Simvastatin Type 2 Diabetes Mellitus

q-value <= 0.05

Up in disease, down with

treatment

AMIT SERUM RESPONSE 40

MCF10A'

'Genes whose expression peaked at 40 min after stimulation of MCF10A cells with serum.'

Top 100 genes negatively associated with T-cell acute lymphoblastic leukemia MLL T-ALL)

FERRANDO T ALL WITH MLL

ENL FUSION DN'

expressing MLL-ENL fusion

[GeneID=4297;4298].'

'Genes down-regulated in T98G cells

RUIZ TNC TARGETS DN' (glioblastoma) by TNC [GeneID=3371].'

'Genes down-regulated in the ANBL-6 cell line

CROONQUIST IL6 (multiple myeloma, MM) after withdrawal of

DEPRIVATION DN' IL6 [GeneID=3569].'

Down in disease, up with

treatment

'Adipocyte genes induced in 3T3-L1 cells

(adipocyte) by constitutively active PPARG

LI ADIPOGENESIS BY [GeneID=5468] or its agonist, TZD

ACTIVATED PPARG' [PubChem=5437].'

Sirolimus AML

q-value <= 0.01

Up in disease, down with

treatment

KEGG SPLICEOSOME' 'Spliceosome'
MIPS SPLICEOSOME' 'Spliceosome'

MIPS C COMPLEX

SPLICEOSOME' 'C complex spliceosome'

REACTOME PROCESSING OF

CAPPED INTRON CONTAINING 'Genes involved in Processing of Capped Intron-

PRE MRNA' Containing Pre-mRNA'

REACTOME TRANSPORT OF

MATURE TRANSCRIPT TO Genes involved in Transport of Mature

CYTOPLASM' Transcript to Cytoplasm'

REACTOME MRNA

PROCESSING' 'Genes involved in mRNA Processing'
REACTOME MRNA SPLICING' 'Genes involved in mRNA Splicing'

REACTOME MRNA 3 END

PROCESSING' 'Genes involved in mRNA 3"-end processing'

REACTOME CLEAVAGE OF

GROWING TRANSCRIPT IN THE 'Genes involved in Cleavage of Growing TERMINATION REGION ' Transcript in the Termination Region '

PICCALUGA 'Down-regulated genes in angioimmunoblastic **ANGIOIMMUNOBLASTIC** lymphoma (AILT) compared to normal T LYMPHOMA DN' lymphocytes.' 'Genes down-regulated in Daudi cells (B lymphocytes) stably expressing CD5 **GARY CD5 TARGETS DN'** [GeneID=921] off a plasmid vector.' 'Genes down-regulated in metastatic breast cancer tumors having type 2 amplification in **GINESTIER BREAST CANCER** the 20q13 region; involves MYBL2, STK6 and ZNF217 [GeneID=4605;6790;7764]' 20013 AMPLIFICATION DN' **RODRIGUES THYROID** 'Genes up-regulated in poorly differentiated thyroid carcinoma (PDTC) compared to normal CARCINOMA POORLY DIFFERENTIATED UP' thyroid tissue.' 'Genes up-regulated in anaplastic thyroid **RODRIGUES THYROID** carcinoma (ATC) compared to normal thyroid CARCINOMA ANAPLASTIC UP' tissue.' 'Genes down-regulated in T helper cells (defines as CD4+) isolated from patients with HAHTOLA MYCOSIS mucosis fungoides compared to those from **FUNGOIDES CD4 DN'** normal control donors.' **ENK UV RESPONSE** 'Genes down-regulated in NHEK cells (normal KERATINOCYTE DN' epidermal keratinocytes) after UVB irradiation.' 'Genes down-regulated in ME-A cells (breast cancer, sensitive to apoptotic stimuli) exposed to doxorubicin [PubChem=31703] in the presence of medium concentrate (MC) from GRAESSMANN RESPONSE TO ME-C cells (breast cancer, resistant to apoptotic MC AND DOXORUBICIN DN' stimuli).' 'Genes up-regulated in T1 cells (primary melanoma, sensitive to TRAIL [GeneID=8743]) compared to G1 cells (metastatic melanoma, HAMAI APOPTOSIS VIA TRAIL UP' resistant to TRAIL).' 'Genes changed in xenograft tumors formed by SEIDEN ONCOGENESIS BY DLD-1 or DKO-4 cells (colon cancer) MET' overexpressing MET [GeneID=4233].' 'Cluster 7: genes up-regulated in B493-6 cells (B lymphocytes) by MYC [GeneID=4609] and down-SCHLOSSER MYC TARGETS regulated by the combination of MYC and REPRESSED BY SERUM' serum.' 'Common down-regulated transcripts in fibroblasts expressing either XP/CS or TDD

mutant forms of ERCC3 [GeneID=2071], after

UVC irradiation.'

DACOSTA UV RESPONSE VIA

ERCC3 COMMON DN'

	regulated at 6 hr dihydrotestosterone
SCHAEFFER PROSTATE	[PubChem=10635]) which are also down-
DEVELOPMENT AND CANCER	regulated in localized vs metastatic prostate
BOX4 DN'	cancers.'
567.151.	'Genes whose expression positively correlated
	with that of SMARCA2 [GeneID=6595] in
SHEN SMARCA2 TARGETS UP'	
SHEN SWARCAZ TARGETS UP	prostate cancer samples.'
	'Genes up-regulated in P493-6 cells (Burkitt''s
	lymphoma) induced to express MYC
UP'	[GeneID=4609].'
	'Genes down-regulated in BJAB cells (B-
PENG GLUTAMINE	lymphoma) after glutamine [PubChem=738]
DEPRIVATION DN'	deprivation.'
ZHAN MULTIPLE MYELOMA	'Top genes up-regulated in MM4 vs MM1
SUBGROUPS'	subgroup of multiple myeloma samples.'
	'Down-regulated genes distinguishing in
MOREAUX MULTIPLE	multiple myeloma (MM) samples with lower
MYELOMA BY TACI DN'	expression of TACI [GeneID=23495].'
	'Genes whose CpG islands showed greatly
	increased histone H4 acetylation in NB4 cells
NOUZOVA TRETINOIN AND H4	·
ACETYLATION'	treatment with tretinoin [PubChem=5538].'
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	'Genes up-regulated in HEK293 cells
	(embryonic kidney) at 6 h, 12 h or 24 h after
DEBIASI APOPTOSIS BY	infection with reovirus strain T3A (known as a
REOVIRUS INFECTION UP'	strong inducer of apoptosis).'
REGVINGS IN ECTION OF	'Genes enriched in embryonic, neural and
RAMALHO STEMNESS UP'	hematopoietic stem cells.'
DAZARD RESPONSE TO UV	•
	'Genes down-regulated in NHEK cells (normal
NHEK DN'	keratinocytes) by UV-B irradiation.'
	'Cluster G6: genes increasingly down-regulated
DAZARD UV RESPONSE	in NHEK cells (normal keratinocyte) after UV-B
CLUSTER G6'	irradiation.'
	'Genes down-regulated in SaOS-2 cells
	(osteosarcoma) upon knockdown of YY1
DE YY1 TARGETS DN'	[GeneID=7528] by RNAi.'
	'Genes up-regulated 9 h after induction of
CHEN HOXA5 TARGETS 9HR	HoxA5 [GeneID=3205] expression in a breast
UP'	cancer cell line.'
	'Genes down-regulated in neural stem cells
ZHANG TLX TARGETS 36HR	(NSC) at 36 h after cre-lox knockout of TLX
DN'	(NR2E1) [GeneID=7101].'

'Early prostate development genes (down-

'Cohesin targets identified by ChIP-chip which WENDT COHESIN TARGETS were up-regulated after knockdown of CTCF UP' and RAD21 [GeneID=10664;5885] by RNAi.' 'Transcripts enriched in pseudopodia of NIH/3T3 cells (fibroblast) in response to MILI PSEUDOPODIA haptotactic migratory stimulus by fibronectin, HAPTOTAXIS UP' FN1 [GeneID=2335].' THILLAINADESAN ZNF217 'Genes bound and activated by ZNF217 TARGETS UP' [GeneID=7764] in MCF7 cells (breast cancer).' 'Genes down-regulated upon overexpression of PARVB [GeneID=29780] in MDA-MB-231 cells JOHNSTONE PARVB TARGETS (breast cancer) cultured in 3D collagen I and 3D 2 DN' Matrigel only.' Down in disease, up with treatment **KEGG GRAFT VERSUS HOST** DISFASE' 'Graft-versus-host disease' REACTOME METABOLISM OF STEROID HORMONES AND 'Genes involved in Metabolism of steroid VITAMINS A AND D' hormones and vitamins A and D' REACTOME EXTRACELLULAR 'Genes involved in Extracellular matrix MATRIX ORGANIZATION' organization' REACTOME PEPTIDE LIGAND 'Genes involved in Peptide ligand-binding **BINDING RECEPTORS'** receptors' **PICCALUGA** 'Up-regulated genes in angioimmunoblastic **ANGIOIMMUNOBLASTIC** lymphoma (AILT) compared to normal T LYMPHOMA UP' lymphocytes.' 'Genes down-regulated in nasopharyngeal **SENGUPTA** carcinoma (NPC) positive for LMP1 NASOPHARYNGEAL [GeneID=9260], a latent gene of Epstein-Barr CARCINOMA WITH LMP1 DN' virus (EBV).' 'Genes down-regulated in metastases from malignant melanoma compared to the primary JAEGER METASTASIS DN' tumors.' 'Genes down-regulated in CD34+ [GeneID=947] hematopoetic cells by expression of NUP98-**TAKEDA TARGETS OF NUP98** HOXA9 fusion [GeneID=4928;3205] off a HOXA9 FUSION 8D DN' retroviral vector at 8 days after transduction.'

'Genes down-regulated in CD34+ [GeneID=947]
hematopoetic cells by expression of NUP98TAKEDA TARGETS OF NUP98 HOXA9 fusion [GeneID=4928;3205] off a
HOXA9 FUSION 10D DN' retroviral vector at 10 days after transduction.'

	'Genes down-regulated in CD34+ [GeneID=947]
	hematopoetic cells by expression of NUP98-
TAKEDA TARGETS OF NUP98	HOXA9 fusion [GeneID=4928;3205] off a
HOXA9 FUSION 16D DN'	retroviral vector at 16 days after transduction.'
SABATES COLORECTAL	'Genes up-regulated in colorectal adenoma
ADENOMA UP'	compared to normal mucosa samples.'
	'Genes down-regulated in CD133+
JAATINEN HEMATOPOIETIC	[GeneID=8842] cells (hematopoietic stem cells,
STEM CELL DN'	HSC) compared to the CD133- cells.'
	'Genes identified by subtractive hybridization
	comparing malignant and benign components
	of a hepatocellular carcinoma (HCC) in a pre-
CAVARD LIVER CANCER	existing liver adenoma in a morphologically
MALIGNANT VS BENIGN'	normal liver.'
TARTE PLASMA CELL VS	'Genes up-regulated in mature plasma cells
PLASMABLAST UP'	compared with plasmablastic B lymphocytes.'
	'Genes down-regulated in hematopoietic
HESS TARGETS OF HOXA9 AND	·
MEIS1 DN'	and MEIS1 [GeneID=3205;4211].'
	'Genes down-regulated during in vitro
	maturation of CD14+ [GeneID=929] monocytes
LENAOUR DENDRITIC CELL	(day 0) into immature (day 7) and mature
MATURATION DN'	dendritic cells (day 14).'
	'Genes up-regulated at 6 months of age in lungs
	from LIPA [GeneID=3988] knockout mice, which
LIAN LIPA TARGETS 6M'	display pulmonary pathology.'
	'Genes up-regulated at 3 months of age in lungs
	from LIPA [GeneID=3988] knockout mice, which
LIAN LIPA TARGETS 3M'	display pulmonary pathology.'
MCLACHLAN DENTAL CARIES	'Genes down-regulated in pulpal tissue
DN'	extracted from carious teeth.'
MCLACHLAN DENTAL CARIES	'Genes up-regulated in pulpal tissue extracted
UP'	from carious teeth.'
SMID BREAST CANCER	'Genes down-regulated in the luminal B
LUMINAL B DN'	subtype of breast cancer.'
	'Genes up-regulated in ex-vivo colonic tissue
ZHENG IL22 SIGNALING UP'	after treatment with IL22 [GeneID=50616].'
	'Tumorigenesis markers of head and neck
	squamous cell carcinoma (HNSCC): up-
	regulated in the "early" tumors vs normal
CDOMED THMODICENECIC HD	annual a l

CROMER TUMORIGENESIS UP' samples.'

'Top 100 probe sets contrubuting to the positive side of the 1st principal component; predominantly associated with spindle cell and pleomorphic sarcoma samples.'

NAKAYAMA SOFT TISSUE TUMORS PCA1 UP'