

hIP\_split\_rna\_cutoffproximal.active.accUp.rnaDown-proximal\_2500bp-true\_GO\_BP



homeostasis myeloid neuron leukocyte  
regulation cell positive differentiation negative development

**Similarity**  
0.8  
0.6  
0.4  
0.2  
0

compound macromolecule rna transcription nitrogen  
process biosynthetic regulation metabolic cellular