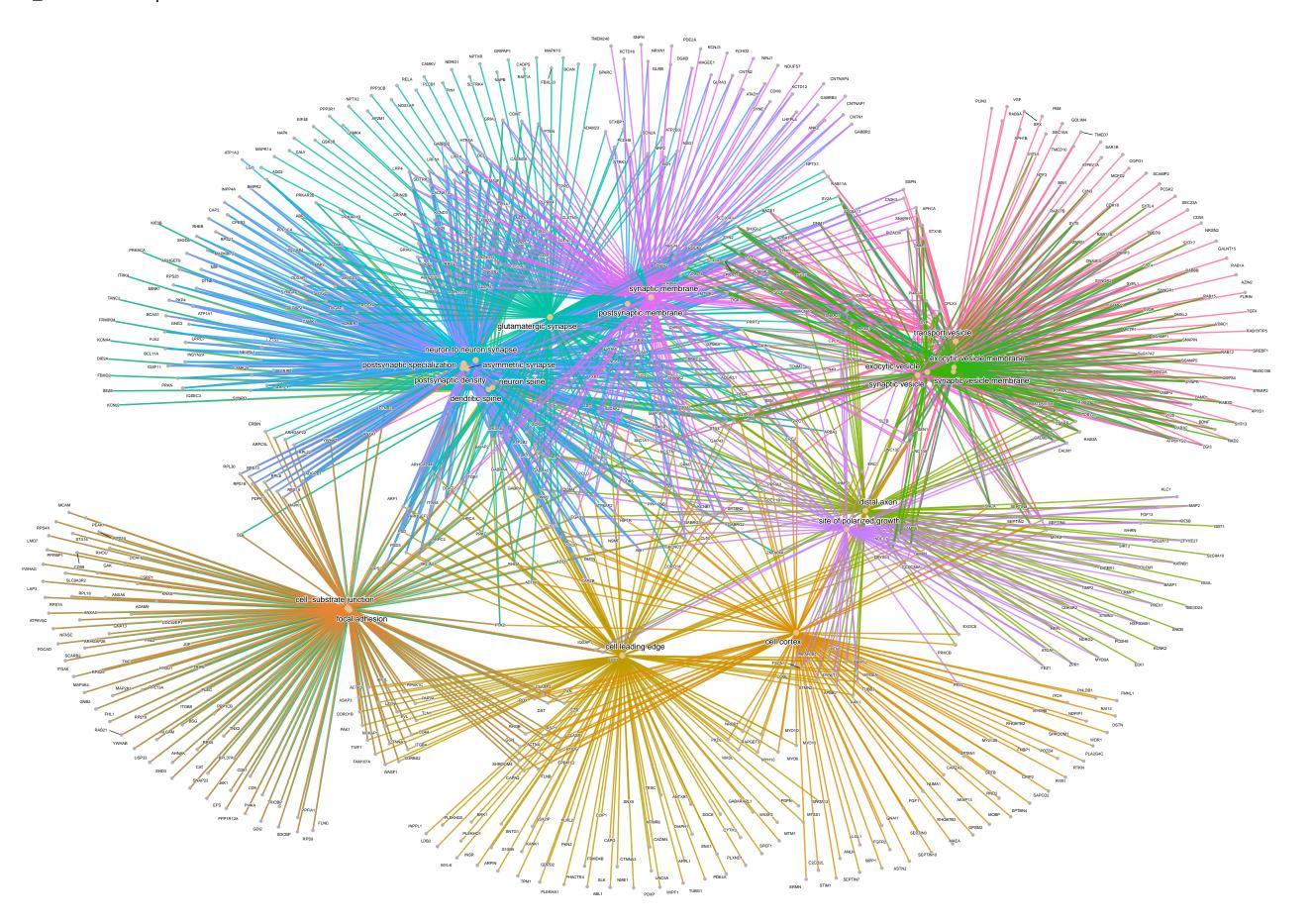


cluster_2 Cellular Compartment Network



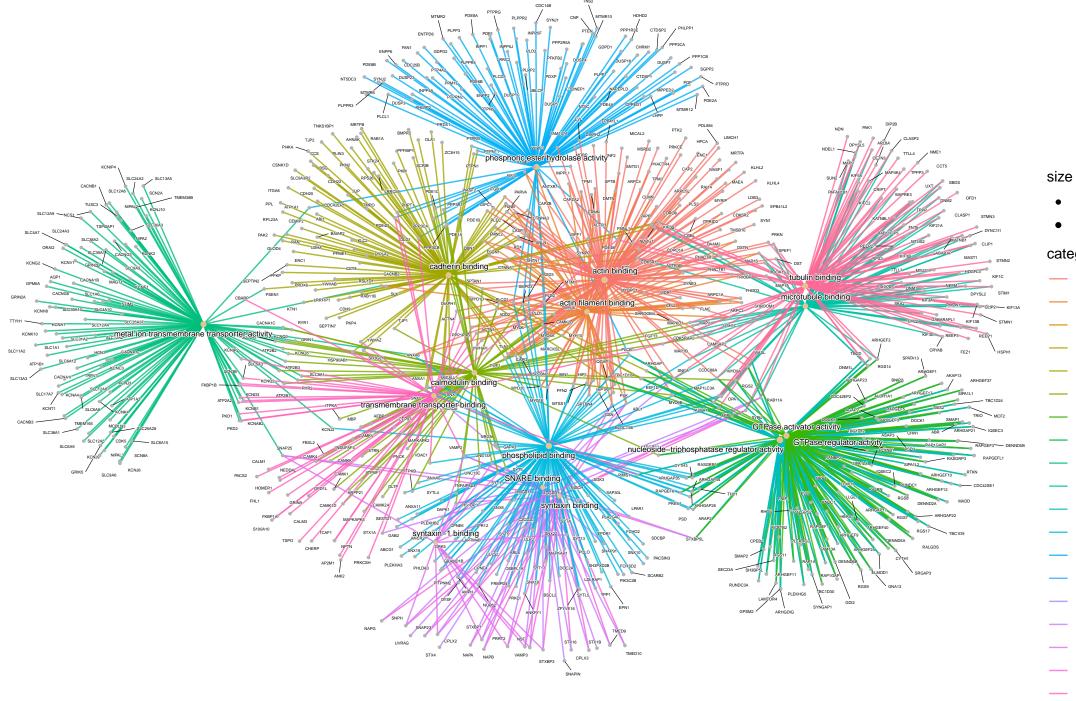
category

- asymmetric synapse
- cell-substrate junction
- cell cortex
- cell leading edge
- dendritic spine
- distal axon
- exocytic vesicle
- exocytic vesicle membrane
- focal adhesion
- glutamatergic synapse
- neuron spine
- neuron to neuron synapse
- postsynaptic density
- postsynaptic membrane
- postsynaptic specialization
- site of polarized growth
- synaptic membrane
- synaptic vesicle
 - o)..ap...o ..o....
- synaptic vesicle membrane
- transport vesicle

size

- 50
- 100
- 150

cluster_2 Molecular Function Network



ATPase activity, coupled to transmembrane movement of ions, rotational mechanism

ATPAT Pase=coupled ion transmembrane transporter activity

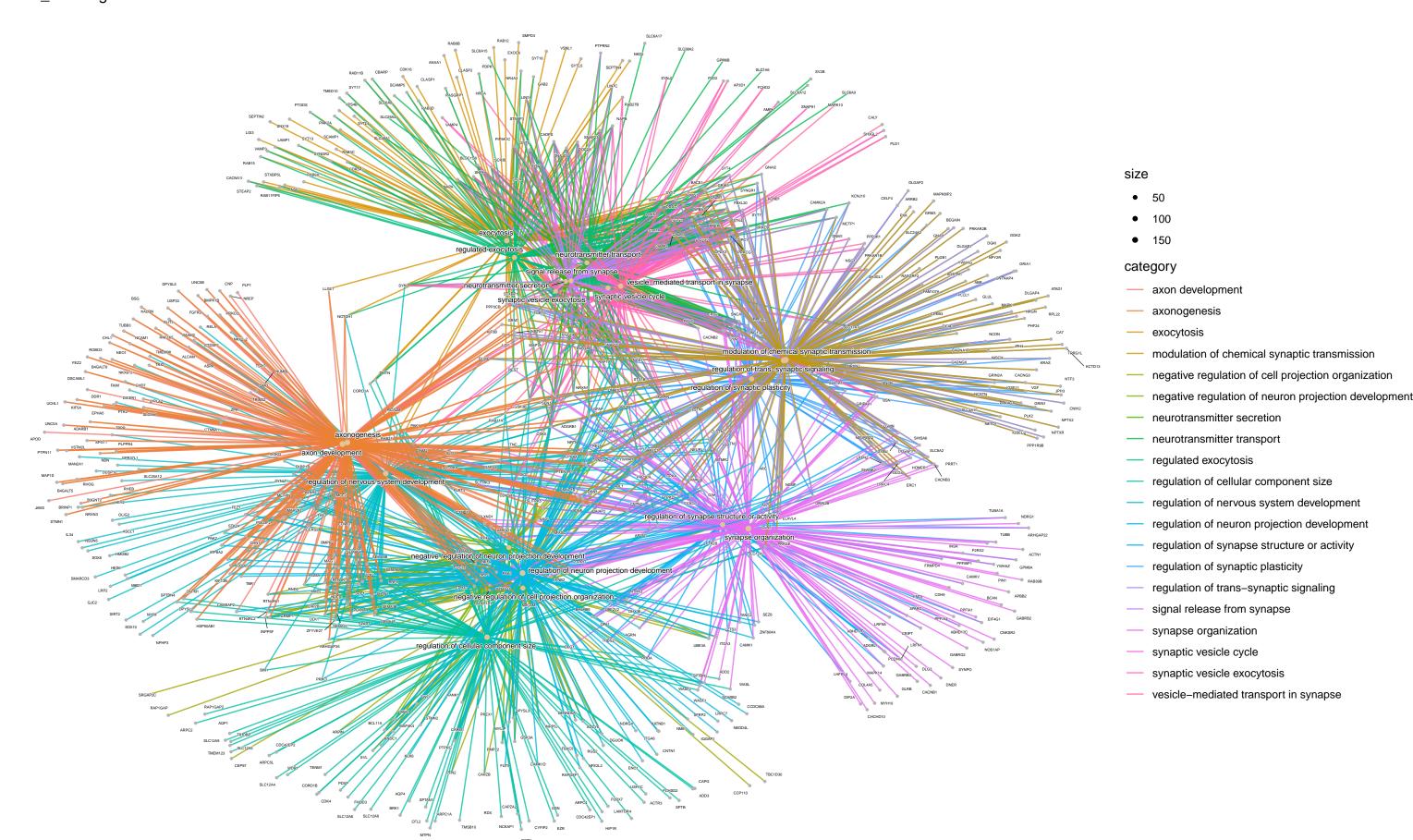
proton-transporting ATPase activity, rotational mechanism

pyrophosphate hydrolysis-driven proton transmembrane transporter activity

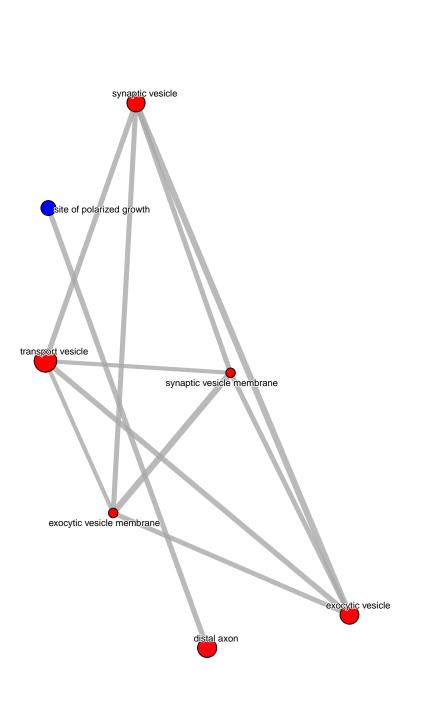
- 50
- 100

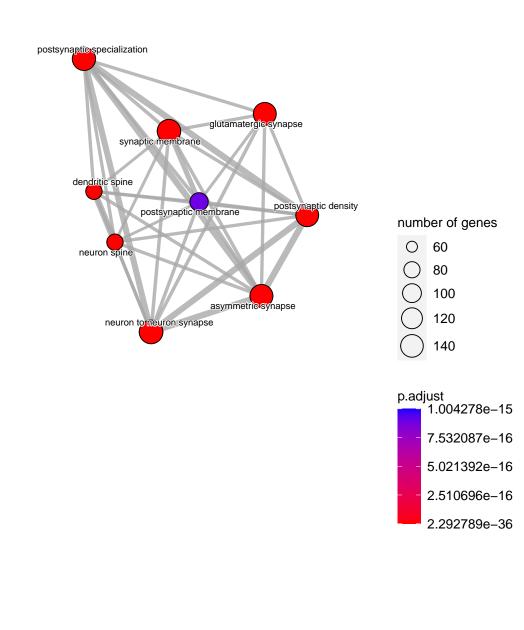
category

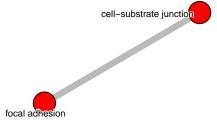
- actin binding
- actin filament binding
- ATPase–coupled ion transmembrane transporter activity
- ATPase activity, coupled to transmembrane movement of ions, rotational mechanism
- cadherin binding
- calmodulin binding
- GTPase activator activity
- GTPase regulator activity
- metal ion transmembrane transporter activity
- microtubule binding
- nucleoside-triphosphatase regulator activity
- phospholipid binding
- phosphoric ester hydrolase activity
- proton-transporting ATPase activity, rotational mechanism
- pyrophosphate hydrolysis-driven proton transmembrane transporter activity
- SNARE binding
- syntaxin-1 binding
- syntaxin binding
- transmembrane transporter binding
- tubulin binding





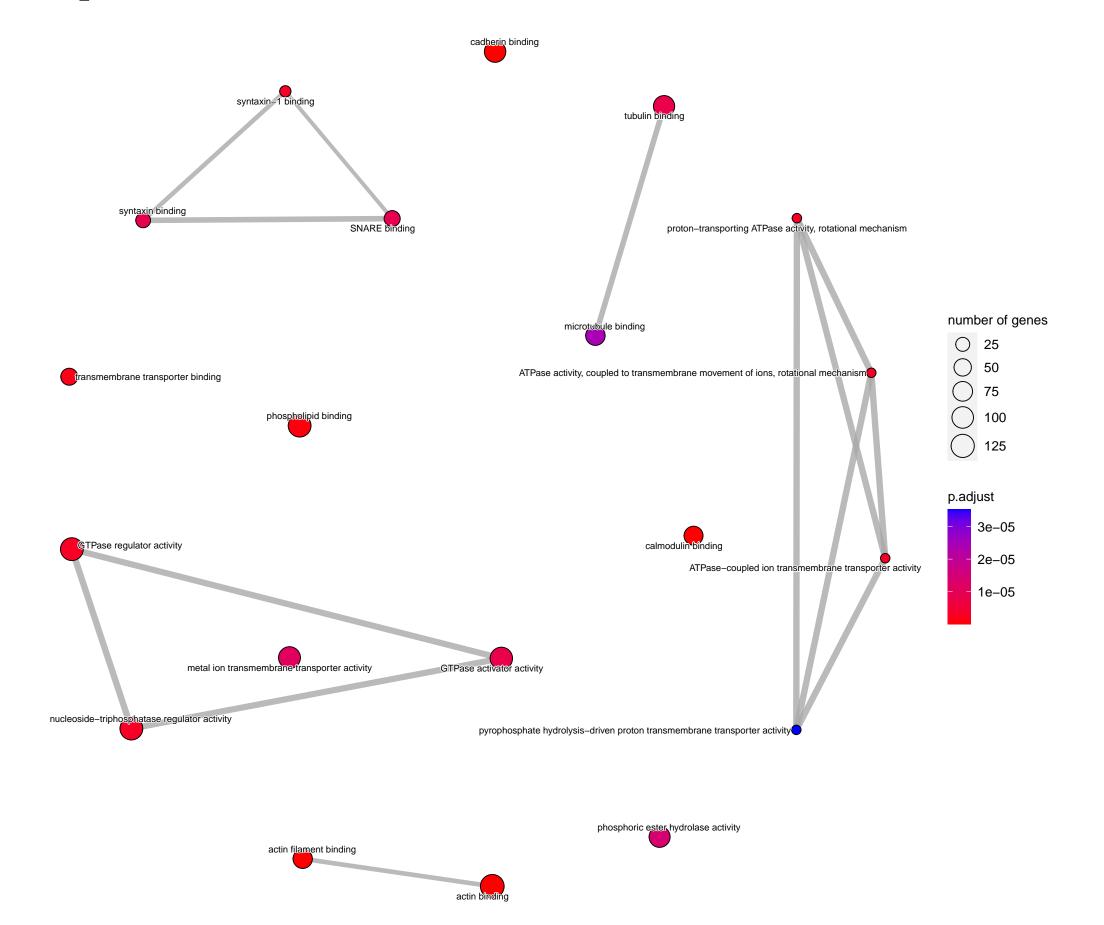








cluster_2 Molecular Function Modules



cluster_2 Biological Process Modules

