with_ChIP_proximal.inactive.accUp.H3K27me3_GO_BP

Similarity
0.5
0.4
0.3
0.2
0.1
0

differentiation cell involved process tubule metanephric formation
development morphogenesis regulation

positive compound macromolecule transcription rna cellular
process regulation metabolic biosynthetic

mechanical peroxide hydrogen acid
response cellular stimulus hexose monosaccharide glucose

cohesion division segregation chromosome maintenance
regulation positive sister chromatid mitotic

cellular apoptotic negative cytolysis
regulation cell process proliferation positive biological