

Metagroup 5 Silhouette: 0.37 P-value: 3.2e-08 Num genes: 11

Axon (CC)
Cell projection (CC)

HCN1 SORT1 APOE THY1 DDN CHRM1 FZD3 NLGN4X NRSN2
STRC ROBO2 HHIP SLC3A2

Metagroup 6 Silhouette: 0.36 P-value: 2.8e-07 Num genes: 13

Dendrite (CC)
Neuronal cell body (CC)
Cell surface (CC)

Metagroup 7 Silhouette: 0.28 P-value: 2.1e-06 Num genes: 9

Postsynaptic membrane (CC)
Synapse (CC)
Neuroactive ligand-receptor interaction

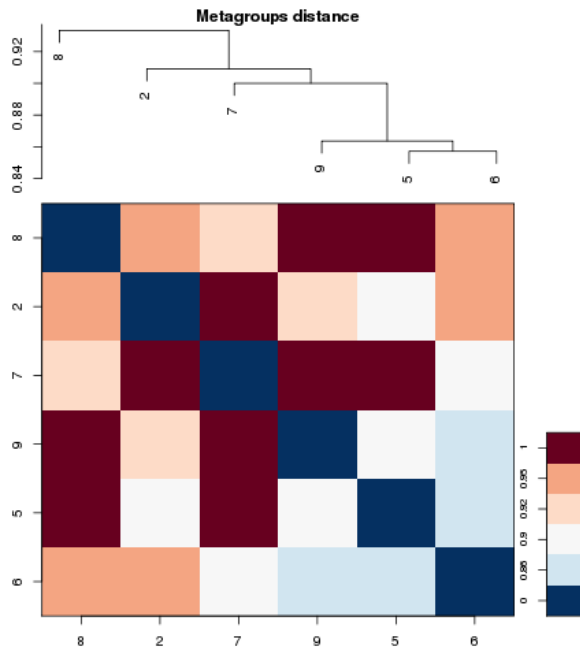
Metagroup 9 Silhouette: 0.27 P-value: 0.00035 Num genes: 12

Enzyme binding (MF)
Microtubule (CC)
Alzheimer's disease
MAPK signaling pathway

[GO terms tree]



Distances between Metagroups:



Distance matrix:

	Mg8	Mg2	Mg7	Mg9	Mg5	Mg6
Mg8	0	0.95	0.93	1	1	0.95
Mg2	0.95	0	1	0.91	0.91	0.96
Mg7	0.93	1	0	1	1	0.9
Mg9	1	0.91	1	0	0.9	0.86
Mg5	1	0.91	1	0.9	0	0.86
Mg6	0.95	0.96	0.9	0.86	0.86	0