with\_ChIP\_bg\_all\_proximal.inactive.accUp.H3K27me3\_GO\_BP



differentiation cell formation involved tubule process metanephric development morphogenesis regulation



**Similarity** 8.0 0.6 0.4

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