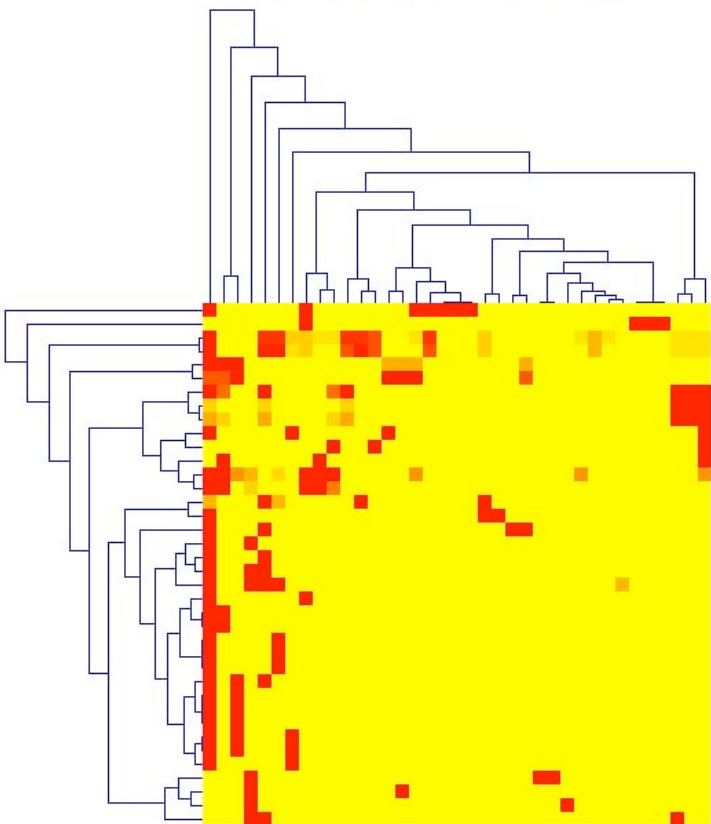




BCL2
 HMCB1
 SLC11A1
 SNCA
 BDNF
 NEFL
 CNAQ
 ONECUT2
 IRS2
 PITX2
 ENH1
 SLC11A1
 CDC25B
 SLC11A1
 DAB1
 NRCAM
 CPNMB
 PKP2
 IGF1
 MYB
 SALL1
 SPRY1
 ALDH18A1
 SCD1
 ACOT10
 NR2F2
 FADS2
 FABP7
 TNFRSF21
 LTBP1
 LTBP3
 TOB1
 NTRK2
 RORB
 VEGFA



GO:0022610, biological_adhesion:3
 GO:0007179, transforming_growth_factor_beta_receptor_signaling_pathway
 GO:0007417, central_nervous_system_development:10
 GO:0030030, cell_projection_organization:8
 GO:0051174, regulation_of_phosphorus_metabolic_process:14
 GO:0051174, regulation_of_phosphorus_metabolic_process:11
 GO:0007423, sensory_organ_development:5
 GO:0048592, eye_morphogenesis:7
 GO:0001654, eye_development:16
 GO:0048469, cell_maturation
 GO:0001569, patterning_of_blood_vessels
 GO:0010463, mesenchymal_cell_proliferation:2
 GO:0040011, locomotion:4
 GO:0051270, regulation_of_cell_motion:12
 GO:0050767, regulation_of_neurogenesis:6
 GO:0000082, G1_S_transition_of_mitotic_cell_cycle
 GO:0001656, metanephros_development:15
 GO:0007006, mitochondrial_membrane_organization
 GO:0046666, retinal_cell_programmed_cell_death
 GO:0042493, response_to_drug
 GO:0051402, neuron_apoptosis:13
 GO:0001952, regulation_of_cell-matrix_adhesion
 GO:0031960, response_to_corticosteroid_stimulus
 GO:0051384, response_to_glucocorticoid_stimulus
 GO:0031103, axon_regeneration
 GO:0031102, neuron_projection_regeneration
 GO:0048678, response_to_axon_injury
 GO:0010623, developmental_programmed_cell_death
 GO:0033032, regulation_of_myeloid_cell_apoptosis
 GO:0033028, myeloid_cell_apoptosis
 GO:0033033, negative_regulation_of_myeloid_cell_apoptosis
 GO:0048066, pigmentation_during_development:1
 GO:0048070, regulation_of_pigmentation_during_development
 GO:0042551, neuron_maturation
 GO:0046394, carboxylic_acid_biosynthetic_process:9
 GO:0042116, macrophage_activation
 GO:0006637, acyl-CoA_metabolic_process
 GO:0014047, glutamate_secretion