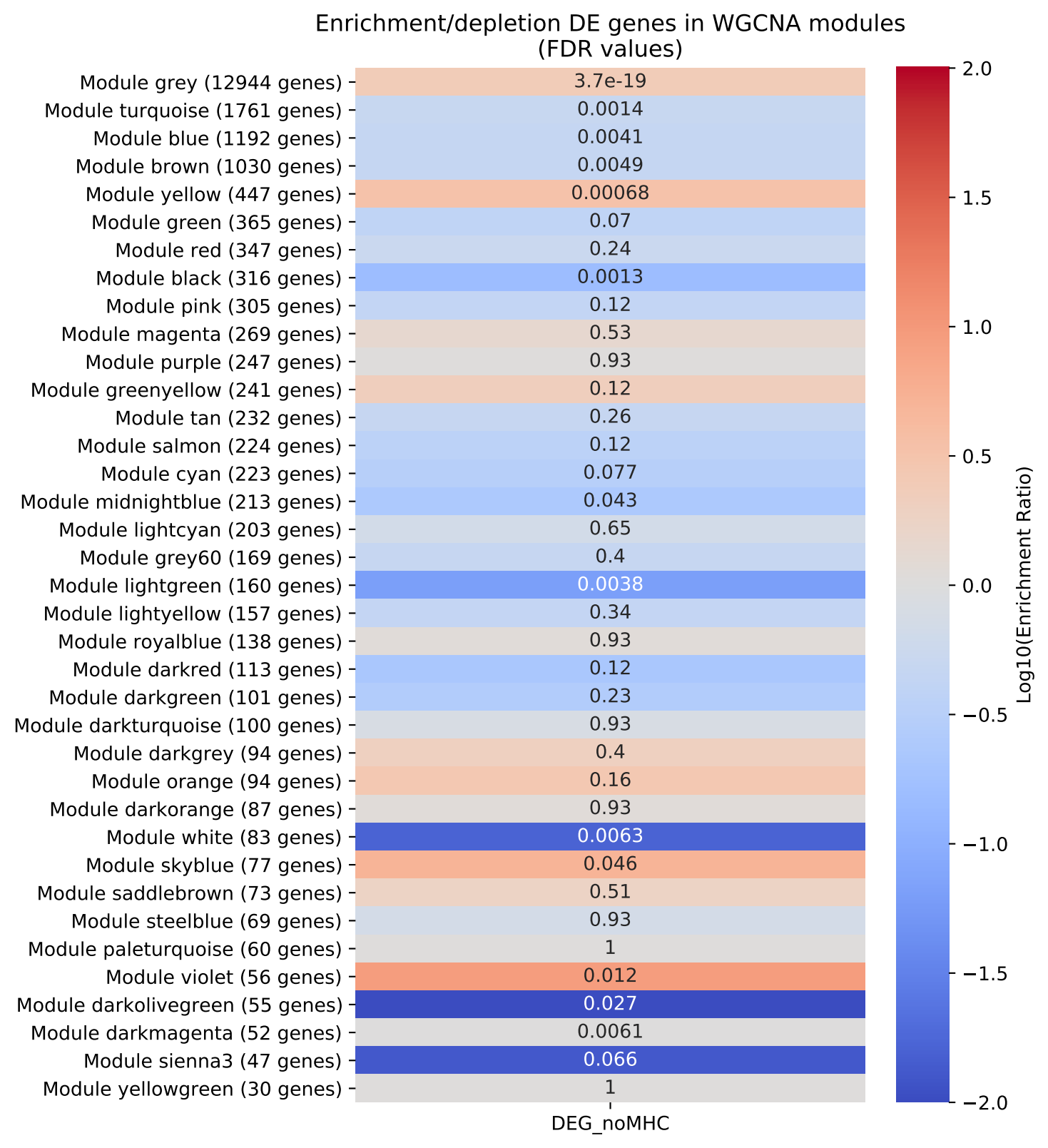
Enrichment/depletion DE genes in WGCNA modules
(FDR values)
Module grey (12944 genes)
Module turquoise (1761 genes)
Module blue (1192 genes)
Module brown (1030 genes)
Module yellow (447 genes)
Module green (365 genes)
Module red (347 genes)
Module black (316 genes)
Module pink (305 genes)
Module magenta (269 genes)
Module purple (247 genes)
Module greenyellow (241 genes)
Module tan (232 genes)
Module salmon (224 genes)
Module cyan (223 genes)
Module midnightblue (213 genes)
Module lightcyan (203 genes)
Module grey60 (169 genes)
Module lightgreen (160 genes)
Module lightyellow (157 genes)
Module royalblue (138 genes)
Module darkred (113 genes)
Module darkgreen (101 genes)
Module darkturquoise (100 genes)
Module darkgrey (94 genes)
Module orange (94 genes)
Module darkorange (87 genes)
Module white (83 genes)
Module skyblue (77 genes)
Module saddlebrown (73 genes)
Module steelblue (69 genes)
Module paleturquoise (60 genes)
Module violet (56 genes)
Module darkolivegreen (55 genes)
Module darkmagenta (52 genes)
Module sienna3 (47 genes)
Module yellowgreen (30 genes)
3.7e-19
0.0014
0.0041
0.0049
0.00068
0.07
0.24
0.0013
0.12
0.53
0.93
0.12
0.26
0.12
0.077
0.043
0.65
0.4
0.0038
0.34
0.93
0.12
0.23
0.93
0.4
0.16
0.93
0.0063
0.046
0.51
0.93
1
0.012
0.027
0.0061
0.066
1
DEG_noMHC
2.0
1.5
1.0
0.5
0.0
−0.5
−1.0
−1.5
−2.0
Log10(Enrichment Ratio)