

n_ChIP_split_bg_all_proximal.inactive.accUp.H3K27me3-proximal_2500bp_GO_BP



differentiation cell involved process tubule formation metanephric
development morphogenesis regulation

Similarity
0.5
0.4
0.3
0.2
0.1
0

positive compound macromolecule transcription rna cellular
process regulation metabolic biosynthetic

cohesion division segregation chromosome maintenance
regulation positive sister chromatid mitotic

glucose peroxide hydrogen acid
response cellular stimulus retinoic monosaccharide hexose

mesenchymal apoptotic negative cytolysis
regulation positive process cell cellular epithelium