Enrichment/depletion DE genes in WGCNA modules (FDR values) 2.0 0.0037 Module grey (6097 genes) -Module turquoise (2299 genes) -0.15 0.15 Module blue (1842 genes) -Module brown (1583 genes) -0.0065 Module yellow (1255 genes) -0.00099 1.5 Module green (1061 genes) -0.00014 Module red (906 genes) -0.0048 Module black (490 genes) -0.011 Module pink (489 genes) -0.0057 0.0013 Module magenta (473 genes) -- 1.0 0.046 Module purple (465 genes) -Module greenyellow (445 genes) -0.0057 0.37 Module tan (425 genes) -0.00023 Module salmon (363 genes) -0.085 Module cyan (317 genes) -- 0.5 Module midnightblue (260 genes) -0.016 og10(Enrichment Ratio 0.00044 Module lightcyan (227 genes) -5.4e-55 Module grey60 (190 genes) -Module lightgreen (182 genes) -0.12 Module lightyellow (156 genes) -0.016 - 0.0 0.085 Module royalblue (154 genes) -0.56 Module darkred (144 genes) -Module darkgreen (127 genes) -0.032 0.67 Module darkturquoise (119 genes) -0.0001 -0.5Module darkgrey (104 genes) -Module orange (97 genes) -0.088 Module darkorange (94 genes) -0.13 0.35 Module white (92 genes) -0.00023 Module skyblue (90 genes) --1.00.0037 Module steelblue (84 genes) -Module saddlebrown (84 genes) -1 Module paleturquoise (76 genes) -0.77 Module violet (63 genes) -0.0036 Module darkolivegreen (56 genes) --1.5Module darkmagenta (52 genes) -0.35 0.34 Module sienna3 (47 genes) -Module yellowgreen (46 genes) -0.48 Module skyblue3 (44 genes) -1 0.48 Module plum1 (42 genes) --2.0

DEG noMHC