

		neg	pos	
ARCH1262_zf_C2H2_ZNF713				— Nvec_vc1.1_XM_001639285.3 KDEL motif–containing protein 1
				— Nvec_vc1.1_XM_032381880.2 Epididymis tissue sperm binding
				— Nvec_vc1.1_XM_032362392.2 Laminin:like:CNTNAP1/CNTNAP2/CNT
				— Nvec_vc1.1_XM_032364563.2
ARCH1681_POU1F1_POU3F2_POU3F1_POU3F4				— Nvec_vc1.1_XM_001624579.3 Prefoldin subunit 4
				— Nvec_vc1.1_XM_001626362.3
				— Nvec_vc1.1_XM_032362924.2 APEX nuclease (Apurinic/apyrimid
				— Nvec_vc1.1_XM_032371836.2 Core–binding factor, beta subuni
				— Nvec_vc1.1_XM_001636807.3 Cornifelin
				— Nvec_vc1.1_XM_001641721.3
ARCH1792_zf_C4_Nuclear_receptors_ftz_f1				— Nvec_vc1.1_XM_032371336.2 zf–CCCH:ZFP36L1/ZFP36L2
				— Nvec_vc1.1_XM_001632345.3 cDNA FLJ33079 fis, clone TRACH20
				— Nvec_vc1.1_XM_001637929.3 RRM:like:RBM23/RBM39/SRSF2/SRSF8
				— Nvec_vc1.1_XM_048729708.1 Pyruvate carboxylase, mitochondr
ARCH1860_islike_FOXL1_FOXC2_Foxc1				— Nvec_vc1.1_XM_048733452.1 Importin–9
				— Nvec_vc1.1_XM_001630762.3 60S ribosomal protein L13a
				— Nvec_vc1.1_XM_001638022.3 Sjogren’s syndrome nuclear autoa
				— Nvec_vc1.1_XM_001641219.2
				— Nvec_vc1.1_XM_032385706.2 FoxQ2d
				— Nvec_vc1.1_XM_048726526.1
ARCH2545_PPARG				— Nvec_vc1.1_XM_001632619.3 cDNA FLJ57219, highly similar to
				— Nvec_vc1.1_XM_048726685.1 Replication factor C subunit 2
				— Nvec_vc1.1_XM_032386221.2
				— Nvec_vc1.1_XM_048723326.1
				— Nvec_vc1.1_XM_001622235.3 G1/S–specific cyclin–D2
				— Nvec_vc1.1_XM_001630513.3 Crk–like protein
ARCH2633_HLH_SREBF2				— Nvec_vc1.1_XM_048719744.1 Solute carrier family 25 member
				— Nvec_vc1.1_XM_048722879.1 Catalase
				— Nvec_vc1.1_XM_001636422.3 sterol_dehydrogenase:HSD17B10
				— Nvec_vc1.1_XM_001631087.3
				— Nvec_vc1.1_XM_001634707.3 CTRL protein
				— Nvec_NVE11610
ARCH2647				— Nvec_vc1.1_XM_001628844.3 Epididymis secretory sperm bindi
				— Nvec_vc1.1_XM_032365481.2 Zinc finger SWIM domain–containi
				— Nvec_vc1.1_XM_032376545.2 cDNA FLJ54017, highly similar to
				— Nvec_vc1.1_XM_048733321.1 Testis derived transcript (3 LIM
				— Nvec_vc1.1_XM_048733365.1 cDNA FLJ58466, highly similar to
				— Nvec_NVE15515
ARCH2655_zf_C2H2_ZNF212				— Nvec_vc1.1_XM_001626816.3 GDP–fucose protein O–fucosyltran
				— Nvec_vc1.1_XM_032370093.2 Death domain–containing protein
				— Nvec_vc1.1_XM_001640933.3 cDNA, FLJ79193, highly similar t
				— Nvec_vc1.1_XM_001641927.3 WD repeat–containing protein 5B
				— Nvec_vc1.1_XM_048724761.1 6–phosphogluconate dehydrogenase
				— Nvec_vc1.1_XM_001629132.3 40S ribosomal protein S16
ARCH2665_zf_C2H2_ZNF785				— 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
ARCH2671_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032367735.2 Piwi:PIWIL1/PIWIL3/PIWIL4
				— Nvec_vc1.1_XM_032384295.2 FA_desaturase:SCD5
				— 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
ARCH268_EBF1				— 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
				— Nvec_vc1.1_XM_032370047.2 PID:NOS1AP
ARCH2696_zf_C2H2_ZNF785				— Nvec_vc1.1_XM_001630052.3 Protein LAP2
				— Nvec_vc1.1_XM_001639073.3 Major facilitator superfamily do
				— Nvec_vc1.1_XM_032378226.2 Vacuolar protein sorting–associa
				— Nvec_vc1.1_XM_032387016.2
				— Nvec_vc1.1_XM_048720203.1 [Protein ADP–ribosylarginine] hy
				— Nvec_vc1.1_XM_032381112.2 SAM:BFAR
ARCH2700_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032381807.2 PMCA:AC010616/ATP12A/ATP1A1/ATP1
				— Nvec_vc1.1_XM_032384009.2 Afadin– and alpha–actinin–bindin
				— Nvec_NVE11782
				— Nvec_NVE3036
				— Nvec_NVE9810
				— Nvec_v1g107040 Copper transport protein ATOX1
ARCH2701_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001619828.3 FAD–dependent oxidoreductase dom
				— Nvec_vc1.1_XM_001622638.3 Very–long–chain (3R)–3–hydroxyac
				— Nvec_vc1.1_XM_001623615.3
				— Nvec_vc1.1_XM_001625350.2 B–cell lymphoma/leukemia 10
				— Nvec_vc1.1_XM_001627312.3
				— Nvec_vc1.1_XM_001628389.3
ARCH2702_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001628947.3 40S ribosomal protein S18
				— Nvec_vc1.1_XM_001633594.3
				— Nvec_vc1.1_XM_001636447.3 zf–C2H2.HG6.0:GLI1/GLI2/GLI3/GLI
				— Nvec_vc1.1_XM_001638182.3
				— Nvec_vc1.1_XM_001639736.3 cDNA FLJ51907, highly similar to
				— Nvec_vc1.1_XM_001640586.3
ARCH2703_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032362308.2 Collagen:like:COLEC10/COLEC11/FC
				— Nvec_vc1.1_XM_032363186.2 Signal transducing adapter molec
				— Nvec_vc1.1_XM_032371460.2 Sarcosine dehydrogenase, mitoch
				— Nvec_vc1.1_XM_032371613.2 cDNA FLJ76474, highly similar to
				— Nvec_vc1.1_XM_032372609.2 Pkinase:OXSR1/STK39
				— Nvec_vc1.1_XM_032373490.2 DTW domain–containing protein 1
ARCH2704_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032375518.2 PMCA:ATP2C1/ATP2C2
				— Nvec_vc1.1_XM_032378379.2 wnt:WNT11
				— Nvec_vc1.1_XM_048720266.1
				— Nvec_vc1.1_XM_048724529.1 Anchor protein
				— Nvec_vc1.1_XM_048727447.1
				— Nvec_vc1.1_XM_048729504.1 cDNA FLJ61218, highly similar to
ARCH2705_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048731331.1 cDNA FLJ77988
				— Nvec_vc1.1_XM_048734658.1 Solute carrier family 20 (Phosph
				— Nvec_NVE25375 Guanine nucleotide–binding protein G(I)/G(S)
				— Nvec_vc1.1_XM_001631707.3 Epididymis secretory protein Li
				— Nvec_vc1.1_XM_001632516.3 Monoamine oxidase A
				— Nvec_vc1.1_XM_001635133.3 Fibroblast growth factor recepto
ARCH2706_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032385176.2 TIR:like:IL1R1/IL1RAP/IL1RAPL1/I
				— Nvec_vc1.1_XM_048725791.1 Sugar_tr:like:SLC22A1/SLC22A2/SL
				— Nvec_vc1.1_XM_001640624.3 N–acylneuraminate–9–phosphatase
				— Nvec_NVE18754 Cell division cycle 26
				— Nvec_vc1.1_XM_032370349.2 SWIRM:like:KDM1A:likeclu:1
				— Nvec_vc1.1_XM_032384105.2 PDZ:SHANK1/SHANK2/SHANK3
ARCH2707_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048719974.1 Protein asteroid homolog 1
				— Nvec_vc1.1_XM_001629587.3 cDNA FLJ53033, highly similar to
				— Nvec_vc1.1_XM_001633809.3 Pkinase:TNK1/TNK2
				— Nvec_vc1.1_XM_001634654.3 Chromosome 22 open reading frame
				— Nvec_vc1.1_XM_001638819.3 Protein dpy–30 homolog
				— Nvec_vc1.1_XM_032383342.2 Protein eva–1 homolog C
ARCH2708_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048722400.1 GPCR–rhodopsin:NA
				— Nvec_v1g206505
				— Nvec_vc1.1_XM_001622895.3
				— Nvec_vc1.1_XM_001632169.3 Ras homolog gene family, member
				— Nvec_vc1.1_XM_001632488.3 Catechol O–methyltransferase dom
				— Nvec_vc1.1_XM_001633368.3 E3 ubiquitin–protein ligase TRIM
ARCH2709_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032361818.2 BTB:ABTB1
				— Nvec_vc1.1_XM_032362029.2 Histone:NA
				— Nvec_vc1.1_XM_032364850.2 Protein C12orf4
				— Nvec_vc1.1_XM_032365233.2 Lethal(2) giant larvae protein h
				— Nvec_vc1.1_XM_032370462.2 General transcription factor 3C
				— Nvec_vc1.1_XM_032382055.2 Glyceraldehyde–3–phosphate dehyd
ARCH2710_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032384172.2 zf–met.HG20.0:NA
				— Nvec_vc1.1_XM_032386449.2 cDNA FLJ16604 fis, clone TEST140
				— Nvec_vc1.1_XM_032387048.2
				— Nvec_vc1.1_XM_048723054.1
				— Nvec_vc1.1_XM_048724922.1 TBC1 domain family, member 15
				— Nvec_vc1.1_XM_048732565.1 E3 ubiquitin–protein ligase Haka
ARCH2711_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048733302.1 HLH.HG3.4:MITF/TFE3/TFEB/TFEC
				— Nvec_vc1.1_XM_001622588.3
				— Nvec_vc1.1_XM_001633727.3 Protein phosphatase 1H
				— Nvec_vc1.1_XM_001637149.3 RAP1A, member of RAS oncogene fa
				— Nvec_vc1.1_XM_001639489.3 Migration–inducing gene 14
				— Nvec_vc1.1_XM_032381270.2 Transcriptional repressor scratc
ARCH2712_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_032386835.2 cDNA FLJ20362 fis, clone HEP1698
				— Nvec_vc1.1_XM_048719723.1 cDNA FLJ55674, highly similar to
				— Nvec_vc1.1_XM_048729023.1 Pkinase:STK32A/STK32B/STK32C
				— Nvec_vc1.1_XM_048729913.1
				— Nvec_vc1.1_XM_001627423.3 HMGBbox_Sox.HG1.4:BBX/CIC
				— Nvec_vc1.1_XM_032383201.2 AN1–type zinc finger protein 2A
ARCH2713_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048728942.1 MMS19 nucleotide excision repair
				— Nvec_vc1.1_XM_001626458.3
				— Nvec_vc1.1_XM_001635684.3 Cyclin–dependent kinase 3
				— Nvec_vc1.1_XM_001636512.3 Dolichyl–diphosphooligosaccharid
				— Nvec_vc1.1_XM_032380235.2 Flavin containing monooxygenase
				— Nvec_vc1.1_XM_048727840.1 Putative uncharacterized protein
ARCH2714_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001638761.3 zf–C2H2.HG13.1:TRERF1/ZNF541
				— Nvec_vc1.1_XM_001624118.3 Glucose–6–phosphatase 2
				— Nvec_vc1.1_XM_032373907.2 Isocitrate dehydrogenase [NAD] s
				— Nvec_vc1.1_XM_001623589.3 Peptidyl–prolyl cis–trans isomer
				— Nvec_vc1.1_XM_048720142.1 Uncharacterized protein KIAA1467
				— Nvec_vc1.1_XM_048722345.1
ARCH2715_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_048732272.1
				— Nvec_vc1.1_XM_001637171.3 Solute carrier family 35, member
				— Nvec_NVE4959
				— Nvec_vc1.1_XM_032365386.2 F–box/LRR–repeat protein 13
				— Nvec_vc1.1_XM_048720479.1
				— Nvec_vc1.1_XM_001622544.3
ARCH2716_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001629168.3 Protein disulfide–isomerase A6
				— Nvec_vc1.1_XM_001630079.3 Tubulin, alpha 2
				— Nvec_vc1.1_XM_032371411.2
				— Nvec_vc1.1_XM_032372812.2 Uncharacterized membrane protein
				— Nvec_vc1.1_XM_048727123.1
				— Nvec_NVE24146
ARCH2717_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001629557.3 2–hydroxyacyl–CoA lyase 1
				— Nvec_vc1.1_XM_001631329.3 TIR:like:SARM1:likeclu:26
				— Nvec_vc1.1_XM_001637508.3 IRF.HG1.4:IRF1/IRF2
				— Nvec_vc1.1_XM_032363779.2
				— Nvec_vc1.1_XM_032376717.2 Tubulin polyglutamylase TTL11
				— Nvec_vc1.1_XM_001628740.3 Cathepsin L
ARCH2718_zf_C2H2_ZIM2				— Nvec_vc1.1_XM_001635053.3
				— Nvec_vc1.1_XM_032375586.2 Mitochondrial basic amino acids
				— Nvec_vc1.1_XM_048729407.1 Dual specificity protein phosph
				— Nvec_vc1.1_XM_032378617.2 Potassium channel subfamily K me
				— Nvec_vc1.1_XM_048720853.1 Eukaryotic elongation factor 2 k
				— Nvec_vc1.1_XM_001630119.3 Glutamate dehydrogenase 2, mitoc
ARCH940_HMGBbox_Sox_SOX7_Sox21b				
	Elav_neg	Fox_neg	Ncol_neg	Elav_pos
				Fox_pos
				Ncol_pos