GO Enrichment Analysis: Biological processes peptidyl-serine dephosphorylation positive regulation of supramolecular fiber organization actin polymerization or depolymerization regulation of supramolecular fiber organization GeneRatio regulation of actin filament organization 0.04 0.08 0.12 RNA splicing 0.16 regulation of RNA splicing p.adjust proteasome-mediated 0.04 ubiquitin-dependent protein catabolic process 0.03 0.02 mRNA processing 0.01 regulation of mRNA metabolic process negative regulation of cell migration negative regulation of locomotion negative regulation of cell motility DEG & DEtxs DEG only DEE only (37)(661)(213)Cluster