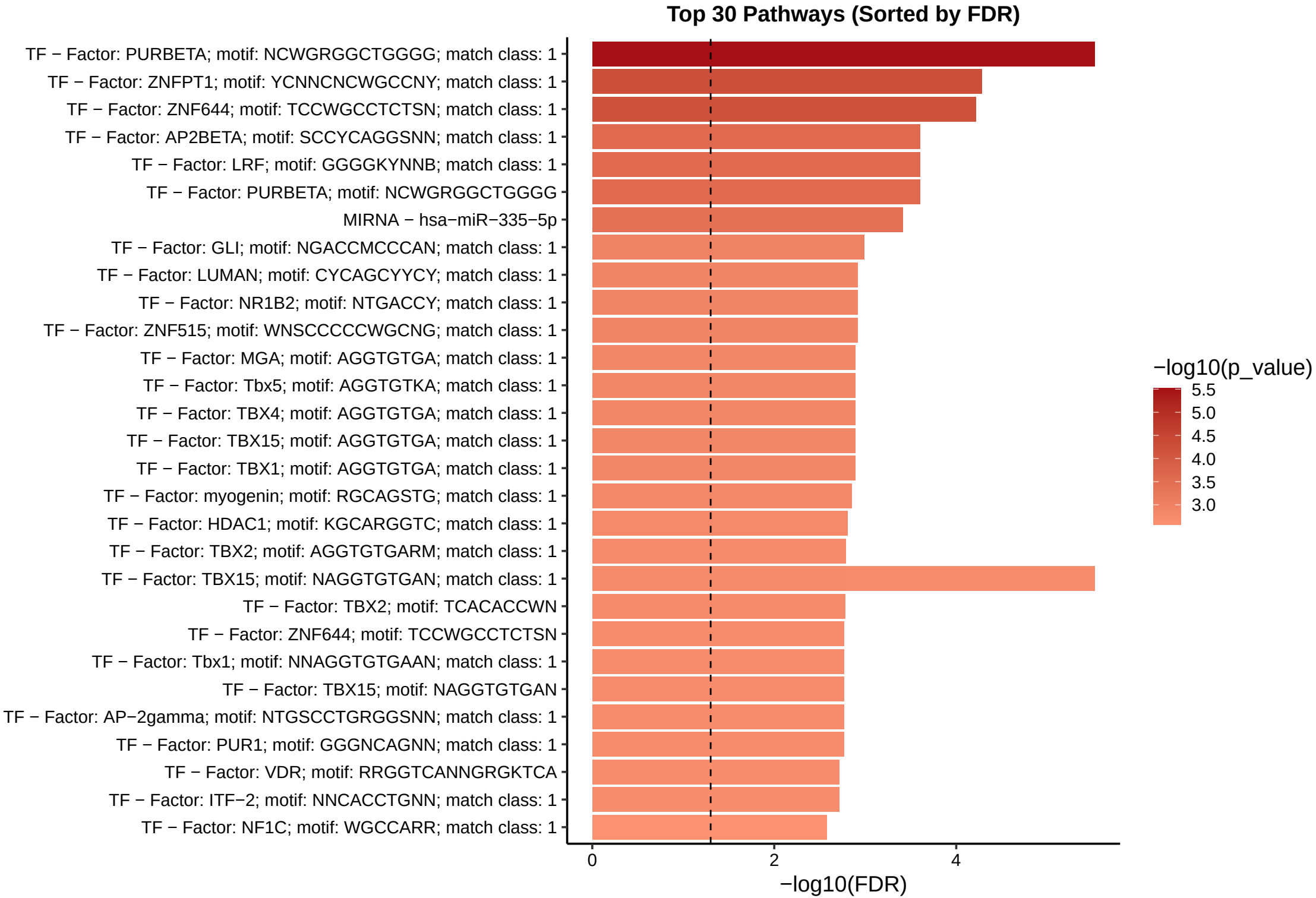
Top 30 Pathways (Sorted by FDR)
TF − Factor: PURBETA; motif: NCWGRGGCTGGGG; match class: 1
TF − Factor: ZNFPT1; motif: YCNNCNCWGCCNY; match class: 1
TF − Factor: ZNF644; motif: TCCWGCCTCTSN; match class: 1
TF − Factor: AP2BETA; motif: SCCYCAGGSNN; match class: 1
TF − Factor: LRF; motif: GGGGKYNNB; match class: 1
TF − Factor: PURBETA; motif: NCWGRGGCTGGGG
MIRNA − hsa−miR−335−5p
TF − Factor: GLI; motif: NGACCMCCCAN; match class: 1
TF − Factor: LUMAN; motif: CYCAGCYYCY; match class: 1
TF − Factor: NR1B2; motif: NTGACCY; match class: 1
TF − Factor: ZNF515; motif: WNSCCCCCWGCNG; match class: 1
TF − Factor: MGA; motif: AGGTGTGA; match class: 1
TF − Factor: Tbx5; motif: AGGTGTKA; match class: 1
TF − Factor: TBX4; motif: AGGTGTGA; match class: 1
TF − Factor: TBX15; motif: AGGTGTGA; match class: 1
TF − Factor: TBX1; motif: AGGTGTGA; match class: 1
TF − Factor: myogenin; motif: RGCAGSTG; match class: 1
TF − Factor: HDAC1; motif: KGCARGGTC; match class: 1
TF − Factor: TBX2; motif: AGGTGTGARM; match class: 1
TF − Factor: TBX15; motif: NAGGTGTGAN; match class: 1
TF − Factor: TBX2; motif: TCACACCWN
TF − Factor: ZNF644; motif: TCCWGCCTCTSN
TF − Factor: Tbx1; motif: NNAGGTGTGAAN; match class: 1
TF − Factor: TBX15; motif: NAGGTGTGAN
TF − Factor: AP−2gamma; motif: NTGSCCTGRGGSNN; match class: 1
TF − Factor: PUR1; motif: GGGNCAGNN; match class: 1
TF − Factor: VDR; motif: RRGGTCANNGRGKTCA
TF − Factor: ITF−2; motif: NNCACCTGNN; match class: 1
TF − Factor: NF1C; motif: WGCCARR; match class: 1
0
2
4
−log10(FDR)
−log10(p_value)
5.5
5.0
4.5
4.0
3.5
3.0