Immune_Cycling_cells - IMM - Colon - ActiveCD_vs_Control T cell activation - GO:0042110 - p: 5.56e-06 *** mononuclear cell differentiation - GO:1903131 - p: 9.10e-06 *** lymphocyte differentiation – GO:0030098 – p: 4.26e–07 *** T cell differentiation - GO:0030217 - p: 4.09e-07 *** chromatin remodeling - GO:0006338 - p: 1.00e-10 *** transcription regulator complex - GO:0005667 - p: 5.08e-04 *** CC perinuclear region of cytoplasm - GO:0048471 - p: 1.14e-06 *** Gene Count tumor necrosis factor receptor binding - GO:0005164 - p: 7.13e-04 *** phosphotransferase activity, alcohol group as acce... - GO:0016773 - p: 1.78e-04 *** protein kinase activity - GO:0004672 - p: 6.49e-05 *** GO Term Cluster DNA-binding transcription factor activity, RNA pol... - GO:0000981 - p: 4.69e-05 *** DNA-binding transcription factor activity - GO:0003700 - p: 2.25e-05 *** RNA polymerase II cis-regulatory region sequence-s... - GO:0000978 - p: 2.36e-05 *** cis-regulatory region sequence-specific DNA bindin... - GO:0000987 - p: 2.36e-05 *** DNA-binding transcription factor binding - GO:0140297 - p: 2.19e-04 *** transcription factor binding - GO:0008134 - p: 1.51e-06 *** RNA polymerase II-specific DNA-binding transcripti... - GO:0061629 - p: 3.56e-05 *** nuclear receptor binding - GO:0016922 - p: 3.42e-04 *** chromatin binding - GO:0003682 - p: 1.00e-10 *** histone binding - GO:0042393 - p: 9.92e-08 *** histone modifying activity - GO:0140993 - p: 2.36e-07 *** 1.4 1.5 1.6 1.7 1.8 Absolute NES