Table 1

Clinical data and genomic abnormalities in colorectal cancer subtypes categorized based on gut microbiota.

	Classification based on gut microbiota			
	Subtype 1	Subtype 2	Subtype 3	Subtype 4
Age		Older	Younger	
Body mass index		Higher		Lower
Smoking history		Less		
Alcohol consumption	Higher			
Hypermutated phenotype		More		
Significantly mutated genes		*ARID1A, B2M, PTEN*		
Mutational signatures	ID1	SBS44, ID2	SBS88, ID18, Novel1	
Copy number alterations	More amplification			
Timing of CNN-LOH / loss+gain		Late	Early	
Immune cell profiling			Regulatory T cells ↑	
Microenvironment			Immunosuppressive	
Consensus molecular subtype (CMS)		CMS1		
Prognosis			Poorer	

CNN-LOH, copy number neutral-loss of heterozygosity