crypt - EPI - Ileum - InactiveCD_vs_Control Up external encapsulating structure organization - GO:0045229 - p: 1.43e-04 ** extracellular structure organization - GO:0043062 - p: 1.43e-04 *** extracellular matrix organization - GO:0030198 - p: 1.43e-04 *** recombinational repair - GO:0000725 - p: 9.93e-05 ** regulation of T cell mediated cytotoxicity - GO:0001914 - p: 7.11e-05 *** animal organ morphogenesis - GO:0009887 - p: 7.19e-05 ** pattern specification process - GO:0007389 - p: 4.66e-04 ** regionalization - GO:0003002 - p: 2.43e-04 ** connective tissue development - GO:0061448 - p: 1.58e-05 *** 6 bone morphogenesis - GO:0060349 - p: 2.32e-04 *** ureteric bud development - GO:0001657 - p: 2.59e-04 ** 6 epidermis development - GO:0008544 - p: 1.99e-04 *** 4 keratinocyte differentiation - GO:0030216 - p: 2.03e-04 *** epidermal cell differentiation - GO:0009913 - p: 1.27e-05 *** regulation of epidermal cell differentiation – GO:0045604 – p: 2.02e-05 ** 4 4 regulation of epidermis development - GO:0045682 - p: 1.19e-05 *** circadian rhythm - GO:0007623 - p: 1.17e-04 ** rhythmic process - GO:0048511 - p: 1.06e-05 *** mitotic cell cycle process - GO:1903047 - p: 1.73e-04 ** Gene Count chromosome organization - GO:0051276 - p: 9.65e-05 ** chromosome segregation - GO:0007059 - p: 9.71e-05 ** nuclear chromosome segregation - GO:0098813 - p: 5.75e-05 ** sister chromatid segregation - GO:0000819 - p: 1.26e-04 *** organelle fission - GO:0048285 - p: 4.62e-06 *** mitotic sister chromatid segregation - GO:0000070 - p: 1.85e-04 *** 뭐 nuclear division - GO:0000280 - p: 3.03e-06 * regulation of mitotic nuclear division - GO:0007088 - p: 5.19e-04 *** mitotic nuclear division - GO:0140014 - p: 2.19e-06 ** Cluster regulation of nuclear division - GO:0051783 - p: 2.32e-04 *** positive regulation of nuclear division - GO:0051785 - p: 1.64e-04 ** immune response-activating signaling pathway - GO:0002757 - p: 4.84e-04 *** response to bacterium - GO:0009617 - p: 2.84e-04 *** regulation of immune response - GO:0050776 - p: 3.20e-05 ** innate immune response - GO:0045087 - p: 1.07e-05 *** activation of immune response - GO:0002253 - p: 2.73e-05 ** positive regulation of immune response - GO:0050778 - p: 2.17e-06 *** immune response-regulating cell surface receptor s... - GO:0002768 - p: 1.85e-04 ** immune effector process - GO:0002252 - p: 3.46e-06 *** immune response-activating cell surface receptor s... - GO:0002429 - p: 9.71e-05 *** defense response to bacterium - GO:0042742 - p: 2.21e-04 *** T cell receptor signaling pathway - GO:0050852 - p: 5.16e-04 *** leukocyte mediated immunity - GO:0002443 - p: 2.22e-06 *** adaptive immune response based on somatic recombin... - GO:0002460 - p: 5.97e-06 ** antigen receptor-mediated signaling pathway - GO:0050851 - p: 3.56e-05 * B cell mediated immunity - GO:0019724 - p: 1.31e-04 *** immunoglobulin mediated immune response - GO:0016064 - p: 1.31e-04 *** adaptive immune response - GO:0002250 - p: 2.85e-07 *** complement activation, classical pathway - GO:0006958 - p: 3.77e-04 ** complement activation - GO:0006956 - p: 2.38e-04 *** lymphocyte mediated immunity - GO:0002449 - p: 7.01e-08 * actin cytoskeleton - GO:0015629 - p: 5.50e-05 *** CC phagocytic vesicle - GO:0045335 - p: 1.24e-04 ** blood microparticle - GO:0072562 - p: 2.01e-05 ** DNA-binding transcription activator activity - GO:0001216 - p: 1.97e-04 ** DNA-binding transcription activator activity, RNA ... - GO:0001228 - p: 2.01e-04 *** transcription coregulator activity - GO:0003712 - p: 1.26e-04 *** transcription corepressor activity - GO:0003714 - p: 4.31e-05 *** transcription regulator inhibitor activity - GO:0140416 - p: 2.08e-04 *** 0 antigen binding - GO:0003823 - p: 4.71e-07 ***

2.2

Absolute NES

2.0

1.6

2.0

2.2

1.8

1.6

1.8

GO Term