Gene ontology in rosy module with 504 genes actomyosin structure organization (GO:0031032) Hits anterograde trans-synaptic signaling (GO:0098916)

chemical synaptic transmission (GO:0007268) regulation of relaxation of cardiac muscle (GO:1901897) regulation of catecholamine secretion (GO:0050433) Combined regulation of regulated secretory pathway (GO:1903305) Score axon guidance (GO:0007411)

protein phosphorvlation (GO:0006468) potassium ion transmembrane transport (GO:0071805) 1.5 2.0 2.5 3.0 3.5 4.0

- 750 - 500 regulation of calcium ion-dependent exocytosis (GO:0017158) -log<sub>10</sub>(Adjusted P-value)