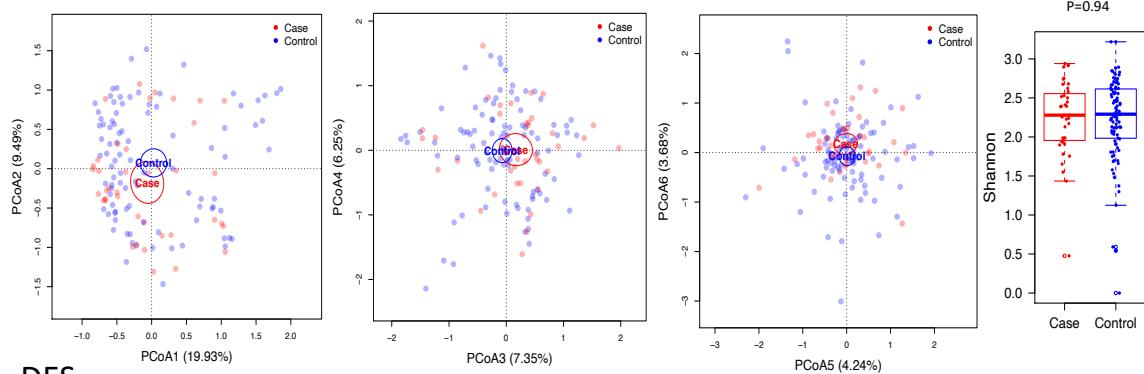


Fig. S1. Selection criteria for the current analysis.

ASC

Case_Ctrl R² = 0.01159 P= 0.09



DES

Case_Ctrl R² = 0.0147 P= 0.031

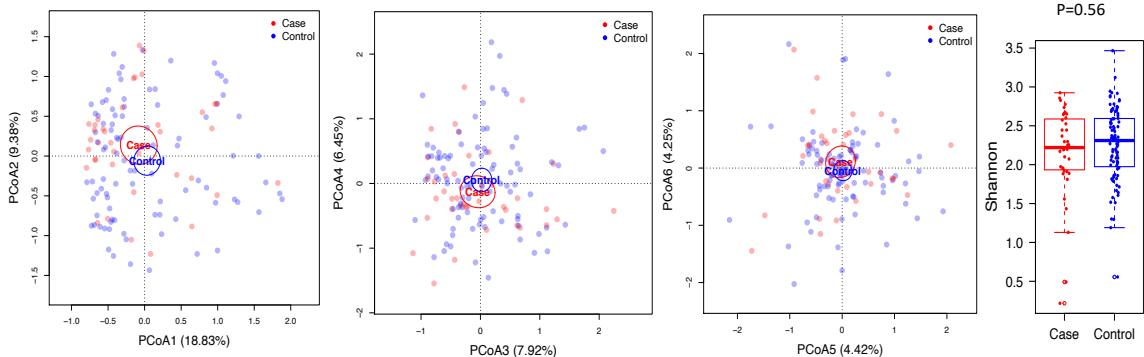


Fig. S2. PCoA plots and alpha-diversity of the ascending (ASC) and descending (DES) colon microbiomes of study participants without antibiotics intake (n=132: 39 cases and 93 controls). The ellipses in PCoA plots indicate 95% confidence limits. The boxplots showed the median, 25th and 75th percentile.

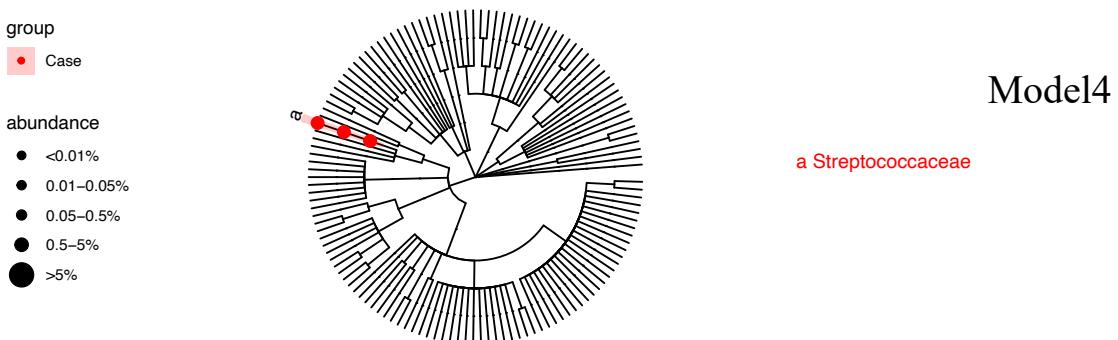
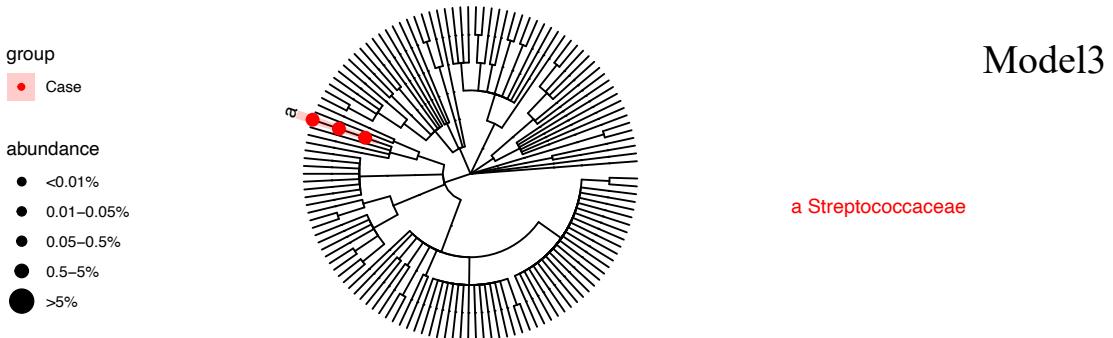
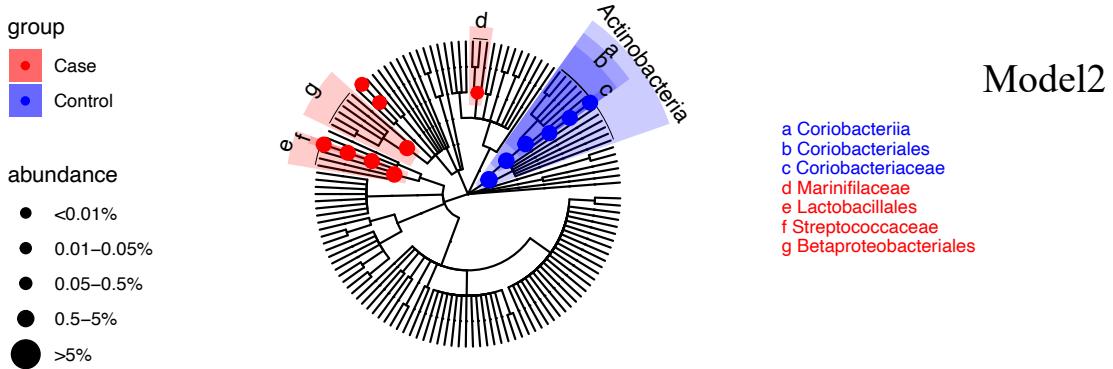
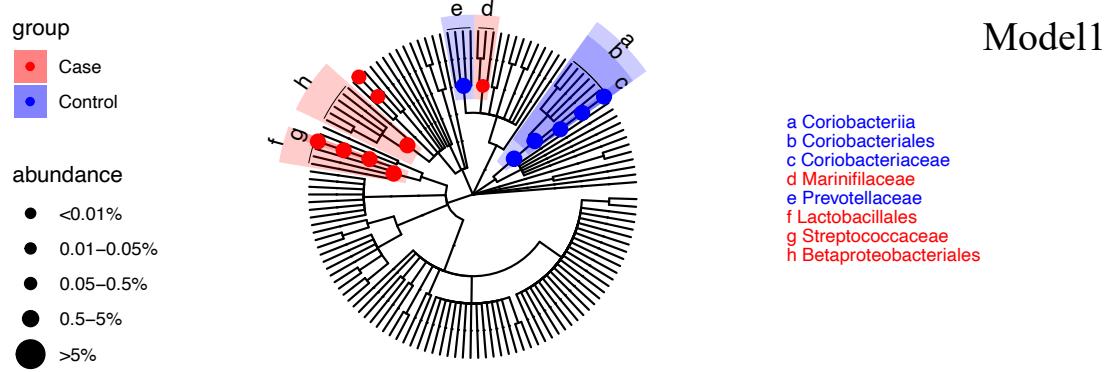


Fig. S3. Significant differential taxa in the descending (DES) colon microbiome between microscopic colitis cases and controls that did not use antibiotics. Model 1 is adjusted for batch, education and PPI use. Model 2 is additionally adjusted for sex. Model 3 is adjusted for age along with the covariates in model 2. Model 4 is adjusted for BMI along with the covariates in model 3. The branches of significant taxa from phylum to family level were highlighted and labeled. The node sizes are proportional to the overall relative abundance of the taxa.

Table S1. The significant associations between individual taxa and case/control adjusted for covariates (see the 4 models in methods) in each of the stratified levels of liquid stools (0-2, 3, 4-5, >6) in the ascending (ASC) and descending (DES) colon microbiome. Insignificant results are not shown.

a.

	ASC, Liquid stools: 0-2							
	model 1				model 2			
	Stats	P	FDR	Enriched in	Stats	P	FDR	Enriched in
g_Eggerthella--s uncultured bacterium	-3.88	3.54E-04	0.087	Case	-4.04	2.23E-04	0.073	Case
f_Marinifilaceae	Insignificant				-3.36	1.68E-03	0.092	Case
f_Muribaculaceae	Insignificant				-3.35	1.69E-03	0.092	Case
f_Christensenellaceae	-3.51	1.07E-03	0.087	Case	-3.48	1.19E-03	0.092	Case
g_Christensenellaceae_R-7_group	-3.58	8.57E-04	0.087	Case	-3.55	9.52E-04	0.092	Case
g_Akkermansia--unclassified	-3.54	9.65E-04	0.087	Case	-3.68	6.53E-04	0.092	Case

b.

	ASC, Liquid stools: 3							
	model 1				model 2			
	Stats	P	FDR	Enriched in	Stats	P	FDR	Enriched in
g_Ruminococcaceae_UCG-002	Insignificant				Insignificant			
g_Ruminococcaceae_UCG-002--unclassified	Insignificant				Insignificant			
g_Ruminococcaceae_UCG-003	-4.41	2.22E-04	0.036	Case	-4.03	6.01E-04	0.096	Case
g_Ruminococcaceae_UCG-003--unclassified	-4.41	2.22E-04	0.036	Case	-4.03	6.01E-04	0.096	Case
o_Betaproteobacteriales	Insignificant				Insignificant			
f_Burkholderiaceae	Insignificant				Insignificant			
	model 4							
	Stats	P	FDR	Enriched in				
g_Ruminococcaceae_UCG-002	-3.97	9.92E-04	0.065	Case				
g_Ruminococcaceae_UCG-002--unclassified	-3.97	9.92E-04	0.065	Case				
g_Ruminococcaceae_UCG-003	-4.55	2.83E-04	0.045	Case				
g_Ruminococcaceae_UCG-003--unclassified	-4.55	2.83E-04	0.045	Case				
o_Betaproteobacteriales	-3.96	1.02E-03	0.065	Case				

f_Burkholderiaceae	-3.85	1.28E-03	0.068	Case				
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c.

		DES, Liquid stools: 0-2							
		model 1				model 2			
		Stats	P	FDR	Enriched in	Stats	P	FDR	Enriched in
f_Marinilaceae		Insignificant			-3.28	2.09E-03	0.080	Case	
g_Odoribacter--unclassified		Insignificant			-3.36	1.66E-03	0.072	Case	
f_Muribaculaceae		Insignificant			-3.17	2.88E-03	0.098	Case	
c_Deltaproteobacteria		Insignificant			-3.55	9.69E-04	0.049	Case	
o_Desulfovibrionales		Insignificant			-3.60	8.27E-04	0.049	Case	
f_Desulfovibrionaceae		Insignificant			-3.60	8.27E-04	0.049	Case	
g_Bilophila	-3.78	4.83E-04	0.060	Case	-4.02	2.37E-04	0.036	Case	
g_Bilophila--unclassified	-3.75	5.28E-04	0.060	Case	-3.89	3.55E-04	0.036	Case	
g_Akkermansia--unclassified	-3.71	5.92E-04	0.060	Case	-3.89	3.53E-04	0.036	Case	
		model 4							
		Stats	P	FDR	Enriched in				
f_Marinilaceae		Insignificant							
g_Odoribacter--unclassified		Insignificant							
f_Muribaculaceae		Insignificant							
c_Deltaproteobacteria	-3.36	1.84E-03	0.096	Case					
o_Desulfovibrionales	-3.46	1.41E-03	0.096	Case					
f_Desulfovibrionaceae	-3.46	1.41E-03	0.096	Case					
g_Bilophila	-3.86	4.55E-04	0.096	Case					
g_Bilophila--unclassified	-3.36	1.85E-03	0.096	Case					
g_Akkermansia--unclassified	-3.35	1.89E-03	0.096	Case					

d.

		DES, Liquid stools: 3							
		model 1				model 2			
		Stats	P	FDR	Enriched in	Stats	P	FDR	Enriched in
g_Roseburia	-3.53	1.88E-03	0.085	Case	Insignificant				
g_Roseburia--unclassified	-3.53	1.88E-03	0.085	Case	Insignificant				
g_Ruminococcaceae_UCG-003	-3.59	1.62E-03	0.085	Case	Insignificant				
g_Ruminococcaceae_UCG-003--unclassified	-3.59	1.62E-03	0.085	Case	Insignificant				
o_Betaproteobacteriales	-5.48	1.66E-05	0.005	Case	-5.06	5.23E-05	0.016	Case	
f_Burkholderiaceae	-5.15	3.63E-05	0.005	Case	-4.76	1.05E-04	0.016	Case	

g__Sutterella	- 3.45	2.29E-03	0.085	Case	Insignificant			
g__Sutterella--unclassified	- 3.50	2.00E-03	0.085	Case	Insignificant			
	DES, Liquid stools: 3							
	model 3					model 4		
	Stats	P	FDR	Enriched in	Stats	P	FDR	Enriched in
g__Roseburia	Insignificant				Insignificant			
g__Roseburia--unclassified	Insignificant				Insignificant			
g__Ruminococcaceae_UCG-003	Insignificant				Insignificant			
g__Ruminococcaceae_UCG-003--unclassified	Insignificant				Insignificant			
o__Betaproteobacteriales	- 5.13	5.92E-05	0.015	Case	- 6.23	9.21E-06	0.003	Case
f__Burkholderiaceae	- 4.90	9.89E-05	0.015	Case	- 5.83	2.01E-05	0.003	Case
g__Sutterella	Insignificant				Insignificant			
g__Sutterella--unclassified	Insignificant				Insignificant			