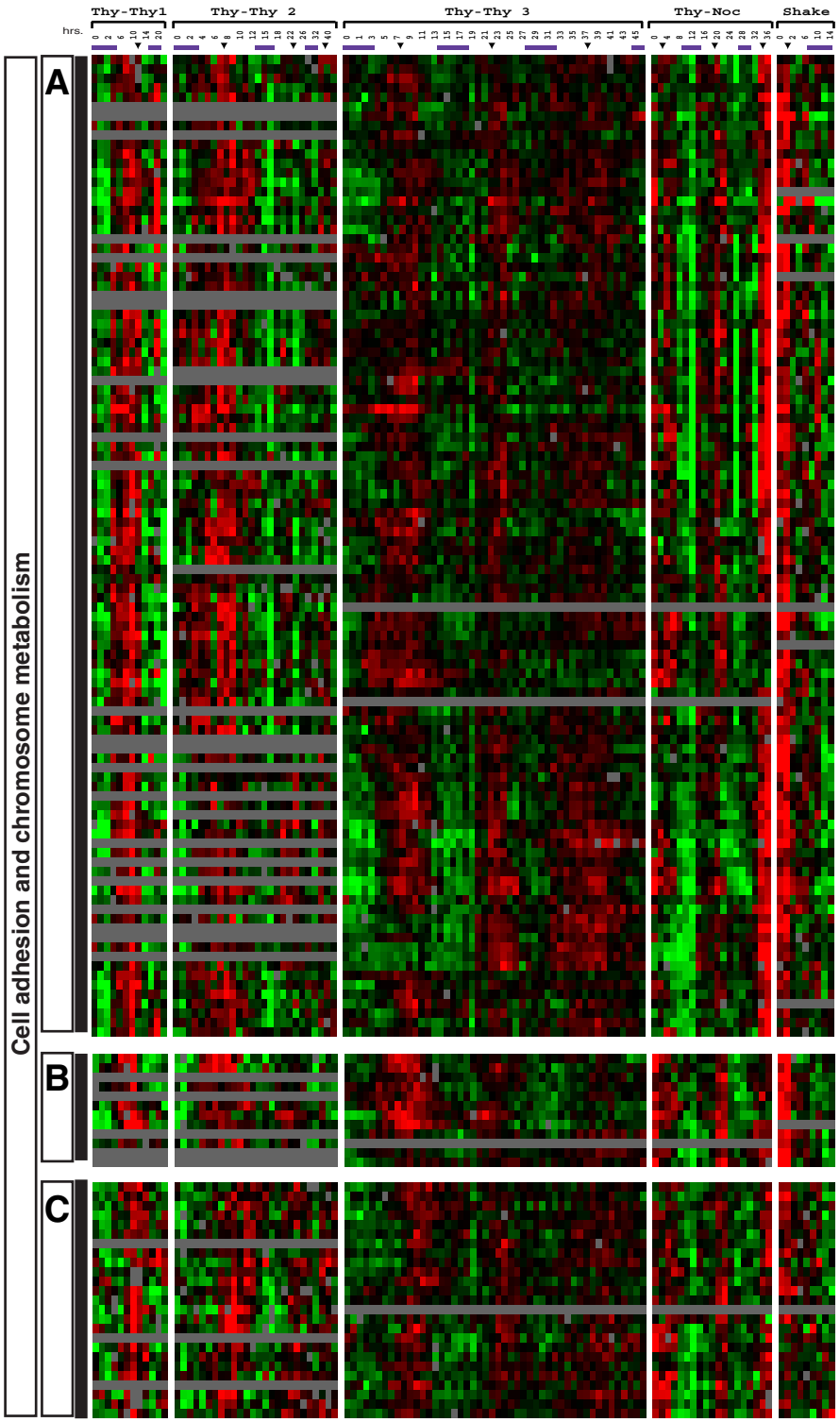

PHASE	NAME	GO BIOLOGICAL PROCESS
M/G1	KRAS2 v-Ki-ras2 Kirsten rat sarcoma 2, N95249	cell cycle control
G2/M	PRPSAP1 phosphoribosyl pyrophosphate synthetase-assoc. 1, R20005	nucleotide metabolism
M/G1	YWHAH (14-3-3 eta), N69107	signal transduction
M/G1	PSEN1 presenilin 1, AA403083	*cell adhesion | chromosome segregation
M/G1	FL89213, AA486770	
G2/M	ESTs, N59330	
M/G1	TERA protein, AA906997	
M/G1	FLJ23260, AA463961	
G2/M	CIA coactivator independent of AF-2, AA521358	
G2/M	FLJ20259 hypothetical protein, T55949	
M/G1	UBE2D3 **ubiquitin-conjugating enzyme E2D3 (UBC4/5 family), AA017199	*ubiquitin-dependent protein degradation
M/G1	HMGE Grp5-like protein cochaperone (mitochondrial), H61884	
M/G1	DKC1 dyskeratosis congenita 1, AA052960	RNA metabolism | Telomere maintenace
M/G1	C1orf2 chromosome 1 open reading frame 2, H11572	
M/G1	PTMS parathymosin, R11526	DNA replication| cell cycle control
M/G1	CDKN3, CDK2-associated dual specificity phosphatase (CDI1/KAP), AA284072	cell cycle control
M/G1	G3BP Ras-GTPase-activating protein, AA449834	signal transduction
M/G1	PPP3CA protein phosphatase 3 catalytic subunit, alpha, W60310	
M/G1	ESTs, R93772	
M/G1	MDM2 p53-binding protein, AI188173	cell proliferation
M/G1	PRKAR1A protein kinase, N36599	signal transduction
G2/M	cDNA DKFZp586I1518, AA287958	
M/G1	TOB2 transducer of ERBB2, 2, AA486088	cell cycle control
M/G1	cDNA FLJ21691, R01288	
M/G1	CRK v-crk avian sarcoma virus CT10 oncogene, H75530	cell motility
M/G1	ESTs, AA862450	
M/G1	clone TA40 untranslated mRNA, AA903913	
M/G1	MRPS2 mitochondrial ribosomal protein S2, AA459419	
G2/M	RNPS1 RNA-binding protein S1, AA496879	RNA metabolism
G2/M	SWI/SNF related, matrix associated, actin dependent regulator of chromatin B1	DNA integration| transcription
G2/M	KIAA0176, R44371	
G2/M	CTNND1 catenin cadherin-associated protein, delta 1 (P120), AA024656	cell adhesion
G2/M	PTPN9 protein tyrosine phosphatase, non-receptor type 9, AA434420	protein modification
G2/M	VCAM1 vascular cell adhesion molecule 1, H16591	
M/G1	MGC5338 hypothetical protein, R83260	
G2/M	SS18 synovial sarcoma translocation, chromosome 18, N59206	
G2/M	SWI/SNF related, matrix associated, actin dependent regulator of chromatin D1	DNA packaging| transcription
G2/M	BIRC2 baculoviral IAP repeat-containing 2, AA702174	signal transduction
G2/M	SET07 PR/SET domain containing protein 07, AA421470	
M/G1	W85890	
M/G1	TERA protein, AA465096	
M/G1	cDNA FLJ14028, AA454682	
M/G1	FLJ20094 hypothetical protein, W24963	
M/G1	FLJ13154 hypothetical protein, AA923560	
M/G1	PRC1 protein regulator of cytokinesis 1, AA449593	cytokinesis
M/G1	NSAP1 NS1-associated protein 1, AA186327	RNA metabolism|Transport
M/G1	FLJ20624 hypothetical protein, AA443698	
M/G1	WS-3 novel RGD-containing protein, AA449975	
G2/M	AA490946	
G2/M	DR1 down-regulator of transcription 1,TBP-binding negative cofactor 2, AA044623	transcription
G2/M	KIF5B kinesin family member 5B, AA608707	cell growth and/or maintenance
M/G1	FBI1 HIV-1 inducer of short transcripts binding protein, R06252	
G2/M	KIAA0978, AA485878	
M/G1	MGC5338 hypothetical protein, AA463627	
M/G1	ESTs, R31512	
M/G1	PPP2CA protein phosphatase 2 alpha, AA599092	cell cycle control
G2/M	IMAGE:3834331, N53024	
G2/M	SFRS3 splicing factor, arginine/serine-rich 3, AA598400	
G2/M	ESTs, N62567	
M/G1	TSC22 TGF beta-stimulated protein TSC-22, AA664389	transcription
M/G1	C4B complement component 4B, AA664406	cell communication
G2/M	INSM1 insulinoma-associated 1, R38640	
G2/M	KIAA0618, H82035	
G2/M	IDI2 diphosphate dimethylallyl diphosphate isomerase 2, R42713	
M/G1	ZNF281 zinc finger protein 281, N47469	transcription
M/G1	VCL vinculin, AA486728	cell shape and cell size control
G2/M	CNN2 calponin 2, AA284856	cytoskeletal organization
G2/M	SPS selenophosphate synthetase, AA488081	protein modification
G2/M	ITPR1 inositol 1,4,5-triphosphate receptor, type 1, AA035450	signal transduction
G2/M	HP1-BP74, N28983	
G2/M	TGIF TGFB-induced factor TALE family homeobox, R83270	transcription
G2/M	ESTs, AA429809	
M/G1	CBX3 chromobox 3, AI139106	
M/G1	ESTs, AI278336	
M/G1	RAN, RAS oncogene family, AA456636	transport
M/G1	G3BP Ras-GTPase-activating protein, AA598628	signal transduction
M/G1	MGC19556 hypothetical protein, AA609463	
M/G1	MRPL19 mitochondrial ribosomal protein L19, AA521243	
G2/M	SFPQ splicing factor, AA426575	RNA metabolism
G2/M	SFPQ splicing factor, AA425853	RNA metabolism
G2/M	RAD51C, R37145	DNA repair| DNA recombination
M/G1	KIAA1265, AA479402	
M/G1	KIAA0952, AA454989	
M/G1	KIAA0952, AA679150	
M/G1	RAD21, AA663102	*sister chromatid cohesion | DNA recombination| DNA repair
M/G1	ZPBP zona pellucida binding protein, AA400474	
M/G1	**EST, AI061169	
M/G1	PTTG1 pituitary tumor-transforming 1, AI362866	*sister chromatid cohesion
M/G1	PTTG1 pituitary tumor-transforming 1, AA430241	*sister chromatid cohesion
M/G1	HRB HIV-1 Rev binding protein, AA485958	RNA localization
M/G1	FREQ frequenin, H16821	
G2/M	CTL2 gene, AA454710	
M/G1	KIAA0306 protein, AA485621	
M/G1	MGC5350 hypothetical protein, AI002036	
M/G1	NUCKS similar to rat nuclear ubiquitous casein kinase 2, AA158346	
G2	NUCKS similar to rat nuclear ubiquitous casein kinase 2, AA927182	
M/G1	NUCKS similar to rat nuclear ubiquitous casein kinase 2, AI053436	
G2/M	E2IG4 estradiol-induced, R13844	
M/G1	DKFZp434D1428 hypothetical protein, AA432275	
M/G1	MGC4604 hypothetical protein, AA211524	
M/G1	HIF1A hypoxia-inducible factor 1 alpha, AA598526	transcription
M/G1	FXR1 fragile X mental retardation 1, N62761	cell death| transport
G2/M	MLLT4 mixed-lineage leukemia trithorax 4, AA010818	cell adhesion
M/G1	TRIP8 thyroid hormone receptor interactor 8, AA425650	transcription
G2/M	NUDE1 LIS1-interacting protein N79612	
M/G1	FLJ11269 hypothetical protein R37888	
M/G1	PSCD3 pleckstrin homology, Sec7 and coiled/coil domains 3 AA629264	*vescile transport
M/G1	CDK7 R22624	cell cycle control
M/G1	CDK7 AI311067	cell cycle control
M/G1	SNK serum-inducible kinase AA460152	
M/G1	TROAP trophinin associated protein (tastin) H94949	cell adhesion
M/G1	TEM8 tumor endothelial marker 8 H58644	
M/G1	FABP1 fatty acid binding protein 1 AA682392	lipid metabolism
M/G1	RHEB2 Ras homolog enriched in brain 2 AA482117	signal transduction
M/G1	cDNA FLJ22844 AA975103	
M/G1	mRNA DKFZp762B195 AA625574	
M/G1	GNB1 guanine nucleotide binding protein G protein, beta1 AA488027	*RAS protein signal transduction
M/G1	KIAA1404, W72798	
M/G1	TIP47 cargo selection protein, AA416787	*vesical transport
M/G1	ESTs, R63922	
M/G1	ILF2 interleukin enhancer binding factor 2, H95712	*transcription
M/G1	CDC42 GTP-binding protein, AA668681	*cytoskeletal regulation
M/G1	TOMM70A translocase of outer mitochondrial membrane 70A, AA702684	
M/G1	**adrenal gland protein AD-004, AA150301	
M/G1	KPNB1 karyopherin (importin) beta 1, AA121732	transport
M/G1	TOP1 topoisomerase I, AA232856	
M/G1	DNAJB6 DnaJ Hsp40 homolog B6, AA431203	
M/G1	RAB3A, member RAS oncogene family, H14230	cell-cell signaling
M/G1	AOC2 amine oxidase, copper containing 2, N50959	amino acid and derivative metabolism
M/G1	MGC10850 hypothetical protein, H46748	
M/G1	ESTs, H50656	
M/G1	ESTs, R27432	
M/G1	KMO kynurenine 3-monooxygenase, AA044326	metabolism
M/G1	KMO kynurenine 3-monooxygenase, AA152183	metabolism
M/G1	Homo sapiens cDNA FLJ11904, W00890	
M/G1	FLJ20425 hypothetical protein, AA424663	
M/G1	KIAA1547, AA136692	
M/G1	ESTs, AA701352	
M/G1	SGK serum/glucocorticoid regulated kinase, AA424756	transport
M/G1	AMD1 S-adenosylmethionine decarboxylase 1, AA504772	
G2/M	KIAA0182, H05099	