with_ChIP_bg_all_proximal.inactive.accUp.H3K27me3_GO_BP metanephric involved tubule epithelial kidney cell nephron development morphogenesis regulation differentiation embryonic cell fate eye cameratype formation development morphogenesis **Similarity** 8.0 0.6 0.4 0.2 positive cellular compound rna macromolecule transcription process regulation metabolic biosynthetic cohesion division segregation chromosome maintenance regulation positive sister chromatid mitotic response cellular acid stimulus retinoic monosaccharide hexose **Nerve** optic vestibulocochlear formation development cranial