T_cells_CD8 - IMM - Ileum - InactiveCD_vs_Control Down supramolecular fiber organization - GO:0097435 - p: 6.54e-05 *** negative regulation of programmed cell death – GO:0043069 – p: 7.54e–06 *** negative regulation of apoptotic process - GO:0043066 - p: 1.03e-05 *** multicellular organismal-level homeostasis - GO:0048871 - p: 1.55e-05 *** apoptotic signaling pathway - GO:0097190 - p: 2.01e-05 *** negative regulation of cell differentiation – GO:0045596 – p: 6.32e–05 *** positive regulation of cytokine production - GO:0001819 - p: 4.50e-05 *** positive regulation of cell differentiation – GO:0045597 – p: 1.69e–06 *** Gene Count regulation of cytokine production – GO:0001817 – p: 1.37e–06 *** cytokine production - GO:0001816 - p: 1.15e-06 *** intrinsic apoptotic signaling pathway - GO:0097193 - p: 2.99e-06 *** Cluster fat cell differentiation - GO:0045444 - p: 6.35e-07 *** response to hormone - GO:0009725 - p: 8.35e-08 *** response to lipid - GO:0033993 - p: 1.28e-07 *** cellular response to hormone stimulus - GO:0032870 - p: 6.55e-07 *** response to steroid hormone - GO:0048545 - p: 4.56e-06 *** cellular response to steroid hormone stimulus - GO:0071383 - p: 1.53e-05 *** cellular response to lipid - GO:0071396 - p: 1.48e-08 *** growth factor receptor binding - GO:0070851 - p: 7.73e-04 *** histone modifying activity - GO:0140993 - p: 9.89e-05 *** kinase binding - GO:0019900 - p: 1.08e-03 ** protein kinase binding - GO:0019901 - p: 1.91e-04 *** 1.6 1.8 2.0 Absolute NES