Enrichment plot: HALLMARK_WNT_BETA_CATENIN_SIGNALING 0.2 Running Enrichment Score (RES) -0.2 Zero crossing at 12612 Peak at 21741 -0.4 9.0--0.8

Gene List Index Number of genes: 25221 (in list), 42 (in gene set)

15000

10000

"cancer"

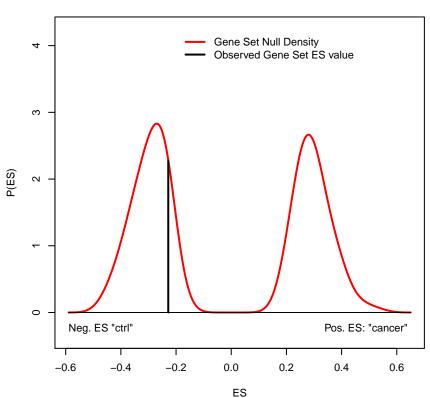
5000

"ctrl"

25000

20000

Gene Set Null Distribution



ES = -0.228 NES = -0.77 Nom. p-val= 0.854 FWER= 1 FDR= 1

Heat Map for Genes in Gene Set cancer ctrl CUL1 WNT6 WNT5B HEY1 HDAC11 CSNK1E MAML1 HDAC2 TCF7 SKP2 FRAT1 ADAM17 NOTCH1 LEF1 FZD1 FZD8 PPARD NCSTN NCOR2 AXIN2 NUMB CTNNB1 NKD1 GNAI1 PTCH1 KAT2A DKK4 RBPJ DLL1 TP53 PSEN2 HDAC5 DVL2 JAG1 MYC JAG2 DKK1 CCND2 NOTCH4 AXIN1 **B**2 **B**3 **B**4 **B**1