with\_ChIP\_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 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

regulation positive cell process proliferation stimulus