T cell Differential Expression negative regulation of miRNA processing positive regulation of execution phase of apoptosis negative regulation of miRNA-mediated gene silencing negative regulation of post-transcriptional gene silencing by regulatory ncRNA negative regulation of post-transcriptional gene silencing negative regulation of gene silencing by regulatory ncRNA regulation of miRNA processing RichFactor regulation of regulatory ncRNA processing 0.200 glial cell apoptotic process **Description** tryptophan catabolic process 0.175 retrograde trans-synaptic signaling 0.150 regulation of sensory perception of pain regulation of sensory perception 0.125 regulation of deoxyribonuclease activity 0.100 positive regulation of peroxisome proliferator activated receptor signaling pathway indole-containing compound catabolic process glucosylceramide catabolic process free ubiquitin chain polymerization ERBB3 signaling pathway bone marrow development adenosine metabolic process 0.000.050.100.150.20

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