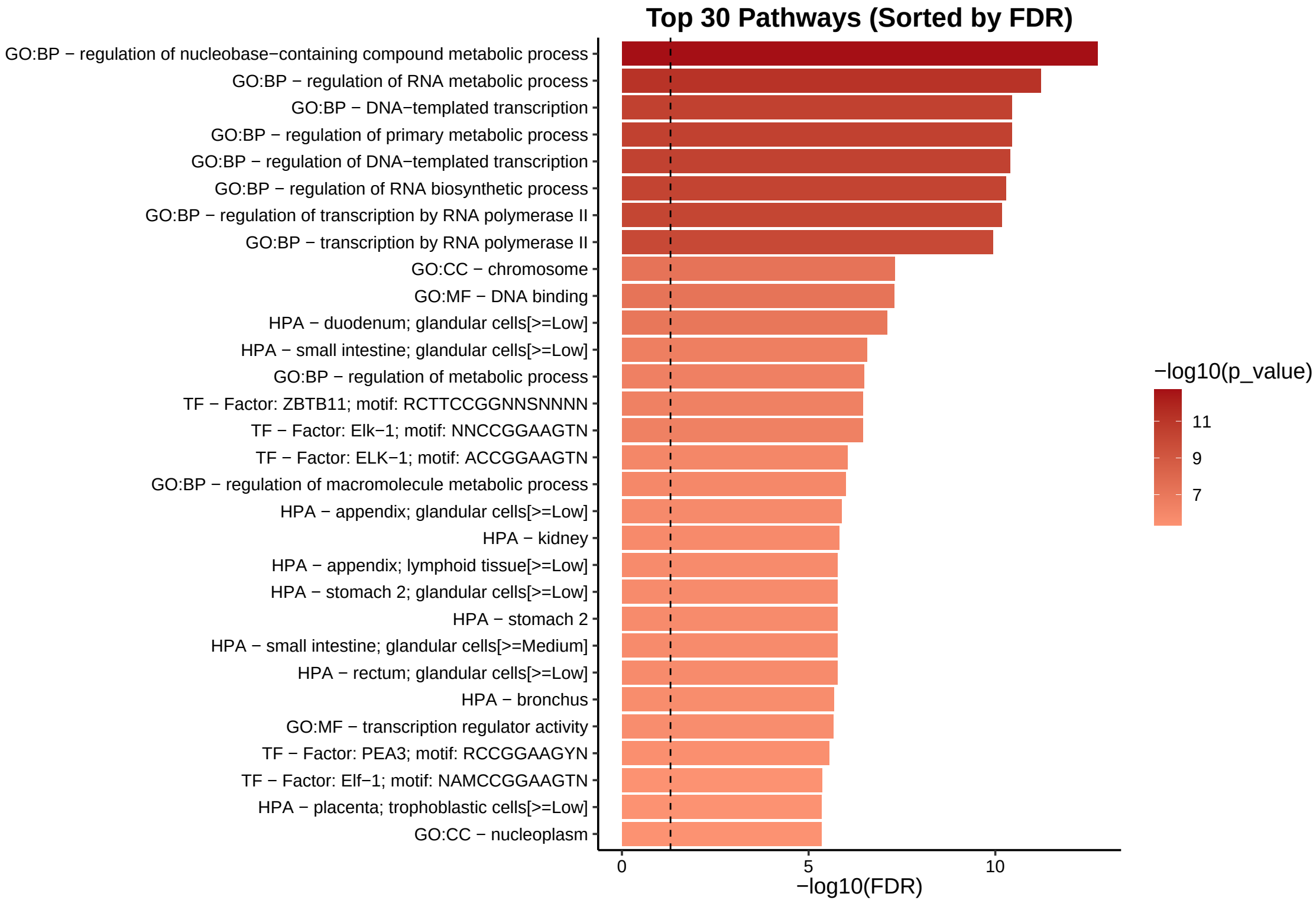
Top 30 Pathways (Sorted by FDR)
GO:BP − regulation of nucleobase−containing compound metabolic process
GO:BP − regulation of RNA metabolic process
GO:BP − DNA−templated transcription
GO:BP − regulation of primary metabolic process
GO:BP − regulation of DNA−templated transcription
GO:BP − regulation of RNA biosynthetic process
GO:BP − regulation of transcription by RNA polymerase II
GO:BP − transcription by RNA polymerase II
GO:CC − chromosome
GO:MF − DNA binding
HPA − duodenum; glandular cells[>=Low]
HPA − small intestine; glandular cells[>=Low]
GO:BP − regulation of metabolic process
TF − Factor: ZBTB11; motif: RCTTCCGGNNSNNNN
TF − Factor: Elk−1; motif: NNCCGGAAGTN
TF − Factor: ELK−1; motif: ACCGGAAGTN
GO:BP − regulation of macromolecule metabolic process
HPA − appendix; glandular cells[>=Low]
HPA − kidney
HPA − appendix; lymphoid tissue[>=Low]
HPA − stomach 2; glandular cells[>=Low]
HPA − stomach 2
HPA − small intestine; glandular cells[>=Medium]
HPA − rectum; glandular cells[>=Low]
HPA − bronchus
GO:MF − transcription regulator activity
TF − Factor: PEA3; motif: RCCGGAAGYN
TF − Factor: Elf−1; motif: NAMCCGGAAGTN
HPA − placenta; trophoblastic cells[>=Low]
GO:CC − nucleoplasm
0
5
10
−log10(FDR)
−log10(p_value)
11
9
7