Gene ontology in darkred module with 381 genes ubiquitin-dependent protein catabolic process (GO:0006511) Hits

regulation of sodium ion transport (GO:0002028) 18 8 regulation of neuron differentiation (GO:0045664)

axon development (GO:0061564)

negative regulation of apoptotic signaling pathway (GO:2001234) Combined positive regulation of protein localization to membrane (GO:1905477) 

Score cellular response to ketone (GO:1901655)

response to forskolin (GO:1904321)

- 150

modification-dependent protein catabolic process (GO:0019941) - 50 myelination (GO:0042552)

0.80 0.82 0.84 0.86 0.88

-log<sub>10</sub>(Adjusted P-value)