



Supplemental Figure S1 Enriched functions for active genes with the highest 10% closeness centralities (CC) in normal colons (A) and active genes in normal colons that underwent significant decrease ($p<0.05$) in CC in CRC (B). For each bar representing a pathway, the fold enrichment was manifested by the bar color. The vertical dashed lines indicate FDR-adjusted p-value equaling 0.05.