Top 30 Pathways (Sorted by FDR)

Top 30 pathways ranked by −log10(FDR), colored by −log10(p_value):

- TF – Factor: AP2BETA; motif: SCCYCAGGSNN; match class: 1
- TF – Factor: AP−2gamma; motif: NTGSCCTGRGGSNN; match class: 1
- TF – Factor: PURBETA; motif: NCWGRGGCTGGGG; match class: 1
- TF – Factor: ZNFPT1; motif: YCNNCNCWGCCNY; match class: 1
- TF – Factor: SALL2; motif: GGGTGGG; match class: 1
- TF – Factor: ZNF644; motif: TCCWGCCTCTSN
- TF – Factor: ZNF644; motif: TCCWGCCTCTSN; match class: 1
- TF – Factor: AP−2gamma; motif: NTGSCCTGRGGSNN
- TF – Factor: PUR1; motif: GGGNCAGNN; match class: 1
- TF – Factor: AP−2alpha; motif: NTNSCCTGRGGSNAN; match class: 1
- TF – Factor: NR1B1:RXR−ALPHA; motif: RRGGTCANNNNNRGGTCA
- TF – Factor: AP−2alpha; motif: NNNGSCCTGRGGSN
- TF – Factor: AP−2alpha; motif: NNNGSCCTGRGGSN; match class: 1
- TF – Factor: ZNF692; motif: SYNGGSCCCASCCNC; match class: 1
- TF – Factor: GKLF; motif: NNNRGGNGNGGSN; match class: 1
- TF – Factor: REST; motif: TTCAGCACCAYGGACAGCKCCN
- TF – Factor: KLF; motif: GGGNGGGG; match class: 1
- TF – Factor: RREB−1; motif: CCCCAAACMMCCCC
- TF – Factor: ZGPAT; motif: GRGGCWGNGGNG; match class: 1
- TF – Factor: ZNF300; motif: SWGGGGGGSN; match class: 1
- TF – Factor: NR1B2; motif: NTGACCY; match class: 1
- TF – Factor: GLI; motif: NGACCMCCCAN
- TF – Factor: NR1B1:RXR−ALPHA; motif: RRGGTCANNNNNRGGTCA; match class: 1
- TF – Factor: sin3A; motif: TGTCCNNGGTGCTG; match class: 1
- TF – Factor: PURBETA; motif: NCWGRGGCTGGGG
- TF – Factor: HNRPUL1; motif: NCNCAGN; match class: 1
- TF – Factor: LKLF; motif: CNCCACCCS
- TF – Factor: AP−2beta; motif: NSCCYNRGGSN
- TF – Factor: HDAC1; motif: KGCARGGTC; match class: 1
- TF – Factor: Ikaros; motif: TGGGAGN; match class: 1

x-axis: −log10(FDR)

Legend: −log10(p_value); values 5, 6, 7