

detected-with\_ChIP\_rna\_cutoff\_bg\_all\_proximal.inactive.accUp.H3K27me3\_GO\_BP



differentiation cell formation involved process tubule 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

glucose peroxide hydrogen acid  
**response** cellular stimulus retinoic monosaccharide hexose

cohesion division segregation chromosome maintenance  
**regulation** positive sister chromatid mitotic

response dna damage cellular  
**regulation** positive cell process proliferation stimulus