T_cells_OGT - IMM - Ileum - InactiveCD_vs_Control Up gamma-tubulin complex - GO:0000930 - p: 1.76e-03 ** mitochondrial protein-containing complex - GO:0098798 - p: 1.46e-03 ** Cul3-RING ubiquitin ligase complex - GO:0031463 - p: 4.83e-04 ** 0 translation preinitiation complex - GO:0070993 - p: 1.26e-03 ** eukaryotic translation initiation factor 3 complex - GO:0005852 - p: 1.81e-03 ** eukaryotic 48S preinitiation complex - GO:0033290 - p: 4.95e-04 *** eukaryotic 43S preinitiation complex - GO:0016282 - p: 3.99e-04 *** 6 synaptic membrane - GO:0097060 - p: 1.92e-03 ** postsynaptic membrane - GO:0045211 - p: 2.34e-04 *** calcium channel complex - GO:0034704 - p: 1.07e-04 *** oxidoreductase complex - GO:1990204 - p: 6.86e-04 *** respiratory chain complex - GO:0098803 - p: 5.78e-04 ** Gene Count transmembrane transporter complex - GO:1902495 - p: 7.26e-05 *** small ribosomal subunit - GO:0015935 - p: 6.27e-08 *** cytosolic small ribosomal subunit - GO:0022627 - p: 2.64e-07 *** cytosolic ribosome - GO:0022626 - p: 1.00e-10 *** ribosomal subunit - GO:0044391 - p: 1.00e-10 *** large ribosomal subunit - GO:0015934 - p: 1.00e-10 *** Cluster cytosolic large ribosomal subunit - GO:0022625 - p: 1.00e-10 *** acyltransferase activity - GO:0016746 - p: 1.52e-03 ** molecular tag activity - GO:0141047 - p: 1.31e-03 * protein tag activity - GO:0031386 - p: 1.31e-03 ** transcription regulator activity - GO:0140110 - p: 3.03e-04 *** double-stranded DNA binding - GO:0003690 - p: 4.40e-04 *** sequence-specific DNA binding - GO:0043565 - p: 1.92e-04 *** sequence-specific double-stranded DNA binding - GO:1990837 - p: 3.17e-04 *** transcription regulatory region nucleic acid bindi... - GO:0001067 - p: 2.21e-04 *** transcription cis-regulatory region binding - GO:0000976 - p: 2.21e-04 *** cis-regulatory region sequence-specific DNA bindin... - GO:0000987 - p: 2.80e-04 *** ₹ RNA polymerase II cis-regulatory region sequence-s... - GO:0000978 - p: 2.44e-04 *** RNA polymerase II transcription regulatory region ... - GO:0000977 - p: 7.39e-05 *** DNA-binding transcription activator activity, RNA ... - GO:0001228 - p: 1.01e-03 ** DNA-binding transcription activator activity - GO:0001216 - p: 1.01e-03 ** DNA-binding transcription factor activity - GO:0003700 - p: 1.81e-06 *** DNA-binding transcription factor activity, RNA pol... - GO:0000981 - p: 1.38e-06 *** ATP-dependent protein folding chaperone - GO:0140662 - p: 2.33e-05 *** protein folding chaperone - GO:0044183 - p: 6.40e-07 *** electron transfer activity - GO:0009055 - p: 1.12e-03 ** rRNA binding - GO:0019843 - p: 6.49e-04 *** structural constituent of ribosome - GO:0003735 - p: 1.00e-10 ***

Absolute NES

GO Term