S4 Fig. Quantification of Chloroflexi (using 16S rRNA gene amplicon sequencing and qFISH). Comparison of data from 16S rRNA amplicon sequencing and FISH regarding abundance of Chloroflexi after 6 and 7 weeks. FISH data is shown as % of EUBmix that are positive with the CFXmix probe (CFX122316 and GNSB94117) and amplicon data (amp) is shown as % of total community reads.