



Analysis Name: CPu-EvC no serum-neuro restrict

Analysis Creation Date: 2015-11-19

Build version: 355958M

Content version: 24718999 (Release Date: 2015-09-14)

Analysis Settings

Reference set: Ingenuity Knowledge Base (Genes Only)

Relationship to include: Direct and Indirect

Includes Endogenous Chemicals

Optional Analyses: My Pathways My List

Filter Summary:

Consider only molecules and/or relationships where

(confidence = Experimentally Observed) AND

(tissues = Stem cells not otherwise specified OR Dorsal Root Ganglion OR Gray Matter OR White Matter OR Sciatic Nerve OR Parietal Lobe OR Neurons not otherwise specified OR Cortical neurons OR Cerebral Ventricles OR Thalamus OR Subventricular Zone OR Corpus Callosum OR Nervous System not otherwise specified OR Amygdala OR Trigeminal Ganglion OR Brain OR Hippocampus OR Microvascular endothelial cells OR Granule cells OR Brainstem OR Purkinje cells OR Pituitary Gland OR Other Neurons OR Stromal cells OR Endothelial cells not otherwise specified OR Nucleus Accumbens OR Olfactory Bulb OR Astrocytes OR Other Nervous System OR Choroid Plexus OR Microglia OR Striatum OR Ventricular Zone OR Hypothalamus OR Substantia Nigra OR Spinal Cord OR Cells not otherwise specified OR Caudate Nucleus OR Cerebral Cortex OR Cerebellum OR Adipocytes OR Medulla Oblongata OR Pyramidal neurons OR Granule Cell Layer OR Putamen OR Other Stem cells)

Top Canonical Pathways

Name	p-value	Overlap
Synaptic Long Term Potentiation	2.26E-05	6.9 % 7/101
Protein Kinase A Signaling	3.84E-05	3.7 % 11/295
Melatonin Signaling	1.48E-04	8.3 % 5/60
Dopamine-DARPP32 Feedback in cAMP Signaling	1.65E-04	5.1 % 7/138
GPCR-Mediated Nutrient Sensing in Enteroendocrine Cells	3.07E-04	7.1 % 5/70

Top Upstream Regulators

Upstream Regulator	p-value of overlap	Predicted Activation
RICTOR	1.08E-05	
L-glutamic acid	1.25E-03	
JAK2	1.84E-03	
HTT	2.08E-03	
PPP3CA	3.22E-03	

Top Diseases and Bio Functions

Diseases and Disorders

Name	p-value	#Molecules
Neurological Disease	4.16E-02 - 1.90E-03	18
Psychological Disorders	1.68E-02 - 1.90E-03	14
Hereditary Disorder	1.68E-02 - 4.03E-03	8
Skeletal and Muscular Disorders	8.20E-03 - 4.03E-03	9
Cancer	8.46E-03 - 8.46E-03	1

Molecular and Cellular Functions

Name	p-value	#Molecules
Cellular Assembly and Organization	4.97E-02 - 4.20E-04	14
Carbohydrate Metabolism	5.16E-04 - 5.16E-04	3
Molecular Transport	1.68E-02 - 5.16E-04	5
Small Molecule Biochemistry	3.52E-02 - 5.16E-04	8
Cell Morphology	4.97E-02 - 8.75E-04	22

Physiological System Development and Function

Name	p-value	#Molecules
Nervous System Development and Function	4.97E-02 - 3.54E-04	37
Tissue Morphology	4.97E-02 - 3.54E-04	17
Organ Morphology	4.97E-02 - 1.04E-03	13
Organismal Development	4.97E-02 - 1.04E-03	20
Endocrine System Development and Function	4.97E-02 - 8.46E-03	3

Top Networks

ID	Associated Network Functions	Score
1	Cell Morphology, Cellular Development, Cellular Growth and Proliferation	21
2	Cell-To-Cell Signaling and Interaction, Nervous System Development and Function, Neurological Disease	17
3	Cardiovascular System Development and Function, Organismal Development, Cellular Movement	15
4	Neurological Disease, Skeletal and Muscular Disorders, Psychological Disorders	13
5	Cellular Growth and Proliferation, Gene Expression, Cell Cycle	6

Top Tox Lists

Name	p-value	Overlap
Negative Acute Phase Response Proteins	6.97E-04	40.0 % 2/5
RAR Activation	2.65E-03	3.7 % 6/164

Mechanism of Gene Regulation by Peroxisome Proliferators via PPAR	6.53E-03	4.5 % 4/88
Hepatic Cholestasis	6.85E-03	3.5 % 5/141
PPAR/RXR Activation	9.32E-03	3.3 % 5/152

Top My Lists

Name	p-value	Overlap
66 EvC only (mostly neuronal) molecules	5.74E-109	98.0 % 50/51

Top Analysis-Ready Molecules

Exp Fold Change up-regulated

Molecules	Exp. Value	Exp. Chart
TRIM32	↑ 43443686634.987	
TLE4	↑ 10296483346.115	
LIAS	↑ 6.968	
SERPINC1	↑ 5.965	
GPX3	↑ 4.745	
TTR	↑ 4.582	
MICALL1	↑ 4.460	
TP53I11	↑ 2.898	
SERPINF2	↑ 2.839	
PLEKHB1	↑ 2.653	

Exp Fold Change down-regulated

Molecules	Exp. Value	Exp. Chart
BID	↓ -7.634	
HAP1	↓ -5.385	
HP	↓ -3.997	
PTPDC1	↓ -3.799	

PLXNC1	↓ -3.310
MYO1B	↓ -3.130
AKAP2	↓ -2.939
EML6	↓ -2.678
RGS14	↓ -2.647
WFS1	↓ -2.482



Analysis Name: CPu-WvE no serum-neuro restrict

Analysis Creation Date: 2015-11-19

Build version: 355958M

Content version: 24718999 (Release Date: 2015-09-14)

Analysis Settings

Reference set: Ingenuity Knowledge Base (Genes Only)

Relationship to include: Direct and Indirect

Includes Endogenous Chemicals

Optional Analyses: My Pathways My List

Filter Summary:

Consider only molecules and/or relationships where

(confidence = Experimentally Observed) AND

(tissues = Stem cells not otherwise specified OR Dorsal Root Ganglion OR Gray Matter OR White Matter OR Sciatic Nerve OR Parietal Lobe OR Neurons not otherwise specified OR Cortical neurons OR Cerebral Ventricles OR Thalamus OR Subventricular Zone OR Corpus Callosum OR Nervous System not otherwise specified OR Amygdala OR Trigeminal Ganglion OR Brain OR Hippocampus OR Microvascular endothelial cells OR Granule cells OR Brainstem OR Purkinje cells OR Pituitary Gland OR Other Neurons OR Stromal cells OR Endothelial cells not otherwise specified OR Nucleus Accumbens OR Olfactory Bulb OR Astrocytes OR Other Nervous System OR Choroid Plexus OR Microglia OR Striatum OR Ventricular Zone OR Hypothalamus OR Substantia Nigra OR Spinal Cord OR Cells not otherwise specified OR Caudate Nucleus OR Cerebral Cortex OR Cerebellum OR Adipocytes OR Medulla Oblongata OR Pyramidal neurons OR Granule Cell Layer OR Putamen OR Other Stem cells)

Top Canonical Pathways

Name	p-value	Overlap
CREB Signaling in Neurons	1.35E-05	6.0 % 9/151
Glutamate Receptor Signaling	1.61E-05	10.5 % 6/57
Neuropathic Pain Signaling In Dorsal Horn Neurons	1.86E-05	8.0 % 7/87
G-Protein Coupled Receptor Signaling	8.71E-05	4.2 % 10/237
Synaptic Long Term Potentiation	4.00E-04	5.9 % 6/101

Top Upstream Regulators

Upstream Regulator	p-value of overlap	Predicted Activation
GRM3	3.04E-06	Activated
HTT	5.26E-06	
L-dopa	7.05E-05	
DRD3	2.51E-04	
NR4A2	4.73E-04	

Top Diseases and Bio Functions

Diseases and Disorders

Name	p-value	#Molecules
Neurological Disease	4.68E-02 - 2.90E-05	30
Psychological Disorders	4.68E-02 - 2.90E-05	26
Metabolic Disease	9.54E-03 - 2.77E-03	8
Hereditary Disorder	3.76E-02 - 8.25E-03	9
Skeletal and Muscular Disorders	3.76E-02 - 8.25E-03	9

Molecular and Cellular Functions

Name	p-value	#Molecules
Amino Acid Metabolism	2.83E-02 - 2.15E-05	8
Cell-To-Cell Signaling and Interaction	4.68E-02 - 2.15E-05	20
Molecular Transport	4.68E-02 - 2.15E-05	14
Small Molecule Biochemistry	4.68E-02 - 2.15E-05	15
Lipid Metabolism	2.00E-02 - 2.88E-05	7

Physiological System Development and Function

Name	p-value	#Molecules
Endocrine System Development and Function	4.68E-02 - 2.88E-05	4
Nervous System Development and Function	4.68E-02 - 6.68E-05	21
Embryonic Development	2.83E-02 - 8.85E-04	7
Organ Development	4.68E-02 - 8.85E-04	3
Organismal Development	9.54E-03 - 8.85E-04	7

Top Networks

ID	Associated Network Functions	Score
1	Behavior, Cell-To-Cell Signaling and Interaction, Drug Metabolism	24
2	Cell Death and Survival, Behavior, Lipid Metabolism	24
3	Neurological Disease, Psychological Disorders, Skeletal and Muscular Disorders	18
4	Cell Death and Survival, Cellular Assembly and Organization, Cellular Development	18
5	Behavior, Neurological Disease, Psychological Disorders	15

Top Tox Lists

Name	p-value	Overlap
FXR/RXR Activation	8.78E-03	4.7 % 4/85
Mitochondrial Dysfunction	1.03E-02	3.6 % 5/138

Positive Acute Phase Response Proteins	1.39E-02	10.5 % 2/19
PXR/RXR Activation	1.40E-02	5.7 % 3/53
Hypoxia-Inducible Factor Signaling	2.04E-02	4.9 % 3/61

Top My Lists

Name	p-value	Overlap
90 WvE only (mostly neuronal) molecules	1.53E-132	92.8 % 64/69

Top Analysis-Ready Molecules

Exp Fold Change up-regulated

Molecules	Exp. Value	Exp. Chart
PDYN	↑ 8.682	
BID	↑ 8.642	
TMSB10/TMSB4X	↑ 4.609	
CNR1	↑ 2.783	
COQ7	↑ 2.745	
MYO1B	↑ 2.635	
SLC6A3	↑ 2.497	
SLC1A3	↑ 2.370	
STK26	↑ 2.325	
WFS1	↑ 2.252	

Exp Fold Change down-regulated

Molecules	Exp. Value	Exp. Chart
LIAS	↓ -7.721	
RPL28	↓ -6.233	
CHERP	↓ -6.140	
SERPINC1	↓ -6.102	

SERPINF2	↓ -4.441
KIAA1217	↓ -4.407
ILF3	↓ -4.204
TTR	↓ -4.120
PJA2	↓ -3.574
GPX3	↓ -3.330



Analysis Name: CPu-WvC (NS,NR)

Analysis Creation Date: 2015-11-27

Build version: 355958M

Content version: 24718999 (Release Date: 2015-09-14)

Analysis Settings

Reference set: Ingenuity Knowledge Base (Genes Only)

Relationship to include: Direct and Indirect

Includes Endogenous Chemicals

Optional Analyses: My Pathways My List

Filter Summary:

Consider only molecules and/or relationships where

(confidence = Experimentally Observed) AND

(tissues = Amygdala OR Cells not otherwise specified OR Cerebellum OR Brainstem OR Hippocampus OR Medulla Oblongata OR Granule Cell Layer OR Granule cells OR Pituitary Gland OR Microvascular endothelial cells OR Stromal cells OR Other Neurons OR Gray Matter OR Cortical neurons OR Other Nervous System OR Astrocytes OR Parietal Lobe OR Neurons not otherwise specified OR Olfactory Bulb OR Striatum OR Cerebral Ventricles OR Corpus Callosum OR Substantia Nigra OR Thalamus OR Caudate Nucleus OR Trigeminal Ganglion OR Brain OR Nervous System not otherwise specified OR Cerebral Cortex OR Putamen OR Sciatic Nerve OR Purkinje cells OR Adipocytes OR Pyramidal neurons OR Other Stem cells OR White Matter OR Dorsal Root Ganglion OR Endothelial cells not otherwise specified OR Nucleus Accumbens OR Choroid Plexus OR Microglia OR Subventricular Zone OR Ventricular Zone OR Hypothalamus OR Stem cells not otherwise specified OR Spinal Cord)

Top Canonical Pathways

Name	p-value	Overlap
Mitochondrial Dysfunction	5.42E-08	10.1 % 14/138
Oxidative Phosphorylation	1.10E-07	13.1 % 11/84
VEGF Signaling	2.35E-04	9.2 % 7/76
Cholecystokinin/Gastrin-mediated Signaling	7.09E-04	7.7 % 7/91
Glycogen Biosynthesis II (from UDP-D-Glucose)	7.88E-04	66.7 % 2/3

Top Upstream Regulators

Upstream Regulator	p-value of overlap	Predicted Activation
RICTOR	7.67E-05	
BDNF	1.85E-04	
ROR1	2.52E-04	
ROR2	2.52E-04	
HTT	3.00E-04	

Top Diseases and Bio Functions

Diseases and Disorders

Name	p-value	#Molecules
Neurological Disease	4.95E-02 - 1.82E-05	39
Organismal Injury and Abnormalities	4.95E-02 - 5.80E-04	21
Psychological Disorders	4.82E-02 - 1.05E-03	27
Skeletal and Muscular Disorders	4.95E-02 - 1.22E-03	20
Inflammatory Response	4.82E-02 - 2.57E-03	5

Molecular and Cellular Functions

Name	p-value	#Molecules
Cell Morphology	4.82E-02 - 2.13E-06	49
Cellular Assembly and Organization	4.82E-02 - 3.98E-06	43
Cellular Function and Maintenance	4.82E-02 - 3.98E-06	40
Cellular Development	4.82E-02 - 7.26E-06	46
Cell-To-Cell Signaling and Interaction	4.95E-02 - 1.14E-04	31

Physiological System Development and Function

Name	p-value	#Molecules
Nervous System Development and Function	4.95E-02 - 5.21E-06	66
Tissue Morphology	4.12E-02 - 5.21E-06	41
Tissue Development	4.82E-02 - 7.26E-06	42
Organ Morphology	4.82E-02 - 1.09E-05	27
Organismal Development	4.82E-02 - 1.09E-05	33

Top Tox Functions**Assays: Clinical Chemistry and Hematology**

Name	p-value	#Molecules
Decreased Levels of Albumin	1.63E-02 - 1.63E-02	1

Cardiotoxicity

Name	p-value	#Molecules
Pulmonary Hypertension	1.63E-02 - 1.63E-02	1

Top Networks

ID	Associated Network Functions	Score
1	Behavior, Cellular Development, Cellular Growth and Proliferation	16
2	Cell Death and Survival, Cellular Development, Nervous System Development and Function	15
3	Neurological Disease, Cell Death and Survival, Psychological Disorders	13
4	Cell Death and Survival, Cancer, Organismal Injury and Abnormalities	13
5	Behavior, Cell Death and Survival, Cellular Movement	13

Top Tox Lists

Name	p-value	Overlap
Mitochondrial Dysfunction	5.42E-08	10.1 % 14/138
Hypoxia-Inducible Factor Signaling	3.14E-03	8.2 % 5/61
RAR Activation	5.49E-03	4.9 % 8/164
Cardiac Fibrosis	2.16E-02	4.1 % 7/170
VDR/RXR Activation	3.28E-02	5.4 % 4/74

Top My Lists

Name	p-value	Overlap
175 WvC only (mostly neuronal) molecules	2.28E-299	99.3 % 149/150

Top Analysis-Ready Molecules

Exp Fold Change up-regulated

Molecules	Exp. Value	Exp. Chart
MICALL1	↑ 4.020	
TRPV2	↑ 3.506	
ENDOG	↑ 2.890	
NDUFS6	↑ 2.726	
MZT2A	↑ 2.580	

COX6A1	↑ 2.387
CAV1	↑ 2.282
TIMM22	↑ 2.269
CLNS1A	↑ 2.259
MOBP	↑ 2.173

Exp Fold Change down-regulated

Molecules	Exp. Value	Exp. Chart
HOMER2	↓ -35054557201.723	
FHL2	↓ -4588652106.352	
BAIAP3	↓ -4.829	
NGB	↓ -4.607	
GYG1	↓ -3.816	
PJA2	↓ -3.573	
DHPS	↓ -3.042	
RPL18	↓ -2.856	
ATF2	↓ -2.807	
AHI1	↓ -2.652	