

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 asterisc are in several metagroups.

Distances between Metagroups:

