Job ID: 20200305-public-4.0.4-nuGxkz Display name: E2.bed **GO Biological Process** -log10(Binomial p value) 10 12 14 16 18 20 regulation of release of cytochrome c from mitochondria regulation of regulatory T cell differentiation regulation of tumor necrosis factor biosynthetic process response to arsenic-containing substance cytoplasmic translation tRNA transport response to muramyl dipeptide intrinsic apoptotic signaling pathway in response to DNA damage by p53 class mediator regulation of interferon-gamma-mediated signaling pathway negative regulation of B cell proliferation negative regulation of lipid storage T cell homeostasis 'de novo' protein folding nucleolytic nuclear-transcribed mRNA catabolic process involved in deadenylation-dependent decay snRNA processing nuclear-transcribed mRNA catabolic process, exonucleolytic 'de novo' posttranslational protein folding positive regulation of vacuole organization cytokinetic process intra-S DNA damage checkpoint