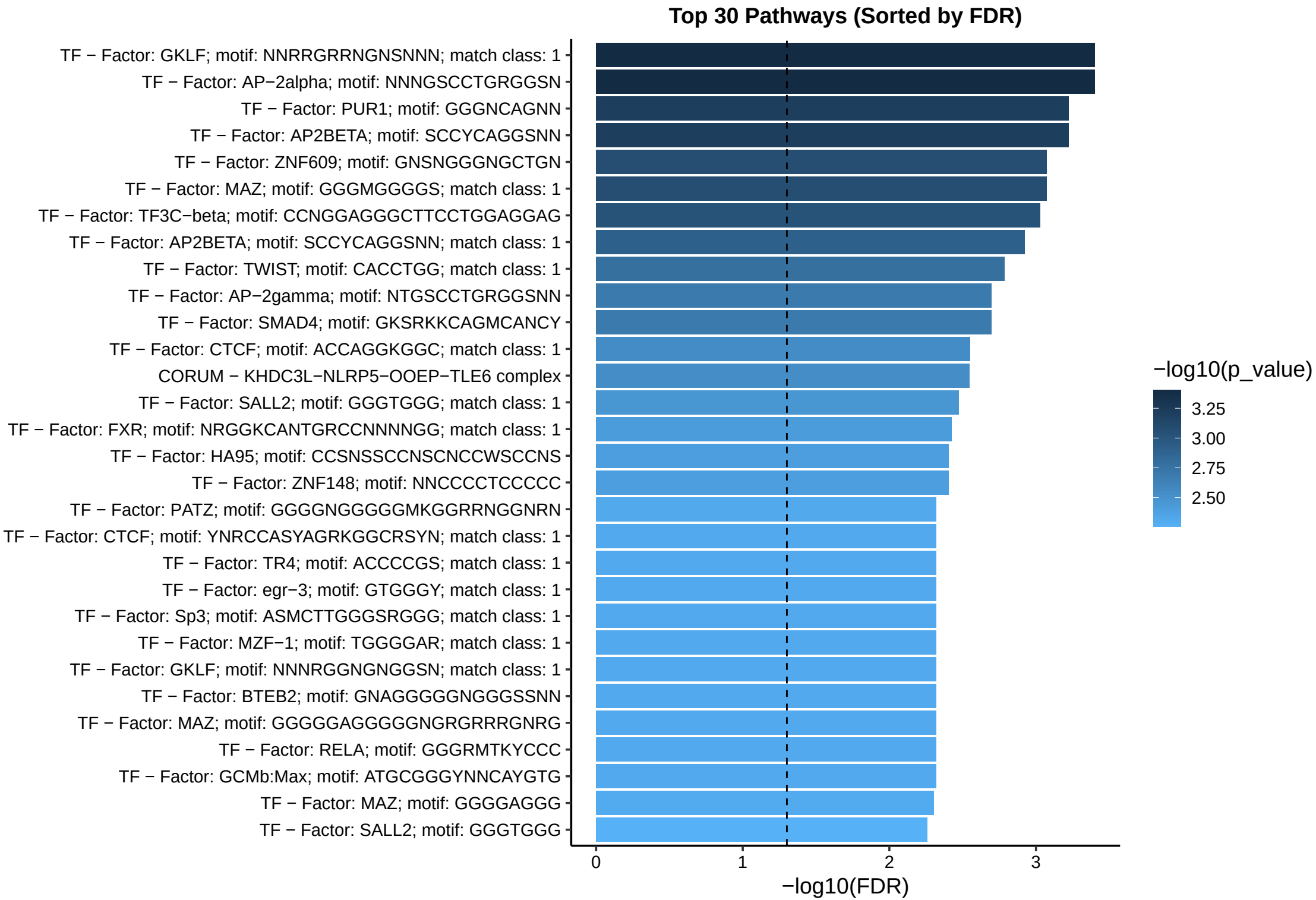
Top 30 Pathways (Sorted by FDR)
TF − Factor: GKLF; motif: NNRRGRRNGNSNNN; match class: 1
TF − Factor: AP−2alpha; motif: NNNGSCCTGRGGSN
TF − Factor: PUR1; motif: GGGNCAGNN
TF − Factor: AP2BETA; motif: SCCYCAGGSNN
TF − Factor: ZNF609; motif: GNSNGGGNGCTGN
TF − Factor: MAZ; motif: GGGMGGGGS; match class: 1
TF − Factor: TF3C−beta; motif: CCNGGAGGGCTTCCTGGAGGAG
TF − Factor: AP2BETA; motif: SCCYCAGGSNN; match class: 1
TF − Factor: TWIST; motif: CACCTGG; match class: 1
TF − Factor: AP−2gamma; motif: NTGSCCTGRGGSNN
TF − Factor: SMAD4; motif: GKSRKKCAGMCANCY
TF − Factor: CTCF; motif: ACCAGGKGGC; match class: 1
CORUM − KHDC3L−NLRP5−OOEP−TLE6 complex
TF − Factor: SALL2; motif: GGGTGGG; match class: 1
TF − Factor: FXR; motif: NRGGKCANTGRCCNNNNGG; match class: 1
TF − Factor: HA95; motif: CCSNSSCCNSCNCCWSCCNS
TF − Factor: ZNF148; motif: NNCCCCTCCCCC
TF − Factor: PATZ; motif: GGGGNGGGGGMKGGRRNGGNRN
TF − Factor: CTCF; motif: YNRCCASYAGRKGGCRSYN; match class: 1
TF − Factor: TR4; motif: ACCCCGS; match class: 1
TF − Factor: egr−3; motif: GTGGGY; match class: 1
TF − Factor: Sp3; motif: ASMCTTGGGSRGGG; match class: 1
TF − Factor: MZF−1; motif: TGGGGAR; match class: 1
TF − Factor: GKLF; motif: NNNRGGNGNGGSN; match class: 1
TF − Factor: BTEB2; motif: GNAGGGGGNGGGSSNN
TF − Factor: MAZ; motif: GGGGGAGGGGGNGRGRRRGNRG
TF − Factor: RELA; motif: GGGRMTKYCCC
TF − Factor: GCMb:Max; motif: ATGCGGGYNNCAYGTG
TF − Factor: MAZ; motif: GGGGAGGG
TF − Factor: SALL2; motif: GGGTGGG
0
1
2
3
−log10(FDR)
−log10(p_value)
3.25
3.00
2.75
2.50