

rna\_cutoff\_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