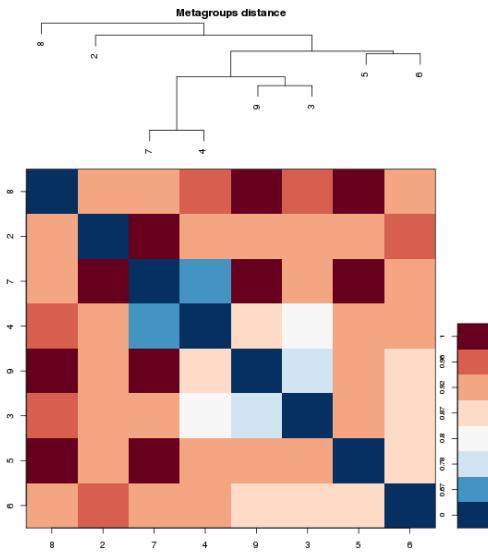


Metagroup 3	Silhouette: 0.2	P-value: 6.4e-10	Genes: 21	
Alzheimer's disease *				Kegg
Calcium ion transport (BP)				GO
Calcium signaling pathway				Kegg
Calmodulin binding (MF)				GO
GnRH signaling pathway				Kegg
Induction of apoptosis by extracellular signals (BP)				GO
Long-term potentiation				Kegg
Melanogenesis				Kegg
Neurotrophin signaling pathway				Kegg
Salivary secretion				Kegg
Tuberculosis				Kegg
Vascular smooth muscle contraction				Kegg
Wnt signaling pathway				Kegg
[GO tree]				
Metagroup 4	Silhouette: 0.06	P-value: 2.1e-09	Genes: 15	
Glutamatergic synapse				Kegg
Postsynaptic density (CC)				GO
Postsynaptic membrane (CC) *				GO
Synapse (CC) *				GO
[GO tree]				

* Terms marked with an asterisk are in several metagroups.

Distances between Metagroups:



Distance matrix:

	Mg8	Mg2	Mg7	Mg4	Mg9	Mg3	Mg5	Mg6
Mg8	0	0.95	0.93	0.95	1	0.96	1	0.95
Mg2	0.95	0	1	0.92	0.91	0.9	0.91	0.96
Mg7	0.93	1	0	0.67	1	0.93	1	0.9
Mg4	0.95	0.92	0.67	0	0.88	0.8	0.92	0.92
Mg9	1	0.91	1	0.88	0	0.78	0.9	0.86
Mg3	0.96	0.9	0.93	0.8	0.78	0	0.93	0.87
Mg5	1	0.91	1	0.92	0.9	0.93	0	0.86
Mg6	0.95	0.96	0.9	0.92	0.86	0.87	0.86	0