

detected-with_ChIP_rna_cutoff_bg_all_proximal.inactive.accUp.H3K27me3_GO_BP



differentiation cell formation involved tubule process metanephric
development morphogenesis regulation

Similarity
0.8
0.6
0.4
0.2
0

positive compound macromolecule transcription rna cellular
regulation process metabolic biosynthetic

cohesion division segregation chromosome maintenance
regulation positive sister chromatid mitotic

glucose peroxide hydrogen
response cellular acid stimulus retinoic monosaccharide hexose

mesenchymal cell apoptotic cytolysis
regulation process negative positive morphogenesis epithelium