## GO Enrichment Analysis: Molecular function mRNA binding transcription coregulator activity GeneRatio 0.02 ubiquitin protein ligase binding 0.03 0.04 0.05 0.06 mRNA 3'-UTR binding p.adjust 0.03 0.02 ubiquitin-like protein ligase binding 0.01 TAP binding peptide antigen binding up (653) down (255)Cluster