Table S3 Read counts in microbiome sample DO-360 broken down by observed SNP allele and by the genotype of genomic DNA sample DO-370.

	allele in DO-360 microbiome			
DO-370 genotype	A (%)		B (%)	
AA	10,324,265	(99.8%)	23,256	(0.2%)
AB	2,083,947	(51.2%)	1,986,380	(48.8%)
BB	5,347	(0.5%)	1,117,994	(99.5%)