

olig_rna_bg_all_proximal.inactive.accUp.H3K27me3-proximal_2500bp-not_detected_GO_BP



differentiation cell involved tubule process formation metanephric
development regulation morphogenesis

Similarity
0.8
0.6
0.4
0.2
0

compound positive cellular macromolecule transcription rna
process regulation biosynthetic metabolic

cohesion division segregation chromosome maintenance
regulation positive sister chromatid mitotic

glucose peroxide hydrogen
response cellular acid stimulus retinoic monosaccharide hexose