

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