IgA_plasma_cell - IMM - Ileum - InactiveCD_vs_Control Down peptidyl-tyrosine phosphorylation - GO:0018108 - p: 5.45e-04 *** peptidyl-tyrosine modification - GO:0018212 - p: 3.31e-04 *** negative regulation of signaling – GO:0023057 – p: 2.12e–04 *** negative regulation of cell communication - GO:0010648 - p: 2.14e-04 *** negative regulation of signal transduction - GO:0009968 - p: 1.02e-04 *** negative regulation of multicellular organismal pr... - GO:0051241 - p: 5.58e-04 *** enzyme-linked receptor protein signaling pathway - GO:0007167 - p: 3.92e-04 *** regulation of multicellular organismal development - GO:2000026 - p: 5.25e-05 *** response to cytokine - GO:0034097 - p: 1.53e-04 *** Gene Count regulation of cell development - GO:0060284 - p: 4.07e-04 *** response to peptide - GO:1901652 - p: 3.96e-05 *** GO Term cell adhesion - GO:0007155 - p: 3.12e-05 *** Cluster negative regulation of cell differentiation - GO:0045596 - p: 3.66e-04 *** cellular response to cytokine stimulus - GO:0071345 - p: 6.77e-05 *** negative regulation of developmental process – GO:0051093 – p: 1.19e–04 *** cellular response to endogenous stimulus - GO:0071495 - p: 5.81e-06 *** response to hormone - GO:0009725 - p: 9.79e-06 *** response to endogenous stimulus – GO:0009719 – p: 7.64e–08 *** enzyme regulator activity - GO:0030234 - p: 4.07e-04 *** molecular function inhibitor activity - GO:0140678 - p: 1.43e-04 *** protein kinase binding - GO:0019901 - p: 6.16e-04 *** kinase binding - GO:0019900 - p: 1.56e-04 *** 1.7 1.5 1.8 1.9 Absolute NES