

Table S2. Average relative abundances (%) of enterotype indicator species in the non-secretors, the secretors and the individuals with FUT2 genotypes AA, AG or GG belonging to the enterotype 1 (ET1) or 3 (ET3).

Taxa	Non-	Non-			Genotype AG (ET3)	Genotype GG (ET3)
	secretors (ET1)	Genotype AG (ET1)	secretors (ET3)	Secretors (ET3)		
<i>Bacteroides fragilis</i> et rel.	1.7	0.6	0.6	1.2	1.4	0.9
<i>Bacteroides ovatus</i> et rel.	1.7	1.1	0.6	1.2	1.3	1.1
<i>Bacteroides vulgatus</i> et rel.	1.3	1.6	0.7	1.7	1.9	1.4
<i>Clostridium symbiosum</i> et rel.	4.0	4.4	3.1	2.0	2.4	1.6
<i>Roseburia intestinalis</i> et rel.	3.8	4.0	2.0	1.4	1.8	0.9
<i>Ruminococcus bromii</i> et rel.	1.1	0.1	2.7	3.3	2.7	3.8
<i>Ruminococcus callidus</i> et rel.	0.9	2.2	1.7	1.4	1.9	0.9
<i>Sporobacter termitidis</i> et rel.	0.3	0.6	1.1	1.6	1.3	1.8
<i>Tannerella</i> et rel.	0.4	0.3	1.2	1.3	1.9	0.7
<i>Uncultured Clostridiales I</i>	0.1	0.01	2.5	5.5	2.8	8.2