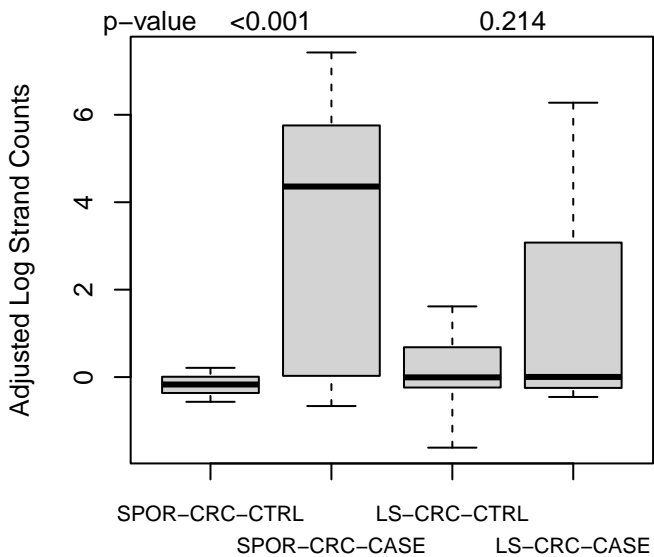
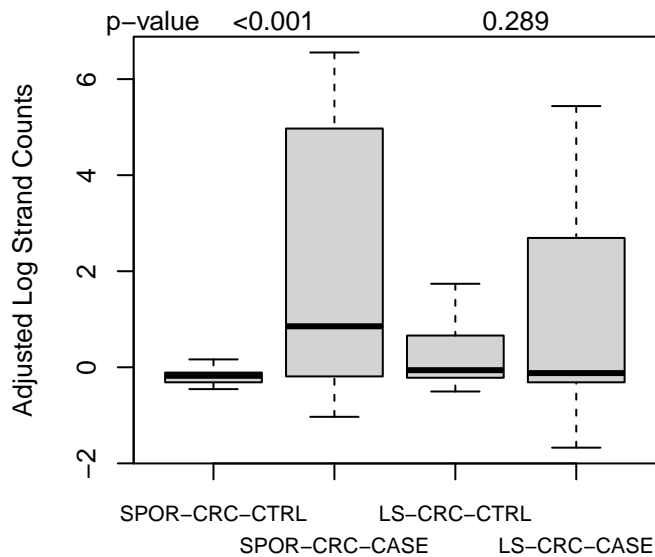


Supplemental Figure 1. Distributions of all methylated DNA markers tested for colorectal cancers and controls, stratified by Lynch syndrome. Log10 strand counts are adjusted based on the level of the control gene, ACTB, in all controls (SPOR-CRC-CTRL and LS-CRC-CTRL). P-value provided to show statistical significance of difference of medians between CASE and CTRL samples within SPOR and LS subgroups.

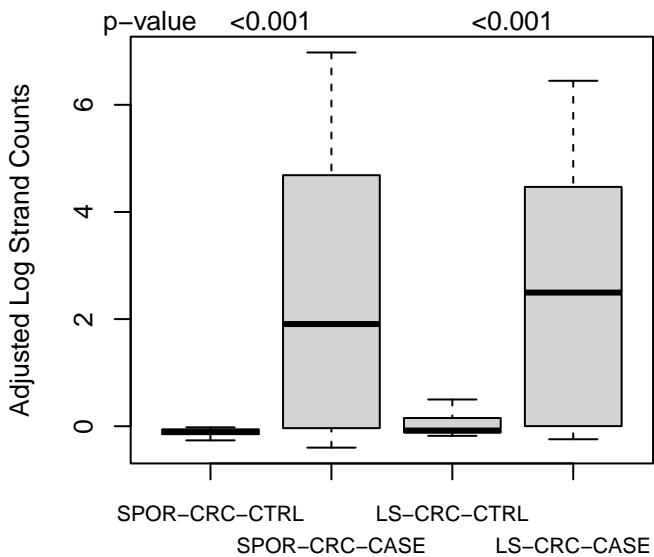
NDRG4



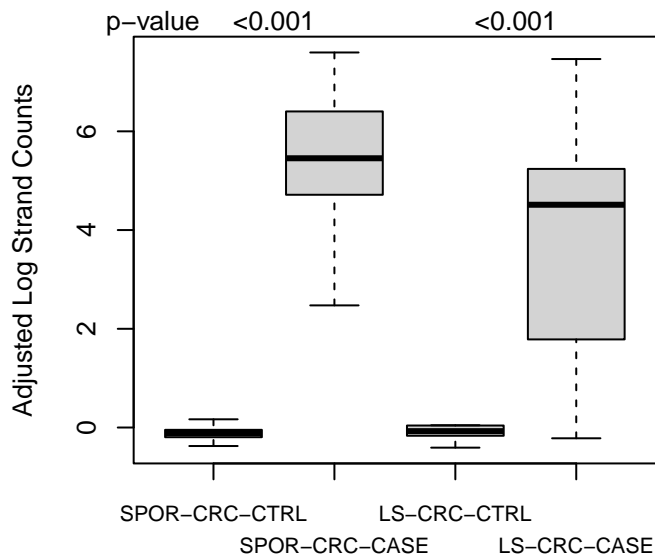
BMP3



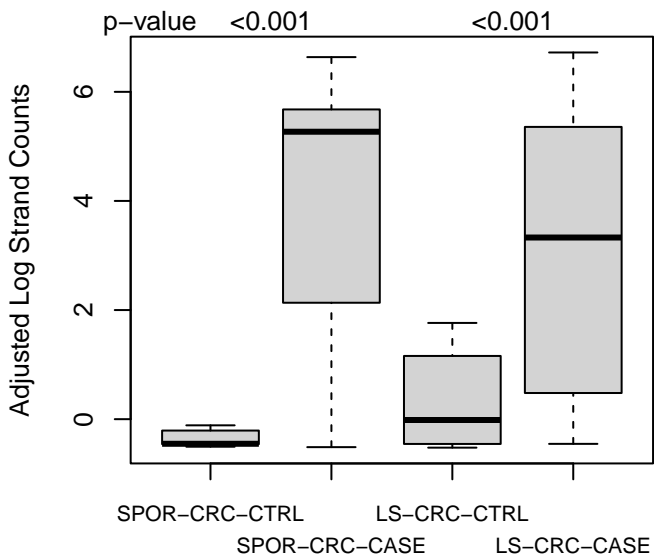
LASS4



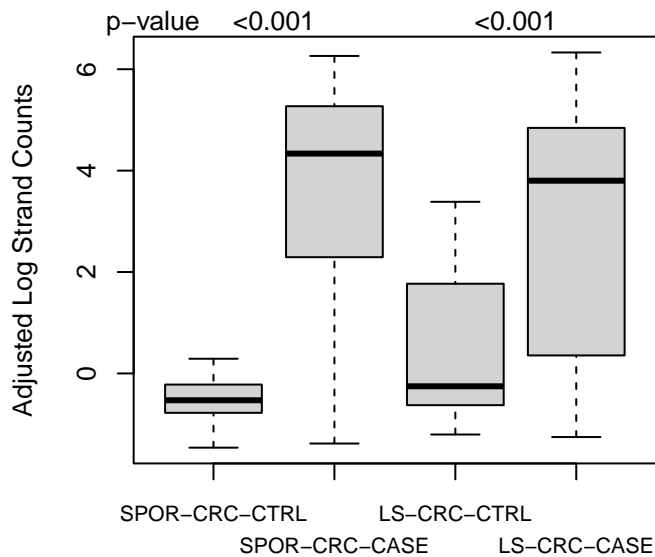
LRRC4



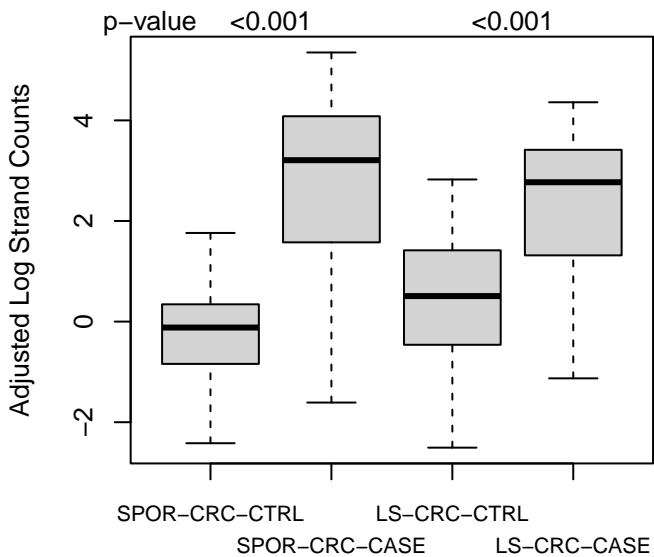
PPP2R5C



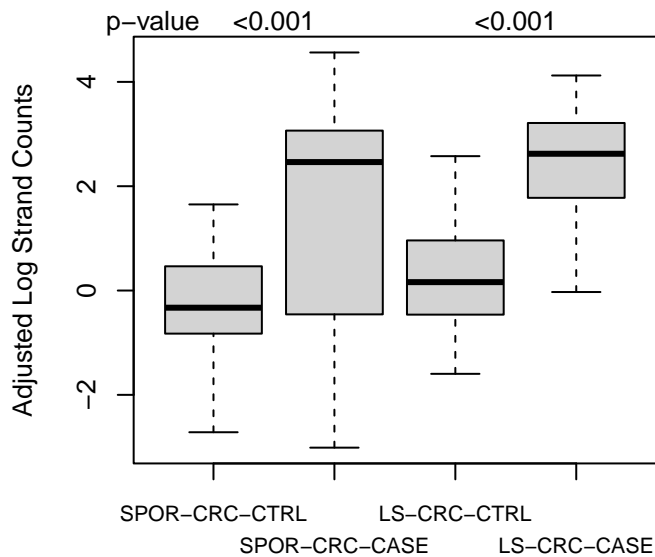
USP44



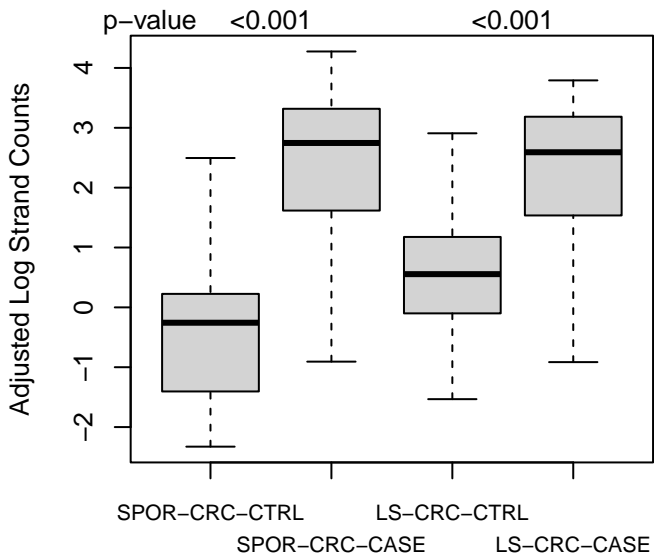
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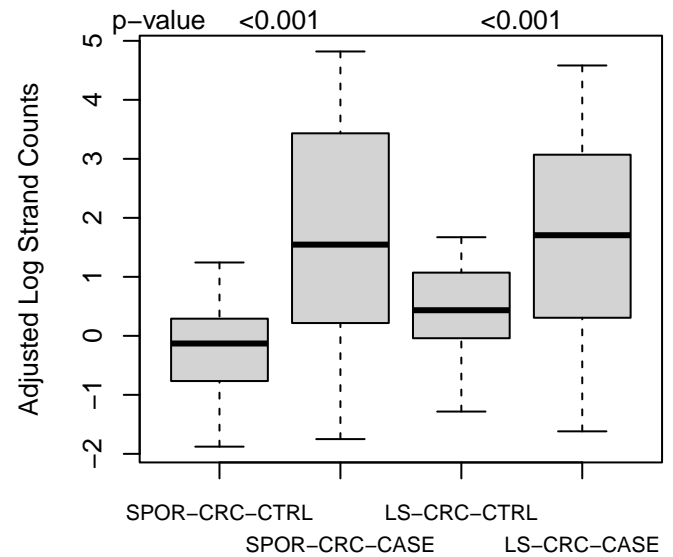
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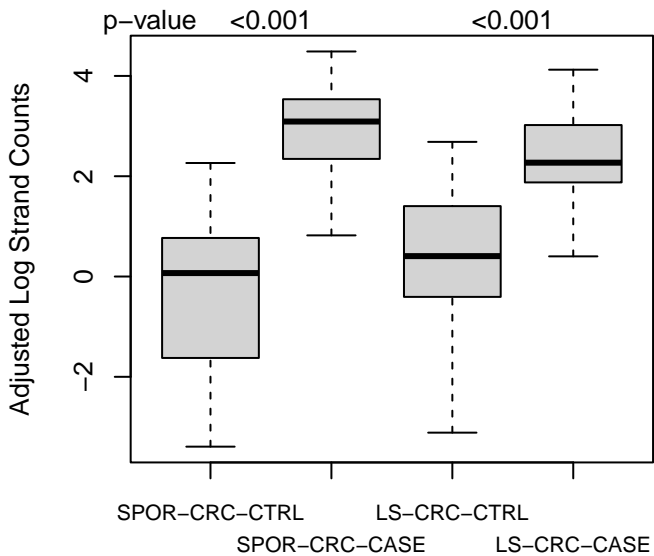
ADCY4



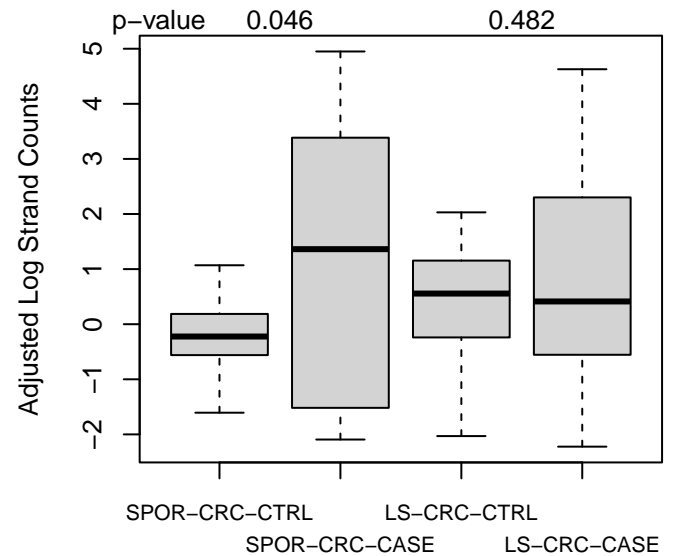
CNTFR



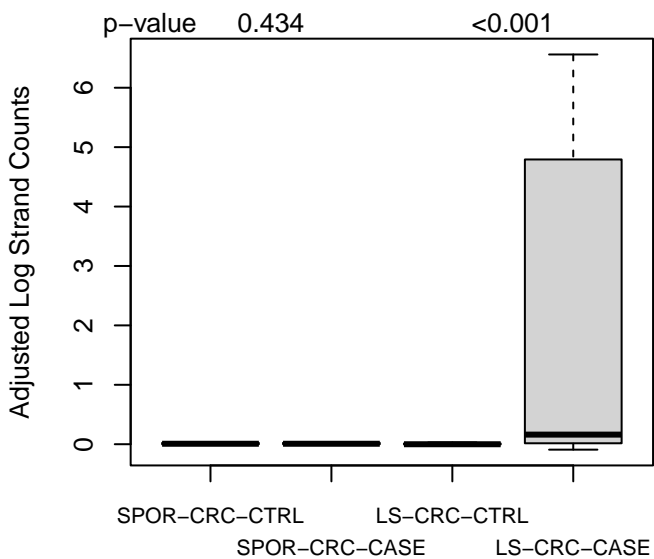
PITX1



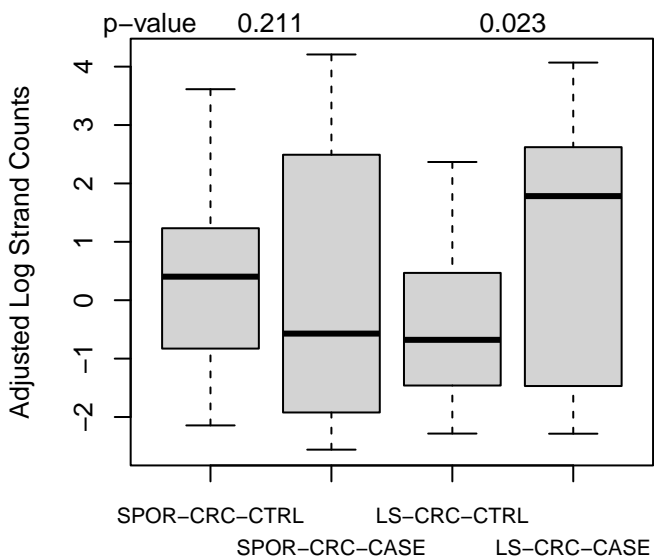
ANTXR1



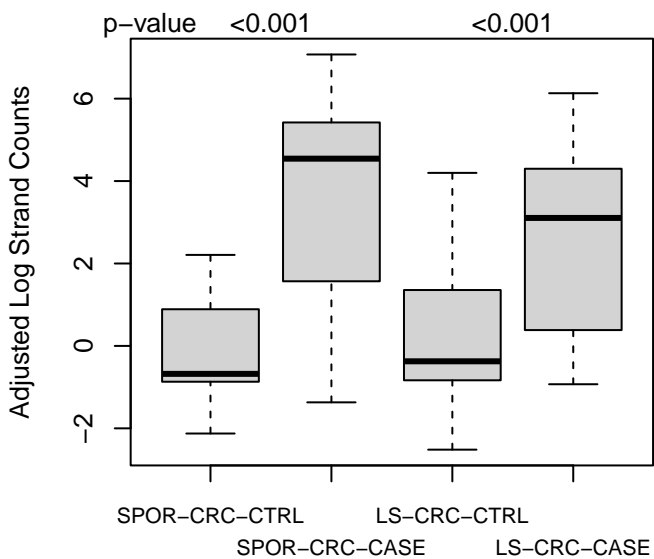
ALKBH5



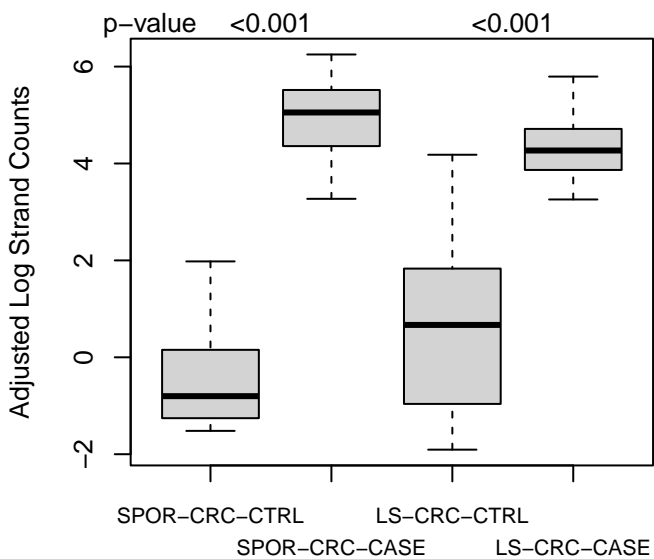
ADM



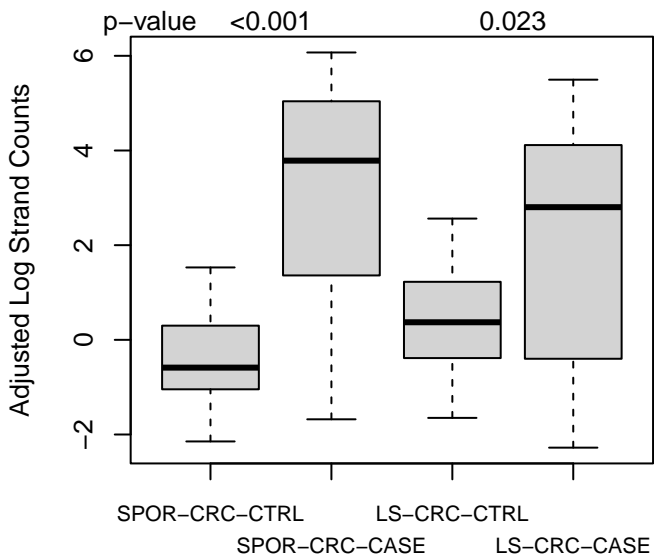
ELMO1



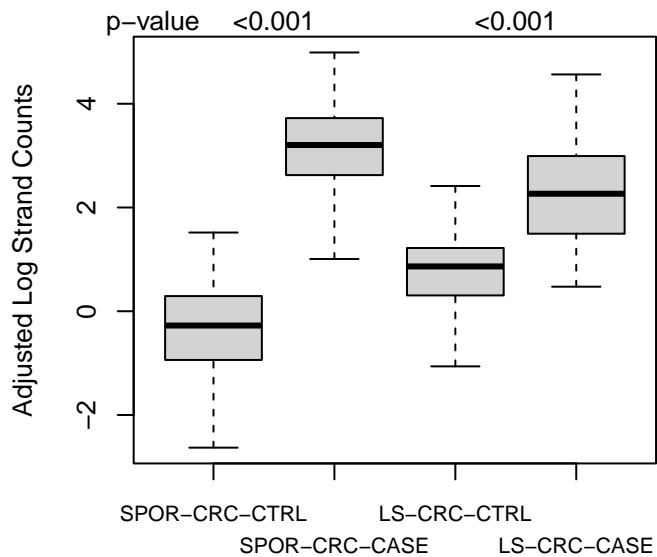
OPLAH



VAV3



ARHGEF4



SFMBT2

