

ot\_detected-with\_ChIP\_rna\_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