

neg			pos				
ARCH1154_RFX_RFX2						— Nvec_vc1.1_XM_001623326.3	Calcium activated nucleotidase 1
						— Nvec_vc1.1_XM_001626082.3	
						— Nvec_vc1.1_XM_001626458.3	
						— Nvec_vc1.1_XM_001634717.3	N-acetyl-D-glucosamine kinase
						— Nvec_vc1.1_XM_048721044.1	Kinesin-like protein
						— Nvec_vc1.1_XM_032379506.2	SET:PRDM6
ARCH1520_ETV6_ETV7_ELF3						— Nvec_vc1.1_XM_032369963.2	
						— Nvec_vc1.1_XM_001631359.3	sterol_dehydrogenase:DHRS12
ARCH1562_zf_C2H2_ZNF334						— Nvec_vc1.1_XM_001640818.3	E3 ubiquitin-protein ligase MARC
						— Nvec_NVE9094	NEDD4-binding protein 2
						— Nvec_vc1.1_XM_001623768.3	Lysosomal thioesterase PPT2
						— Nvec_vc1.1_XM_001624599.3	60S ribosomal protein L24
						— Nvec_vc1.1_XM_001629013.3	Y_phosphatase:like:DUSP1/DUSP4/D
						— Nvec_vc1.1_XM_001632911.3	Golgi integral membrane protein
						— Nvec_vc1.1_XM_001641206.3	GNAS complex locus
						— Nvec_vc1.1_XM_032379058.2	RIO kinase 3 (Yeast)
						— Nvec_vc1.1_XM_032381795.2	Sodium/hydrogen exchanger 9
						— Nvec_vc1.1_XM_048724299.1	Protein DENND6B
						— Nvec_vc1.1_XM_048733370.1	
						— Nvec_vc1.1_XM_032381571.2	Carbonic anhydrase I
						— Nvec_vc1.1_XM_048730893.1	cDNA FLJ31473 fis, clone NT2NE20
						— Nvec_vc1.1_XM_001633090.3	Charged multivesicular body prot
						— Nvec_vc1.1_XM_001629132.3	40S ribosomal protein S16
ARCH2647						— Nvec_vc1.1_XM_001636538.3	Vacuolar protein sorting-associa
						— Nvec_vc1.1_XM_001638847.3	B9 domain-containing protein 2
						— Nvec_vc1.1_XM_001639748.3	Disrupted in renal carcinoma pro
						— Nvec_vc1.1_XM_001640352.3	HCG1994130
						— Nvec_vc1.1_XM_001640975.3	60S ribosomal protein L37a
						— Nvec_vc1.1_XM_032367462.2	60S ribosomal protein L10a
						— Nvec_vc1.1_XM_032367735.2	Piwi:PIWIL1/PIWIL3/PIWIL4
						— Nvec_vc1.1_XM_048723656.1	Cullin 1
						— Nvec_vc1.1_XM_048730672.1	Ankyrin repeat domain-containing
						— Nvec_vc1.1_XM_048731269.1	KH:FUBP1/FUBP3/KHSRP
						— Nvec_vc1.1_XM_001629210.3	F-box/LRR-repeat protein 8
						— Nvec_vc1.1_XM_032383104.2	Protein YIPF
						— Nvec_vc1.1_XM_048727269.1	28S ribosomal protein S30, mitoc
						— Nvec_vc1.1_XM_001641062.3	
						— Nvec_vc1.1_XM_032371473.2	High affinity cGMP-specific 3',5
						— Nvec_vc1.1_XM_048727040.1	Glycerophosphodiester phosphodie
ARCH2655_zf_C2H2_ZNF212							
ARCH2665_zf_C2H2_ZNF785						— Nvec_vc1.1_XM_032370047.2	PID:NOS1AP
ARCH395_HRB98DE						— Nvec_NVE14377	cDNA FLJ60705, highly similar to F-box/LRR-r
						— Nvec_NVE18637	
						— Nvec_NVE21518	
						— Nvec_NVE3414	
						— Nvec_NVE9713	
						— Nvec_v1g127536	Thiosulfate sulfurtransferase/rhodanese-lik
						— Nvec_v1g204395	Protein shisa-4
						— Nvec_v1g240440	
						— Nvec_v1g243618	
						— Nvec_v1g245805	Putative 60S ribosomal protein L39-like 5
						— Nvec_vc1.1_XM_001625275.3	Arf-GAP with coiled-coil, ANK re
						— Nvec_vc1.1_XM_001627627.3	
						— Nvec_vc1.1_XM_001628384.3	Pkinase:AC006486/GSK3A/GSK3B
						— Nvec_vc1.1_XM_001629004.3	cDNA FLJ56853, highly similar to
						— Nvec_vc1.1_XM_001629485.3	Cell division cycle 42 (GTP bind
						— Nvec_vc1.1_XM_001630351.3	RRM:like:ZCRB1:likeclu:1
						— Nvec_vc1.1_XM_001631508.3	YTH:YTHDC1
						— Nvec_vc1.1_XM_001632836.3	Replication factor C (Activator
						— Nvec_vc1.1_XM_001633942.3	Nucleoporin p54
						— Nvec_vc1.1_XM_001635214.3	
						— Nvec_vc1.1_XM_001635307.3	NNMT protein
						— Nvec_vc1.1_XM_001636541.3	Ras-related protein Rab-35
						— Nvec_vc1.1_XM_001638251.3	COesterase:like:NLGN1/NLGN2/NLGN
						— Nvec_vc1.1_XM_001638761.3	zf-C2H2.HG13.1:TRERF1/ZNF541
						— Nvec_vc1.1_XM_001639598.3	
						— Nvec_vc1.1_XM_001641906.3	Coiled-coil domain-containing pr
						— Nvec_vc1.1_XM_032362924.2	APEX nuclease (Apurinic/apyrimid
						— Nvec_vc1.1_XM_032363814.2	
						— Nvec_vc1.1_XM_032364464.2	cDNA FLJ33589 fis, clone BRAMY20
						— Nvec_vc1.1_XM_032366066.2	
						— Nvec_vc1.1_XM_032370541.2	SMAD_MH1.HG2.1:SMAD4
						— Nvec_vc1.1_XM_032375400.2	CTD-binding SR-like protein rA9
						— Nvec_vc1.1_XM_032375473.2	Myb.HG7.0:TADA2B
						— Nvec_vc1.1_XM_032377962.2	GPCR-rhodopsin:like:GPR176:likec
						— Nvec_vc1.1_XM_032379019.2	zf-CCCH:MBNL1/MBNL2/MBNL3
						— Nvec_vc1.1_XM_032380477.2	V-type proton ATPase subunit S1
						— Nvec_vc1.1_XM_032381227.2	cDNA FLJ50631, highly similar to
						— Nvec_vc1.1_XM_032382661.2	Fukutin related protein
						— Nvec_vc1.1_XM_048719451.1	
						— Nvec_vc1.1_XM_048721748.1	
						— Nvec_vc1.1_XM_048724582.1	Phosphatase and tensin-like prot
						— Nvec_vc1.1_XM_048724864.1	Elav1
						— Nvec_vc1.1_XM_048725472.1	
						— Nvec_vc1.1_XM_048725680.1	Centrosomal protein of 104 kDa
						— Nvec_vc1.1_XM_048726248.1	Phosphatase and actin regulator
						— Nvec_vc1.1_XM_048726512.1	
						— Nvec_vc1.1_XM_048728255.1	
						— Nvec_vc1.1_XM_048729664.1	Tetraspanin
						— Nvec_vc1.1_XM_048729671.1	cDNA FLJ55635, highly similar to
						— Nvec_vc1.1_XM_001623052.3	Uncharacterized protein C1orf50
						— Nvec_vc1.1_XM_001628819.3	cDNA FLJ51881, highly similar to
						— Nvec_vc1.1_XM_001630781.3	
						— Nvec_vc1.1_XM_001633221.3	Homer:like:ENAH/EVL/SPRED1/SPRED
						— Nvec_vc1.1_XM_001637226.3	Mannose-P-dolichol utilization d
						— Nvec_vc1.1_XM_001638404.2	Activating signal cointegrator 1
						— Nvec_vc1.1_XM_001641405.3	
						— Nvec_vc1.1_XM_032362411.2	ion-NaKCa:KCNH3/KCNH4/KCNH8
						— Nvec_vc1.1_XM_032363255.2	T-cell immunomodulatory protein
						— Nvec_vc1.1_XM_032363896.2	Filamin-A
						— Nvec_vc1.1_XM_032380187.2	ATP-binding cassette sub-family
						— Nvec_vc1.1_XM_032382958.2	Protein phosphatase 1 regulatory
						— Nvec_vc1.1_XM_032385226.2	Histone deacetylase complex subu
						— Nvec_vc1.1_XM_032385496.2	HLH.HG1.21:PTF1A
						— Nvec_vc1.1_XM_048724767.1	F-box/WD repeat-containing prote
						— Nvec_vc1.1_XM_048729375.1	Calmodulin
						— Nvec_vc1.1_XM_001633102.3	Palmitoyltransferase
						— Nvec_vc1.1_XM_048732170.1	SET:KMT2A/KMT2B
						— Nvec_vc1.1_XM_001629423.3	7-methylguanosine phosphate-spec
						— Nvec_vc1.1_XM_001640456.3	Desumoylating isopeptidase 1
						— Nvec_vc1.1_XM_032364202.2	
						— Nvec_vc1.1_XM_032367105.2	Prickle-like protein 2
						— Nvec_vc1.1_XM_032375602.2	Insm1
						— Nvec_vc1.1_XM_032384465.2	Probable sodium-coupled neutral
						— Nvec_vc1.1_XM_032385246.1	
						— Nvec_vc1.1_XM_032387437.2	
						— Nvec_v1g204213	
						— Nvec_vc1.1_XM_001633494.3	Tigger transposable element-deri
						— Nvec_vc1.1_XM_001635410.3	
						— Nvec_vc1.1_XM_032362457.2	INVS protein
						— Nvec_vc1.1_XM_032363386.2	Ankyrin-2
						— Nvec_vc1.1_XM_032372078.2	zf-C4_Nuclear_receptors.HG1.29:N
						— Nvec_vc1.1_XM_032375586.2	Mitochondrial basic amino acids
						— Nvec_vc1.1_XM_032381586.2	zf-C2H2.HG4.31:like:KLF5:likeclu
						— Nvec_vc1.1_XM_032382907.2	
						— Nvec_vc1.1_XM_032384295.2	FA_desaturase:SCD5
						— Nvec_vc1.1_XM_048722945.1	
						— Nvec_vc1.1_XM_048724522.1	
						— Nvec_vc1.1_XM_048726933.1	
						— Nvec_vc1.1_XM_048727988.1	cDNA FLJ75589, highly similar to
						— Nvec_vc1.1_XM_048729959.1	Glycogen phosphorylase, muscle f
ARCH512_zf_C2H2_ZSCAN16						— Nvec_vc1.1_XM_001634007.3	Tripartite motif-containing prot
						— Nvec_vc1.1_XM_032383224.2	Breast carcinoma amplified seque
						— Nvec_vc1.1_XM_001641734.3	
						— Nvec_vc1.1_XM_032386711.2	GPCR-rhodopsin:like:GPR27/GPR50/
Elav_neg			Elav_pos				
Fox_neg			Fox_pos				
Ncol_neg			Ncol_pos				