Mature_Goblet - EPI - Ileum - InactiveCD_vs_Control Up 6 mucosal immune response - GO:0002385 - p: 2.93e-04 *** nucleobase-containing small molecule metabolic pro... - GO:0055086 - p: 2.48e-04 * nucleoside phosphate metabolic process - GO:0006753 - p: 3.11e-04 * purine-containing compound metabolic process - GO:0072521 - p: 1.96e-04 *** nucleotide metabolic process - GO:0009117 - p: 7.18e-05 *** small molecule catabolic process - GO:0044282 - p: 2.42e-04 ** regulation of cell adhesion - GO:0030155 - p: 1.61e-04 ** regulation of cell-cell adhesion - GO:0022407 - p: 1.07e-04 *** leukocyte cell-cell adhesion - GO:0007159 - p: 1.11e-04 ** regulation of leukocyte cell-cell adhesion - GO:1903037 - p: 3.48e-05 * 2 acylglycerol homeostasis - GO:0055090 - p: 1.29e-05 ** triglyceride homeostasis - GO:0070328 - p: 2.50e-06 ** regulation of immune response - GO:0050776 - p: 3.23e-04 * positive regulation of immune response - GO:0050778 - p: 3.05e-04 ** response to bacterium - GO:0009617 - p: 9.70e-05 ** immune effector process - GO:0002252 - p: 1.82e-06 ** ВР regulation of leukocyte mediated immunity - GO:0002703 - p: 1.98e-04 ** Gene Count cell killing - GO:0001906 - p: 1.39e-04 ** defense response to bacterium - GO:0042742 - p: 1.23e-05 ** regulation of respiratory burst - GO:0060263 - p: 6.64e-05 ** adaptive immune response - GO:0002250 - p: 2.16e-09 * antimicrobial humoral response - GO:0019730 - p: 6.82e-05 ** leukocyte mediated immunity - GO:0002443 - p: 6.41e-09 ** humoral immune response - GO:0006959 - p: 1.00e-06 ** complement activation, classical pathway - GO:0006958 - p: 5.99e-05 ** 0 Cluster adaptive immune response based on somatic recombin... - GO:0002460 - p: 4.64e-08 * 0 humoral immune response mediated by circulating im... - GO:0002455 - p: 1.03e-04 ** antibacterial humoral response - GO:0019731 - p: 2.77e-05 *** lymphocyte mediated immunity - GO:0002449 - p: 8.79e-09 * 0 complement activation - GO:0006956 - p: 6.10e-06 ** B cell mediated immunity - GO:0019724 - p: 6.61e-08 ** 1 immunoglobulin mediated immune response - GO:0016064 - p: 1.43e-08 ** postsynaptic specialization membrane - GO:0099634 - p: 1.08e-03 spliceosomal complex - GO:0005681 - p: 7.85e-04 * mitochondrial membrane - GO:0031966 - p: 3.14e-04 *** mitochondrial matrix - GO:0005759 - p: 2.55e-04 ** organelle inner membrane - GO:0019866 - p: 3.73e-05 * ribosomal subunit - GO:0044391 - p: 8.63e-04 * ribosome - GO:0005840 - p: 2.11e-04 ** mitochondrial protein-containing complex - GO:0098798 - p: 3.14e-04 ** mitochondrial inner membrane - GO:0005743 - p: 4.04e-06 * mitochondrial ribosome - GO:0005761 - p: 3.73e-04 ** organellar ribosome - GO:0000313 - p: 3.73e-04 ** small ribosomal subunit - GO:0015935 - p: 4.61e-04 *** mitochondrial small ribosomal subunit - GO:0005763 - p: 3.95e-04 ** 6 0 organellar small ribosomal subunit - GO:0000314 - p: 3.95e-04 ** lumenal side of membrane - GO:0098576 - p: 1.14e-03 lumenal side of endoplasmic reticulum membrane - GO:0098553 - p: 8.67e-04 ** MHC protein complex - GO:0042611 - p: 8.14e-04 ** blood microparticle - GO:0072562 - p: 5.08e-06 * antigen binding - GO:0003823 - p: 1.15e-08 ***

2.6 1.4

Absolute NES

1.6

1.8

2.0

2.2

2.4

2.6

1.4

1.6

2.0

2.2

2.4

Terr

.09