
- bin.266:d Bacteria;p Firmicutes_A;c Clostridia;o Lachnospirales;f Lachnospiraceae;g Herbinix;s Herbinix sp012523775
- bin.220:d Archaea;p Halobacteriota;c Methanosarcinia;o Methanosarcinales;f Methanosarcinaceae;g Methanosarcina;s__Methanosarcina sp002499445
- bin.320:d Bacteria;p Firmicutes_B;c Syntrophomonadales;f Syntrophomonadaceae;g ;s__
- bin.187:d Bacteria;p Firmicutes;c Bacilli;o ML615J-28;f CAG-313;g UBA1427;s UBA1427 sp002329485
- bin.155:d Bacteria;p Firmicutes_G;c SHA-98;o UBA4971;f UBA4971;g UBA4971;s UBA4971 sp900019985
- bin.63:d Bacteria;p Bacteroidota;c Bacteroidia;o Bacteroidales;f VadinHA17;g SR-FBR-E99;s SR-FBR-E99 sp009881065