

Table S1. Effects of different antibiotics on taxa in the faecal microbiome. Only genera or higher taxon present in at least 20% of the samples are shown. The colored cells indicate the time points at which a significant difference in the relative abundance of the taxon was observed in the antibiotic group compared to the respective placebo group. Red – decrease, green – increase compared to the placebo (Mann-Whitney test, $p<0.005$). Only taxa in the dark-colored cells remained significant after Bonferroni correction for multiple comparisons. BL – baseline, W1 – week 1, M1 – month 1, M2 – month 2, M4 – month 4, M12 – month 12. Mean - average relative abundance of each taxon per study.

Taxon	KI study								HP study								Mean (%)	
	Ciprofloxacin				Clindamycin				Amoxicillin				Minocycline					
	BL	W1	M1	M2	M4	M12	BL	W1	M1	M2	M4	M12	BL	W1	M1	M2	M4	M12
Actinobacteria	Actinobacteria;Bifidobacteriales;Bifidobacteriaceae;Bifidobacterium																0.17	0.25
	Actinobacteria;Bifidobacteriales;Bifidobacteriaceae;Gardnerella																0.16	0.36
	Coriobacteria;Coriobacteriales;Coriobacteriaceae;Adlercreutzia																0.02	0.03
	Coriobacteria;Coriobacteriales;Coriobacteriaceae;Collinsella																0.01	0.04
	Coriobacteria;Coriobacteriales;Coriobacteriaceae;Eggerthella																0.19	0.03
	Coriobacteria;Coriobacteriales;Coriobacteriaceae;Other																0.01	0.03
Bacteroidetes	Coriobacteria;Coriobacteriales;Coriobacteriaceae;Uncultured																0.05	0.03
	Bacteroidia;Bacteroidales;Bacteroidaceae;Bacteroides																20.2	18.6
	Bacteroidia;Bacteroidales;Other;Other																6.05	12.5
	Bacteroidia;Bacteroidales;Porphyromonadaceae;Barnesiella																0.63	0.76
	Bacteroidia;Bacteroidales;Porphyromonadaceae;Butyrimonas																0.05	0.07
	Bacteroidia;Bacteroidales;Porphyromonadaceae;Orobacter																0.28	0.28
	Bacteroidia;Bacteroidales;Porphyromonadaceae;Parabacteroides																1.81	2.58
	Bacteroidia;Bacteroidales;Prevotellaceae;Paraprevotella																0.23	0.31
	Bacteroidia;Bacteroidales;Prevotellaceae;Prevotella																5.6	10.4
	Bacteroidia;Bacteroidales;Ratan060301c;Ratan060301c																5.2	3.58
	Bacteroidia;Bacteroidales;Rikenellaceae;Alistipes																2.6	2.66
	Bacteroidia;Bacteroidales;S24-7;S24-7																0.90	0.42
Firmicutes	Bacilli;Lactobacillales;Streptococcaceae;Streptococcus																0.37	0.42
	Clostridia;Clostridiales;Clostridiaceae;Clostridium																0.04	0.05
	Clostridia;Clostridiales;Family_xiii_incertae_sedis;Other																0.03	0.03
	Clostridia;Clostridiales;Family_xiii_incertae_sedis;Uncultured																0.12	0.16
	Clostridia;Clostridiales;Lachnospiraceae;Anaerostipes																1.0	0.92
	Clostridia;Clostridiales;Lachnospiraceae;Blautia																2.1	2.07
	Clostridia;Clostridiales;Lachnospiraceae;Christensenella																0.06	0.03
	Clostridia;Clostridiales;Lachnospiraceae;Coprococcus																0.74	0.66
	Clostridia;Clostridiales;Lachnospiraceae;Dorea																0.50	0.93
	Clostridia;Clostridiales;Lachnospiraceae;Incertae_sedis																0.26	0.18
	Clostridia;Clostridiales;Lachnospiraceae;Lachnospira																2.67	1.56
	Clostridia;Clostridiales;Lachnospiraceae;Lachnospiraceae																0.14	0.13
	Clostridia;Clostridiales;Lachnospiraceae;Marvinbryantia																0.06	0.04
	Clostridia;Clostridiales;Lachnospiraceae;Other																8.25	6.95
	Clostridia;Clostridiales;Lachnospiraceae;Pseudobutyryrivibrio																3.38	3.09
	Clostridia;Clostridiales;Lachnospiraceae;Roseburia																3.12	1.96
	Clostridia;Clostridiales;Lachnospiraceae;Uncultured																1.53	1.64
	Clostridia;Clostridiales;Other;Other																0.06	0.07
	Clostridia;Clostridiales;Peptococcaceae;Peptococcus																0.02	0.03
	Clostridia;Clostridiales;Peptostreptococcaceae;Incertae_sedis																0.12	0.21
	Clostridia;Clostridiales;Ruminococcaceae;Anaerotruncus																0.25	0.27
	Clostridia;Clostridiales;Ruminococcaceae;Faecalibacterium																11.8	9.78
	Clostridia;Clostridiales;Ruminococcaceae;Incertae_sedis																0.89	0.38
	Clostridia;Clostridiales;Ruminococcaceae;Oscillibacter																0.02	0.06
	Clostridia;Clostridiales;Ruminococcaceae;Other																3.78	3.52
	Clostridia;Clostridiales;Ruminococcaceae;Ruminococcus																0.85	0.89
	Clostridia;Clostridiales;Ruminococcaceae;Subdoligranulum																4.27	3.17
	Clostridia;Clostridiales;Ruminococcaceae;Uncultured																3.02	1.79
	Clostridia;Clostridiales;Veillonellaceae;Dialister																0.27	0.20
	Clostridia;Clostridiales;Veillonellaceae;Phascolarctobacterium																0.36	0.29
	Clostridia;Clostridiales;Veillonellaceae;Veillonella																0.04	0.06
	Clostridia;Other;Other;Other																0.03	0.05
	Erysipelotrichi;Erysipelotrichales;Erysipelotrichaceae;Holdemania																0.04	0.03
	Erysipelotrichi;Erysipelotrichales;Erysipelotrichaceae;Incertae_sedis																0.74	0.59
	Erysipelotrichi;Erysipelotrichales;Erysipelotrichaceae;Turicibacter																0.04	0.03
	Erysipelotrichi;Erysipelotrichales;Erysipelotrichaceae;Uncultured																0.04	0.02
Proteobacteria	Alphaproteobacteria;Rhodospirillales;Rhodospirillaceae;Thalassospira																0.13	0.11
	Betaproteobacteria;Burkholderiales;Alcaligenaceae;Parasutterella																0.15	0.07
	Betaproteobacteria;Burkholderiales;Alcaligenaceae;Sutterella																0.36	0.37
	Deltaproteobacteria;Desulfovibrionales;Desulfovibrionaceae;Bilophila																0.07	0.12
	Deltaproteobacteria;Desulfovibrionales;Desulfovibrionaceae;Desulfovibrio																0.1	0.10
	Gammaproteobacteria;Enterobacteriales;Enterobacteriaceae;Escherichia-shigella																0.05	0.22
	Tenericutes;Mollicutes;Rf9;Uncultured_bacterium;Rf9																0.08	0.19
	Unclassified Bacteria																0.20	0.14
	Verrucomicrobia;Verrucomicrobiae;Verrucomicrobiales;Verrucomicrobiaceae;Akkermansia																0.47	0.41