Top 30 Pathways (Sorted by FDR)

Y-axis (top to bottom):
- TF – Factor: HDAC1; motif: KGCARGGTC; match class: 1
- TF – Factor: PURBETA; motif: NCWGRGGCTGGGG; match class: 1
- TF – Factor: p73; motif: NNRCAWGYCCARRCWTGYC
- TF – Factor: HNRPUL1; motif: NCNCAGN; match class: 1
- TF – Factor: NR1B2; motif: NTGACCY; match class: 1
- TF – Factor: ZNF644; motif: TCCWGCCTCTSN; match class: 1
- TF – Factor: ER–beta; motif: NRGGTCANKSTGACCTNN; match class: 1
- TF – Factor: Sox–8; motif: NGAACAATRN; match class: 1
- TF – Factor: LRF; motif: GGGGKYNNB; match class: 1
- TF – Factor: myogenin; motif: RGCAGSTG; match class: 1
- TF – Factor: NF–KAPPAB1; motif: NGGKRNTTYCCCN
- TF – Factor: NR1B2; motif: NTGACCY
- TF – Factor: ZNFPT1; motif: YCNNCNCWGCCNY; match class: 1
- TF – Factor: SREBP–1; motif: CACSCCA
- GO:MF – monoatomic cation channel activity
- TF – Factor: AP2BETA; motif: SCCYCAGGSNN
- TF – Factor: GLI; motif: NGACCMCCCAN; match class: 1
- TF – Factor: NR1B2; motif: RAGGTCATGACCTN
- TF – Factor: sin3A; motif: TGTCCNNGGTGCTG; match class: 1
- TF – Factor: GATAD2A; motif: CCTKTG; match class: 1
- TF – Factor: AP–2alpha; motif: NTNSCCTGRGGSNAN
- TF – Factor: TR4; motif: NCCCTGACCYB
- TF – Factor: VDR; motif: RRGGTCANNGRGKTCA
- CORUM – HAS1–HAS2 complex
- CORUM – ARTN–GFRA3–RET complex
- CORUM – FOXP3–TRIM28–ZFP90 complex
- CORUM – LPAR2–SIVA1 complex
- CORUM – LPAR2–TRIP6–NHERF2 complex, LPA stimulated
- CORUM – mGluR3–mGluR4 complex

X-axis: −log10(FDR)

Legend: −log10(p_value) — 1.5, 2.0, 2.5, 3.0, 3.5