Macrophages_LYVE1 - 06_TI_IMM_ActiveCD_vs_Control Up adenylate cyclase-activating G protein-coupled rec... - GO:0007189 - p-val: 1.31e-04 *** (3) regulation of immune system process - GO:0002682 - p-val: 6.82e-05 *** innate immune response - GO:0045087 - p-val: 1.24e-04 *** intracellular monoatomic ion homeostasis - GO:0006873 - p-val: 8.15e-05 *** 2 2 monoatomic ion homeostasis - GO:0050801 - p-val: 3.14e-05 *** 2 intracellular monoatomic cation homeostasis - GO:0030003 - p-val: 2.51e-05 *** 뭐 2 monoatomic cation homeostasis - GO:0055080 - p-val: 3.36e-05 *** leukocyte mediated cytotoxicity - GO:0001909 - p-val: 1.32e-04 *** 2 cell killing - GO:0001906 - p-val: 1.22e-06 *** 2 cytoplasmic translation - GO:0002181 - p-val: 6.36e-08 *** ribosomal large subunit biogenesis - GO:0042273 - p-val: 8.48e-06 *** 0 extracellular matrix - GO:0031012 - p-val: 5.56e-03 ** **(5)** external encapsulating structure - GO:0030312 - p-val: 5.56e-03 ** **(5)** intermediate filament cytoskeleton - GO:0045111 - p-val: 3.43e-03 ** 4 organelle subcompartment - GO:0031984 - p-val: 5.32e-03 ** Golgi apparatus - GO:0005794 - p-val: 3.39e-03 ** nuclear outer membrane-endoplasmic reticulum membr... - GO:0042175 - p-val: 3.06e-03 ** endoplasmic reticulum subcompartment - GO:0098827 - p-val: 1.39e-03 ** endoplasmic reticulum membrane - GO:0005789 - p-val: 1.21e-03 ** endosome - GO:0005768 - p-val: 2.49e-04 *** endosome membrane - GO:0010008 - p-val: 1.71e-03 ** lysosome - GO:0005764 - p-val: 2.45e-04 *** Cluster lytic vacuole - GO:0000323 - p-val: 2.45e-04 *** vacuole - GO:0005773 - p-val: 1.17e-04 *** transmembrane transporter complex - GO:1902495 - p-val: 3.60e-03 ** secretory granule - GO:0030141 - p-val: 2.95e-05 *** cell surface - GO:0009986 - p-val: 4.68e-04 *** external side of plasma membrane - GO:0009897 - p-val: 4.83e-03 ** secretory vesicle - GO:0099503 - p-val: 4.32e-06 *** GO Term coated vesicle - GO:0030135 - p-val: 2.05e-03 ** azurophil granule - GO:0042582 - p-val: 5.36e-03 ** 2 Gene Count primary lysosome - GO:0005766 - p-val: 5.36e-03 ** 2 coated vesicle membrane - GO:0030662 - p-val: 3.57e-03 ** 2 clathrin-coated vesicle - GO:0030136 - p-val: 2.19e-03 ** 2 specific granule - GO:0042581 - p-val: 3.88e-03 ** 2 vesicle membrane - GO:0012506 - p-val: 4.15e-08 *** cytoplasmic vesicle membrane - GO:0030659 - p-val: 2.69e-08 *** 2 clathrin-coated vesicle membrane - GO:0030665 - p-val: 3.63e-03 ** 2 vacuolar lumen - GO:0005775 - p-val: 2.10e-03 ** tertiary granule membrane - GO:0070821 - p-val: 3.89e-03 ** 2 2 secretory granule membrane - GO:0030667 - p-val: 8.73e-05 *** proton-transporting two-sector ATPase complex - GO:0016469 - p-val: 5.45e-03 ** 2 lysosomal lumen - GO:0043202 - p-val: 2.44e-03 ** 2 specific granule membrane - GO:0035579 - p-val: 1.27e-03 ** 2 tertiary granule - GO:0070820 - p-val: 3.45e-04 *** 2 endocytic vesicle membrane - GO:0030666 - p-val: 1.56e-04 *** 2 proton-transporting V-type ATPase complex - GO:0033176 - p-val: 1.84e-03 ** 0 phagocytic vesicle membrane - GO:0030670 - p-val: 7.46e-04 *** 2 ATPase dependent transmembrane transport complex - GO:0098533 - p-val: 1.11e-03 ** cation-transporting ATPase complex - GO:0090533 - p-val: 1.11e-03 ** vacuolar proton-transporting V-type ATPase complex - GO:0016471 - p-val: 1.66e-03 ** ribosome - GO:0005840 - p-val: 3.17e-04 *** ribosomal subunit - GO:0044391 - p-val: 3.76e-05 *** cytosolic small ribosomal subunit - GO:0022627 - p-val: 3.60e-03 ** 1 large ribosomal subunit - GO:0015934 - p-val: 1.44e-04 *** cytosolic ribosome - GO:0022626 - p-val: 2.39e-09 *** cytosolic large ribosomal subunit - GO:0022625 - p-val: 7.87e-10 *** 1 1 structural constituent of ribosome - GO:0003735 - p-val: 6.75e-06 *** ≤ rRNA binding - GO:0019843 - p-val: 1.14e-04 *** 1 Absolute NES

100

200

300

400