

Table S6 Primers used for molecular community analysis

Target	Primer name	Primer sequence (universal tag sequence in bold)	amplicon size
Universal tag primers	TAG-for	Cyanine5-AGGTGACACTATAGAATA	
	TAG-rev	GTACGACTCACTATAGGGAA	
<i>Corynebacterium glutamicum</i> DSMZ 1412	tag-CGTerC_F	AGGTGACACTATAGAATAACCACTGAGGGCGATCTTGT	183
	tag-CGTerC_R	GTACGACTCACTATAGGGAAATGTGCTGGCTCCGAAGA	
<i>Faecalibacterium prausnitzii</i> A2-165	tag_Fprau(CG447_RS01330F	AGGTGACACTATAGAATAACAGCCCCAGCATCTCTTG	135
	tag_Fprau(CG447_RS01330R	GTACGACTCACTATAGGGAGGATACCGAGGGAAAGAGTGGAC	
<i>Subdoligranulum variabile</i> DSM 15176	Svar_SUBVAR_05219F	AGGTGACACTATAGAATAATGGAACCGATTCCTGCTTG	170
	Svar_SUBVAR_05219R	GTACGACTCACTATAGGAAAGCAATGAGCTGTCCGATGA	
<i>Eubacterium rectale</i> A1-86 (=DSM 17629)	tag_Erect_EUR_16390F	AGGTGACACTATAGAATAATGCCCTCAGGCACCACTA	147
	tag_Erect_EUR_16390R	GTACGACTCACTATAGGATGGCCTTAAGGTGGCAACA	
<i>Roseburia intestinalis</i> M50/1	Rint_ROI_11740F	AGGTGACACTATAGAATAATCGCAATGAATGCCCTGC	254
	Rint_ROI_11740R	GTACGACTCACTATAGGAAAGATGTCCGCCCTCACAG	
<i>Roseburia faecis</i> M72/1	tag-RFTerC_F_1	AGGTGACACTATAGAATAAACAGTTCCGCTCCCTTGG	339
	tag-RFTerC_R_1	GTACGACTCACTATAGGACCATTGGTCCGGTGTGTA	
<i>Roseburia inulinivorans</i> A2-194	tag_Rinul_ROSEINA2194_04328F	AGGTGACACTATAGAATAACGAGCAGATTTACAGCGT	275
	tag_Rinul_ROSEINA2194_04328R	GTACGACTCACTATAGGAAACCTCATGCCCTGCAATCG	
<i>Coprococcus eutactus</i> ART55/1	tag-CETerC_F	AGGTGACACTATAGAATAAGGAAAACCAGCTCGTGA	165
	tag-CETerC_R	GTACGACTCACTATAGGACCCCTGCTGGAACTTCAAC	
<i>Streptococcus thermophilus</i> CNCM I-3862	tag_S.thermo - F	AGGTGACACTATAGAATAATTGGAAAGGGCAATTGCT	318
	tag_S.thermo - R	GTACGACTCACTATAGGAGTGAACCTCCACTCTCACAC	
<i>Lactobacillus paracasei</i> CNCM I-1518	tag_OFF 2201 ex	AGGTGACACTATAGAATAIIGTAGCACCCTAAAGACG	322
	tag_OFF 2202 ex	GTACGACTCACTATAGGAIIGCCATAAGCGTGTAGCCG	
<i>Bifidobacterium bifidum</i> CNCM I-3650	tag_GeXP_Bif_genus_F1	AGGTGACACTATAGAATAACGCGTCYGGTGTGAAAG	165
	tag_GeXP_Bif_genus_Rev	GTACGACTCACTATAGGAGGTGTTCTCCGATATCTACA	