**Supplementary figures**



**Supplementary Figure 1**. Comparison between Relative Abundance of *P. copri* by PCR and proportion of full metagenomic reads mapped to *Prevotella copri* DSM 18205. A. There was absolute agreement between the *P. copri* PCR and the unbiased NGS with 4/7 of the samples not detecting *P. copri*. B. Of three samples with detectable *P. copri* the two methods correlated almost perfectly (r2=0.99, p=0.001).