APÊNDICE B – Tabela 2

 $Tabela\ 2\ -\ Total\ de\ transcritos\ anotados\ das\ três\ espécies\ estudadas,\ identificados\ como\ diferencialmente\ expressos.$

Identificação TRINITY_DN16283_c3_g1_i11	Anotação	Fold change
•		
	Uromodulin	-14,53
TRINITY_DN12684_c1_g2_i5	Venom phosphodiesterase 1	-12,53
TRINITY_DN15138_c1_g1_i2	Peroxisome proliferator-activated receptor alpha	-10,91
TRINITY_DN15261_c2_g4_i2	Sterol 26-hydroxylase, mitochondrial	-10,66
TRINITY_DN13545_c2_g1_i19	Probable palmitoyltransferase ZDHHC8	-10,59
TRINITY_DN13986_c0_g2_i4	Calpain-5	-10,55
TRINITY_DN14327_c3_g1_i10	Pleckstrin homology-like domain family B member 2	-10,54
TRINITY_DN16482_c2_g2_i6	Glycine cleavage system H protein, mitochondrial	-10,50
TRINITY_DN17133_c2_g1_i8	Sterile alpha motif domain-containing protein 9	-10,26
TRINITY_DN11794_c0_g1_i3	Protein tweety homolog 3	-10,21
TRINITY_DN16614_c3_g3_i6	Histone deacetylase 7	-10,21
TRINITY_DN12227_c1_g1_i5	General vesicular transport factor p115	-10,02
TRINITY_DN12771_c1_g1_i8	Mitochondrial basic amino acids transporter	-9,81
TRINITY_DN14905_c3_g1_i11	CMP-N-acetylneuraminate-beta-galactosamide-alpha-2,3-sialy	-9,73
TRINITY_DN12727_c2_g2_i4	Peroxisomal acyl-coenzyme A oxidase 1	-9,72
TRINITY_DN13654_c3_g1_i5	ATP synthase mitochondrial F1 complex assembly factor 1	-9,62
TRINITY_DN15026_c0_g1_i19	Nucleolar protein 6	-9,42
TRINITY_DN12925_c2_g2_i1	Copper chaperone for superoxide dismutase	-9,41
Transcritos upregulated	D.L. (ADD. 11. A. A. A.	0.40
TRINITY_DN16728_c2_g1_i6	Poly [ADP-ribose] polymerase 9	9,49
TRINITY_DN13237_c1_g2_i6	Ras-related protein Rab-11B	9,64
TRINITY_DN13211_c1_g1_i12	Alanine aminotransferase 2-like	9,85
TRINITY_DN14793_c0_g1_i14	Ubiquinone biosynthesis monooxygenase COQ6, mitochondrial	9,86
TRINITY_DN16283_c3_g1_i14	Deleted in malignant brain tumors 1 protein	9,95
TRINITY_DN14812_c2_g3_i1	Receptor-type tyrosine-protein phosphatase kappa	10,01
TRINITY_DN13894_c0_g1_i16	Inter-alpha-trypsin inhibitor heavy chain H2	10,02
TRINITY_DN16670_c0_g1_i2	RalBP1-associated Eps domain-containing protein 2	10,09
TRINITY_DN13596_c3_g2_i13	Serinepyruvate aminotransferase	10,10
TRINITY_DN15412_c3_g1_i4	Paired amphipathic helix protein Sin3a	10,11
TRINITY_DN17260_c3_g1_i4	Amine sulfotransferase	10,15
TRINITY_DN12088_c0_g2_i9	Isocitrate dehydrogenase [NADP], mitochondrial	10,26
TRINITY_DN15412_c3_g1_i3	Paired amphipathic helix protein Sin3a	10,27
TRINITY_DN14573_c4_g2_i1	NAD kinase	10,36
TRINITY_DN13038_c2_g4_i2	Growth arrest and DNA damage-inducible protein GADD45 ga	10,37
TRINITY_DN15650_c0_g1_i6	Cation-independent mannose-6-phosphate receptor	10,49
TRINITY_DN14905_c3_g1_i8	CMP-N-acetylneuraminate-beta-galactosamide-alpha-2,3-sialy	10,54
TRINITY_DN13160_c3_g1_i7	Glycerol-3-phosphate dehydrogenase [NAD(+)], cytoplasmic	10,55
TRINITY_DN13210_c3_g1_i13	Scavenger receptor class F member 1 Claudin-10	10,67
TRINITY_DN16616_c2_g1_i1		10,68
TRINITY_DN16205_c1_g1_i23	Complement C1q-like protein 2	11,10
TRINITY_DN13172_c1_g2_i6	E3 ubiquitin-protein ligase MARCH5	11,16
TRINITY_DN15048_c1_g1_i6	Cholinephosphotransferase 1	12,10
TRINITY_DN15618_c2_g1_i10	Adenosylhomocysteinase B	12,55
TRINITY_DN16513_c0_g2_i4	Heterogeneous nuclear ribonucleoprotein A0	12,73
TRINITY_DN16662_c0_g1_i4	Adenylate kinase 2, mitochondrial	12,81
TRINITY_DN12616_c3_g1_i6 P. semilunaris	Rho-related GTP-binding protein Rho6	12,90
Transcritos downregulated		
TRINITY_DN12784_c3_g1_i3	Heterogeneous nuclear ribonucleoprotein Q	-11,70
TRINITY_DN11601_c0_g8_i4	Eukaryotic translation initiation factor 2 subunit 1	-11,70
TRINITY_DN13834_c3_g4_i1	Group XIIB secretory phospholipase A2-like protein	-11,50 -11,50
1 KIIVII 1 _DIVI 3034_C3_g4_II		
TRINITY DNIAROS AS AL :12		
TRINITY_DN14802_c2_g1_i12 TRINITY_DN12476_c1_g4_i11	Complement C1q-like protein 4 Cullin-9	-11,47 -11,05

TRINITY_DN13821_c5_g3_i7	Dysferlin	-10,97
TRINITY_DN13051_c2_g1_i2	Myocardial zonula adherens protein	-10,94
TRINITY_DN13629_c2_g1_i28	Vitamin K-dependent protein C	-10,93
TRINITY_DN12945_c6_g2_i16	KAT8 regulatory NSL complex subunit 3	-10,55
TRINITY_DN14008_c4_g1_i9	Membrane-associated guanylate kinase, WW and PDZ domain-	-10,49
TRINITY_DN11861_c4_g3_i1	Forkhead box protein O3	-10,39
TRINITY_DN14610_c1_g2_i10	Cation-independent mannose-6-phosphate receptor	-10,26
TRINITY_DN7773_c0_g1_i1	LINE-1 retrotransposable element ORF2 protein	-10,22
TRINITY_DN14508_c3_g1_i8	Cell cycle control protein 50A	-10,09
TRINITY_DN12259_c4_g1_i5	Transmembrane 6 superfamily member 2	-10,05
TRINITY_DN11476_c4_g1_i9	Rho guanine nucleotide exchange factor 12	-10,01
TRINITY_DN14236_c1_g1_i7	Nicotinate-nucleotide pyrophosphorylase [carboxylating]	-10,00
TRINITY_DN13811_c2_g1_i2	Zinc finger and BTB domain-containing protein 1	-9,93
TRINITY_DN12862_c4_g2_i4	von Willebrand factor (Fragment)	-9,87
TRINITY_DN12476_c1_g4_i10	Cullin-9	-9,77
TRINITY_DN10909_c0_g1_i2	Lysosomal protective protein	-9,70
TRINITY_DN13638_c1_g6_i1	TATA box-binding protein-associated factor RNA polymerase I	-9,55
TRINITY_DN13029_c0_g1_i14	Epoxide hydrolase 1	-9,52
TRINITY_DN10996_c0_g1_i8	E3 ubiquitin-protein ligase ubr3	-9,50
TRINITY_DN11598_c3_g1_i1	Aryl hydrocarbon receptor nuclear translocator-like protein 1	-9,36
TRINITY_DN14023_c0_g2_i6	SUN domain-containing	-9,35
TRINITY_DN12820_c4_g4_i1	Ubiquitin-associated protein 1	-9,30
TRINITY_DN13043_c5_g2_i3	Serine/arginine-rich splicing factor 9	-9,30
TRINITY_DN11564_c0_g1_i3	Leucine-rich repeats and immunoglobulin-like domains protein	-9,29
TRINITY_DN12541_c1_g2_i3	ATP-binding cassette sub-family C member 9	-9,27
TRINITY_DN12872_c8_g2_i2	CUGBP Elav-like family member 2	-9,27
TRINITY_DN14312_c0_g3_i4	Glutamate dehydrogenase 1, mitochondrial	-9,26
TRINITY_DN14692_c0_g3_i3	Histone H3.3 (Fragment)	-9,19
TRINITY_DN13638_c1_g6_i2	TATA box-binding protein-associated factor RNA polymerase I	-9,14
TRINITY_DN11281_c0_g1_i4	Ubiquitin-protein ligase E3A	-9,12
TRINITY_DN13653_c5_g1_i8	Alpha-mannosidase 2C1	-9,11
TRINITY_DN11732_c0_g1_i6	Protein Smaug homolog 2	-9,10
TRINITY_DN11911_c7_g2_i1	Arginine/serine-rich coiled-coil protein 2	-9,10
TRINITY_DN13774_c2_g3_i1	Dual specificity mitogen-activated protein kinase kinase 7	-9,09
TRINITY_DN12668_c5_g1_i4	SAM and SH3 domain-containing protein 1	-9,08
TRINITY_DN11406_c4_g2_i12	Dystonin	-9,02
TRINITY_DN12477_c3_g3_i10	Thyroid hormone receptor-associated protein 3	-9,02
TRINITY_DN11926_c1_g3_i5	Pre-rRNA-processing protein TSR1 homolog	-9,00
TRINITY_DN10624_c0_g1_i7	DnaJ homolog subfamily C member 3	-8,97
TRINITY_DN12221_c2_g1_i8	F-box DNA helicase 1	-8,95
TRINITY_DN13190_c3_g1_i3	Complement C3 (Fragment) White pricing a graph and the pricing I but dealers a 10	-8,93
TRINITY_DN11347_c1_g4_i2 TRINITY_DN14868_c3_g1_i1	Ubiquitin carboxyl-terminal hydrolase 19 Enoyl CoA dolta isomerase 2, mitochondrial	-8,91 -8,87
TRINITY_DN14312_c0_g3_i15	Enoyl-CoA delta isomerase 2, mitochondrial Glutamate dehydrogenase 1, mitochondrial	-8,84
TRINITY_DN10936_c0_g1_i2	Crooked neck-like protein 1	-8,84
TRINITY_DN11275_c2_g2_i4	UPF0472 protein C16orf72 homolog	-8,83
TRINITY_DN11996_c5_g1_i14	Ankyrin repeat domain-containing protein 22	-8,80
TRINITY_DN13941_c2_g3_i7	Collagen alpha-1(IV) chain	-8,78
TRINITY_DN13941_c2_g3_i2	Collagen alpha-1(IV) chain	-8,75
TRINITY_DN10938_c0_g2_i7	Aryl hydrocarbon receptor nuclear translocator	-8,74
TRINITY_DN12668_c5_g1_i2	SAM and SH3 domain-containing protein 1	-8,71
TRINITY_DN14846_c3_g2_i2	Next to BRCA1 gene 1 protein	-8,63
TRINITY_DN13560_c2_g1_i10	Regulator of microtubule dynamics protein 2	-8,63
TRINITY_DN11218_c4_g3_i3	Circadian locomoter output cycles protein kaput	-8,61
TRINITY_DN12711_c2_g3_i2	Copper-transporting ATPase 2	-8,56
TRINITY_DN13770_c3_g1_i13	Transmembrane protein 268	-8,53
TRINITY_DN12893_c3_g2_i4	5'-nucleotidase	-8,49
TRINITY_DN14780_c2_g3_i1	Nicotinate-nucleotide pyrophosphorylase [carboxylating] (Fras	-8,49
TRINITY_DN13815_c1_g2_i7	Selenoprotein N	-8,48
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TRINITY_DN12764_c1_g1_i12	Annexin A1	-8,47
TRINITY_DN12404_c0_g1_i7	Muscleblind-like protein 1	-8,45
TRINITY_DN9791_c0_g1_i7	Glucocorticoid modulatory element-binding protein 2	-8,45
TRINITY_DN13394_c2_g13_i9	CCAAT/enhancer-binding protein gamma	-8,43
TRINITY_DN12317_c2_g1_i4	Rho guanine nucleotide exchange factor 25	-8,42
TRINITY_DN11577_c1_g1_i3	Cysteine and histidine-rich domain-containing protein 1	-8,42
TRINITY_DN14508_c3_g1_i2	Cell cycle control protein 50A	-8,41
TRINITY_DN11409_c0_g2_i4	Mitochondrial proton/calcium exchanger protein	-8,38
TRINITY_DN14140_c1_g1_i15	Thymidine phosphorylase	-8,34
TRINITY_DN14754_c2_g5_i16	Serine/threonine-protein kinase ULK2	-8,32
TRINITY_DN11830_c3_g1_i1	UPF0565 protein C2orf69 homolog	-8,26
TRINITY_DN12489_c5_g2_i11	BRISC and BRCA1-A complex member 1	-8,22
TRINITY_DN8254_c0_g1_i4	Rho GTPase-activating protein 11B	-8,21
TRINITY_DN13620_c0_g1_i1	Sodium-dependent multivitamin transporter	-8,20
TRINITY_DN13411_c0_g1_i19	Magnesium transporter MRS2 homolog, mitochondrial	-8,19
TRINITY_DN14610_c1_g2_i15	Cation-independent mannose-6-phosphate receptor	-8,16
TRINITY_DN12711_c2_g3_i27	Copper-transporting ATPase 2	-8,11
TRINITY_DN13680_c2_g2_i3	Calcium/calmodulin-dependent protein kinase type II subunit gc	-8,11
TRINITY_DN10996_c0_g1_i6	E3 ubiquitin-protein ligase ubr3	-8,07
TRINITY_DN12124_c1_g1_i26	Feline leukemia virus subgroup C receptor-related protein 2	-8,05
TRINITY_DN10897_c0_g1_i5	Cholinephosphotransferase 1	-8,03
TRINITY_DN11395_c0_g1_i7	FYVE, RhoGEF and PH domain-containing protein 6	-8,03
TRINITY_DN13630_c0_g1_i13	Ankyrin repeat and SAM domain-containing protein 6	-8,01
TRINITY_DN13713_c0_g4_i5	60S ribosomal protein L7-like 1	-8,00
TRINITY_DN11418_c4_g1_i4	MICAL-like protein 2	-7,96
TRINITY_DN12473_c4_g1_i4	D-erythrulose reductase	-7,96
TRINITY_DN13629_c2_g1_i10	Vitamin K-dependent protein C	-7,94
TRINITY_DN11237_c0_g1_i12	NACHT, LRR and PYD domains-containing protein 1a	-7,88
TRINITY_DN13858_c1_g1_i3	Transcriptional protein SWT1	-7,79
TRINITY_DN14758_c3_g2_i2	Pseudouridine-5'-phosphate glycosidase	-7,68
TRINITY_DN11377_c3_g1_i1	Kelch-like ECH-associated protein 1	-7,66
TRINITY_DN12837_c1_g2_i8	cAMP-dependent protein kinase catalytic subunit	-7,59
TRINITY_DN11324_c4_g1_i4	Flavin-containing monooxygenase FMO GS-OX1	-7,58
TRINITY_DN14236_c1_g1_i1	Nicotinate-nucleotide pyrophosphorylase [carboxylating]	-7,50
TRINITY_DN14256_c0_g1_i16	Vang-like protein 1	-7,06
TRINITY_DN13255_c1_g1_i3	Cyclin-dependent kinase 19	-6,99
TRINITY_DN11411_c4_g2_i3	Plakophilin-1	-6,89
TRINITY_DN11601_c0_g1_i7	Muscleblind-like protein 2a	-6,75
TRINITY_DN12342_c3_g2_i10	F-box only protein 31	-6,72
TRINITY_DN13029_c0_g1_i18	Epoxide hydrolase 1	-6,38
TRINITY_DN14321_c4_g2_i1	Protein Wiz	-6,35
TRINITY_DN14223_c0_g2_i8	Voltage-dependent anion-selective channel protein 2	-6,29
TRINITY_DN12052_c4_g2_i2	Neurabin-1	-6,02
TRINITY_DN12751_c0_g1_i4	Heterogeneous nuclear ribonucleoprotein L	-5,82
TRINITY_DN11455_c3_g1_i8	KH homology domain-containing protein 4	-5,72
TRINITY_DN13886_c2_g1_i4	Acyl-CoA synthetase family member 2, mitochondrial	-5,07
TRINITY_DN12006_c2_g1_i2	Armadillo repeat-containing protein 1	-4,95
TRINITY_DN12645_c2_g1_i20	Cold shock domain-containing protein E1	-4,83
TRINITY_DN13986_c1_g2_i5	Amine sulfotransferase	-4,15
TRINITY_DN13628_c3_g6_i5	Replication protein A 70 kDa DNA-binding subunit	-3,97
TRINITY_DN13746_c3_g1_i2	Integrin beta-1	-3,95
TRINITY_DN11092_c0_g1_i2	Serpin H1	-3,75
TRINITY_DN14624_c0_g1_i16	Acyl-coenzyme A thioesterase 4	-3,45
TRINITY_DN12117_c4_g2_i2	GATA zinc finger domain-containing protein 1	-3,31
TRINITY_DN13968_c2_g1_i7	Leucine-rich PPR motif-containing protein, mitochondrial	-2,92
TRINITY_DN12031_c3_g1_i7	Microtubule-associated tumor suppressor 1 homolog A	-2,90
TRINITY_DN13875_c0_g2_i7	Alpha-actinin-1	-2,72
TRINITY_DN7332_c0_g1_i2	DnaJ homolog subfamily A member 4	-2,54
TRINITY_DN12629_c0_g1_i1	GTP-binding protein SAR1b	-2,44

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TRINITY_DN13892_c0_g3_i1	Heat shock protein HSP 90-alpha	-2,22
TRINITY_DN13545_c1_g5_i3	Heterogeneous nuclear ribonucleoprotein C	-2,21
TRINITY_DN14663_c1_g5_i1	Transposable element Tcb2 transposase	-2,00
Transcritos upregulated	Donation and in the Management of the Property	2.10
TRINITY_DN13854_c2_g3_i5	Protein arginine N-methyltransferase 1-B	2,10
TRINITY_DN14853_c2_g1_i4	L-rhamnose-binding lectin SML	2,14
TRINITY_DN10562_c0_g1_i2	Ornithine decarboxylase	2,18
TRINITY_DN11435_c6_g2_i5	TATA-binding protein-associated factor 2N	2,46
TRINITY_DN14842_c0_g1_i6	Solute carrier family 22 member 1	2,90
TRINITY_DN10600_c0_g1_i1	ATP-binding cassette sub-family F member 2	3,09
TRINITY_DN14784_c1_g2_i8	Protein SSXT	3,36
TRINITY_DN12825_c4_g1_i1	Calcium-activated potassium channel subunit alpha-1	4,75
TRINITY_DN13787_c1_g1_i11	Enhancer of polycomb homolog 1	5,20
TRINITY_DN10916_c0_g1_i3	CCR4-NOT transcription complex subunit 1	5,90
TRINITY_DN11727_c0_g1_i6	Catechol O-methyltransferase domain-containing protein 1	6,31
TRINITY_DN13946_c0_g1_i1	Receptor expression-enhancing protein 3	6,33
TRINITY_DN14782_c1_g1_i20	RNA-directed DNA polymerase from mobile element jockey	6,45
TRINITY_DN9174_c0_g2_i1	Partitioning defective 6 homolog gamma	7,79
TRINITY_DN11070_c0_g1_i32	ATP-dependent RNA helicase DDX55	7,83
TRINITY_DN13965_c7_g1_i1	Endoribonuclease ZC3H12A	7,87
TRINITY_DN12078_c1_g3_i1	Lysine-specific demethylase 6B	7,89
TRINITY_DN13091_c1_g2_i9	Carboxypeptidase M	7,90
TRINITY_DN9989_c0_g1_i2	Laminin subunit beta-2	7,96
TRINITY_DN14133_c9_g1_i18	Carabin	7,96
TRINITY_DN14223_c0_g2_i4	Voltage-dependent anion-selective channel protein 2	7,90 7,97
TRINITY_DN10596_c0_g1_i1	Transcriptional regulator ATRX	8,02
TRINITY_DN11963_c1_g2_i8	Beta-crystallin A3	8,05
TRINITY_DN11353_c3_g1_i5	Annexin A6	8,05
TRINITY_DN14636_c0_g2_i16	Neoverrucotoxin subunit beta	8,06
TRINITY_DN12435_c5_g1_i2	pre-mRNA 3' end processing protein WDR33	8,07
TRINITY_DN12854_c0_g1_i9	Nucleoporin NUP188 homolog	8,14
TRINITY_DN13344_c0_g1_i9	GTPase IMAP family member 4	8,21
TRINITY_DN10009_c0_g1_i3	FAD synthase	8,24
TRINITY_DN12901_c3_g1_i7	Hydroxysteroid 11-beta-dehydrogenase 1-like protein	8,24
TRINITY_DN13541_c5_g2_i10	Canalicular multispecific organic anion transporter 2	8,25
TRINITY_DN14188_c1_g1_i13	Carnitine O-acetyltransferase	8,25
TRINITY_DN13051_c2_g1_i1	Myocardial zonula adherens protein	8,26
TRINITY_DN12720_c3_g18_i1	C-terminal-binding protein 2	8,29
TRINITY_DN12006_c2_g1_i16	Armadillo repeat-containing protein 1	8,31
TRINITY_DN12307_c3_g2_i13	Gamma-tubulin complex component 4	8,31
TRINITY_DN12711_c2_g3_i29	Copper-transporting ATPase 2	8,31
TRINITY_DN13426_c1_g1_i4	Hepatocyte growth factor-like protein	8,32
TRINITY_DN11084_c0_g1_i2	Ras-related protein Rab-40C	8,38
TRINITY_DN11222_c0_g2_i3	Retrovirus-related Pol polyprotein from transposon RE1	8,42
TRINITY_DN13485_c1_g6_i3	Cingulin-like protein 1	8,44
TRINITY_DN12874_c1_g2_i5	Transcriptional repressor CTCF	8,44
TRINITY_DN13485_c1_g6_i8	Cingulin-like protein 1	8,49
TRINITY_DN13394_c2_g13_i10	CCAAT/enhancer-binding protein gamma	8,51
TRINITY_DN12304_c0_g3_i11	Thrombospondin type-1 domain-containing protein 4	8,51
TRINITY_DN14304_c1_g6_i1	Desmin	8,53
TRINITY_DN11944_c4_g2_i3	Claudin domain-containing protein 1	8,57
TRINITY_DN13962_c0_g1_i9	KAT8 regulatory NSL complex subunit 1-like protein	8,57
TRINITY_DN13302_c0_g1_i9 TRINITY_DN13332_c1_g2_i8	Transcription factor SPT20 homolog	8,58
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TRINITY_DN11700_c3_g1_i13	Complement component C8 alpha chain	8,58
TRINITY_DN14227_c1_g2_i3	Nuclear receptor-binding protein	8,62
TRINITY_DN13748_c2_g1_i10	Transmembrane protein 269	8,63
TRINITY_DN8412_c0_g1_i2	Protein NRDE2 homolog	8,70
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TRINITY_DN13751_c2_g1_i28 TRINITY_DN13089_c1_g1_i14	RUN and FYVE domain-containing protein 2 Neutral and basic amino acid transport protein rBAT	8,71 8,73

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TRINITY_DN13968_c2_g1_i28	Leucine-rich PPR motif-containing protein, mitochondrial	8,80
TRINITY_DN11721_c1_g1_i13	3'(2'),5'-bisphosphate nucleotidase 1	8,85
TRINITY_DN14472_c2_g3_i7	Afadin- and alpha-actinin-binding protein	8,90
TRINITY_DN13819_c1_g1_i8	Arf-GAP with SH3 domain, ANK repeat and PH domain-contain	8,91
TRINITY_DN10456_c0_g1_i6	ATP-dependent RNA helicase DDX42	8,92
TRINITY_DN11917_c1_g1_i10	Vascular endothelial growth factor A	8,93
TRINITY_DN13846_c0_g1_i15	Serine/threonine-protein kinase LATS1	8,94
TRINITY_DN13941_c2_g5_i12	Carbohydrate-responsive element-binding protein	8,97
TRINITY_DN12476_c1_g4_i9	Cullin-9	9,04
TRINITY_DN14582_c1_g2_i14	Phenylalanine-4-hydroxylase	9,08
TRINITY_DN11529_c7_g1_i6	Ubiquitin-conjugating enzyme E2 Q2	9,10
TRINITY_DN9970_c0_g1_i3	Sestrin-1	9,15
TRINITY_DN13746_c3_g1_i7	Integrin beta-1	9,16
TRINITY_DN14310_c5_g6_i5	GTP-binding protein 2	9,17
TRINITY_DN13629_c2_g1_i39	Vitamin K-dependent protein C	9,19
TRINITY_DN13525_c7_g1_i6	Ubinuclein-2	9,25
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TRINITY_DN13969_c4_g1_i11	tRNA wybutosine-synthesizing protein 5	9,26
TRINITY_DN13148_c2_g1_i1	Sprouty-related, EVH1 domain-containing protein 2	9,26
TRINITY_DN13600_c3_g1_i5	Intersectin-1	9,29
TRINITY_DN14312_c0_g3_i11	Glutamate dehydrogenase 1, mitochondrial	9,31
TRINITY_DN13394_c2_g13_i3	CCAAT/enhancer-binding protein gamma	9,37
TRINITY_DN13394_c2_g7_i1	CCAAT/enhancer-binding protein gamma	9,37
TRINITY_DN12490_c1_g1_i2	BRISC complex subunit Abraxas 2	9,41
TRINITY_DN13968_c2_g1_i2	Leucine-rich PPR motif-containing protein, mitochondrial	9,41
TRINITY_DN12435_c5_g1_i4	pre-mRNA 3' end processing protein WDR33	9,42
TRINITY_DN9317_c0_g1_i2	ATP-dependent RNA helicase SUPV3L1, mitochondrial	9,42
TRINITY_DN12983_c5_g4_i7	Electron transfer flavoprotein beta subunit lysine methyltransfe	9,43
TRINITY_DN11730_c3_g1_i17	Phytanoyl-CoA dioxygenase, peroxisomal	9,48
TRINITY_DN10028_c0_g1_i5	Importin-4	9,50
TRINITY_DN13086_c0_g1_i8	Zinc finger protein 609	9,51
TRINITY_DN14877_c8_g1_i7	Ephrin type-B receptor 4	9,52
TRINITY_DN14299_c3_g3_i4	Protein turtle homolog A	9,55
TRINITY_DN11547_c3_g4_i8	Fibrinogen alpha chain	9,58
TRINITY_DN12304_c0_g3_i7	Thrombospondin type-1 domain-containing protein 4	9,69
TRINITY_DN13674_c4_g5_i4	Cholesterol 7-alpha-monooxygenase	9,74
TRINITY_DN11758_c2_g5_i2	Ketimine reductase mu-crystallin	9,76
TRINITY_DN12124_c1_g1_i29	Feline leukemia virus subgroup C receptor-related protein 2	9,89
TRINITY_DN11881_c3_g2_i9	Microprocessor complex subunit DGCR8	9,92
TRINITY_DN11727_c0_g1_i3	Catechol O-methyltransferase domain-containing protein 1	9,96
TRINITY_DN11769_c0_g2_i15	E3 ubiquitin-protein ligase RNF139	9,96
TRINITY_DN11529_c7_g1_i9	Ubiquitin-conjugating enzyme E2 Q2	9,98
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TRINITY_DN11986_c2_g4_i5	Transmembrane channel-like protein 6	9,99
TRINITY_DN14299_c3_g3_i2	Protein turtle homolog A	10,06
TRINITY_DN14550_c1_g2_i4	Heparan-alpha-glucosaminide N-acetyltransferase	10,06
TRINITY_DN12872_c8_g2_i4	CUGBP Elav-like family member 2	10,07
TRINITY_DN12711_c2_g3_i13	Copper-transporting ATPase 2	10,08
TRINITY_DN12521_c0_g2_i3	E3 ubiquitin-protein ligase NEDD4-like	10,13
TRINITY_DN12862_c4_g2_i3	von Willebrand factor (Fragment)	10,29
TRINITY_DN12673_c0_g1_i4	Phosphatidylcholine-sterol acyltransferase	10,29
TRINITY_DN12646_c3_g3_i1	Methylenetetrahydrofolate reductase	10,36
TRINITY_DN12417_c6_g1_i20	Apoptosis-inducing factor 1, mitochondrial	10,39
TRINITY_DN12185_c3_g1_i7	Putative deoxyribonuclease TATDN1	10,41
TRINITY_DN13751_c2_g1_i1	RUN and FYVE domain-containing protein 2	10,49
TRINITY_DN13485_c1_g6_i2	Cingulin-like protein 1	10,51
TRINITY_DN14312_c0_g3_i17	Glutamate dehydrogenase 1, mitochondrial	10,51
TRINITY_DN13504_c0_g2_i4	Kinesin-1 heavy chain	10,62
TRINITY_DN10842_c0_g1_i2	NHL repeat-containing protein 3	10,66
TRINITY_DN13600_c3_g1_i1	Intersectin-1	10,81
TRINITY_DN13433_c3_g6_i3	Protein FAM210B, mitochondrial	10,98
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TTD 11 11 12 12 12 12 12 12 12 12 12 12 12		44.45
TRINITY_DN13968_c2_g1_i3	Leucine-rich PPR motif-containing protein, mitochondrial	11,47
TRINITY_DN12784_c3_g1_i4	Heterogeneous nuclear ribonucleoprotein Q	11,53
TRINITY_DN13968_c2_g1_i8	Leucine-rich PPR motif-containing protein, mitochondrial	11,71
TRINITY_DN13051_c2_g1_i19	Myocardial zonula adherens protein	11,79
TRINITY_DN14418_c1_g1_i14	H-2 class I histocompatibility antigen, L-D alpha chain	13,23
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TRINITY_DN32828_c2_g1_i19	Glucagon receptor	-13,87
TRINITY_DN32734_c1_g2_i12	Myosin-9	-12,40
TRINITY_DN32192_c2_g2_i2	Fatty acid desaturase 2	-12,33
TRINITY_DN31514_c1_g2_i6	G patch domain-containing protein 8	-12,32
TRINITY_DN29887_c3_g1_i1	ERO1-like protein beta	-12,22
TRINITY_DN25606_c0_g2_i1	Major histocompatibility complex class I-related gene protein	-12,17
TRINITY_DN32835_c0_g2_i6	Fibronectin type-III domain-containing protein 3a	-12,16
TRINITY_DN28588_c6_g4_i2	Sialomucin core protein 24	-11,99
TRINITY_DN27228_c0_g1_i5	Diamine acetyltransferase 1	-11,76
TRINITY_DN25394_c0_g1_i31	Class I histocompatibility antigen, F10 alpha chain	-11,58
TRINITY_DN31079_c0_g1_i4	Stimulated by retinoic acid gene 6 protein-like	-11,51
TRINITY_DN28512_c0_g2_i13	Major histocompatibility complex class I-related gene protein	-11,48
TRINITY_DN32745_c0_g2_i7	Transferrin receptor protein 1	-11,44
TRINITY_DN26453_c0_g1_i10	Multidrug resistance-associated protein 5	
_	•	-11,38
TRINITY_DN29810_c0_g1_i5	GlutaminetRNA ligase	-11,38
TRINITY_DN30294_c1_g2_i3	Cathepsin F	-11,37
TRINITY_DN32477_c0_g1_i2	Myosin light chain kinase, smooth muscle (Fragment)	-11,35
TRINITY_DN29377_c2_g1_i2	Apolipoprotein B-100	-11,35
TRINITY_DN24543_c6_g2_i1	Lipocalin	-11,30
TRINITY_DN32756_c0_g1_i22	Phosphatidate phosphatase LPIN1	-11,29
TRINITY_DN33079_c1_g1_i22	Solute carrier family 41 member 1	-11,28
TRINITY_DN28410_c2_g1_i12	Immunoglobulin mu heavy chain	-11,22
TRINITY_DN31058_c0_g1_i9	Ezrin	-11,22
TRINITY_DN32403_c1_g1_i12	Dedicator of cytokinesis protein 8	-11,21
TRINITY_DN24918_c3_g1_i6	Ribonuclease-like 3	-11,19
TRINITY_DN28512_c0_g1_i4	Major histocompatibility complex class I-related gene protein	-11,19
TRINITY_DN31027_c2_g2_i4	Erlin-2	-11,15
TRINITY_DN33324_c15_g1_i2	LINE-1 retrotransposable element ORF2 protein	-11,13
TRINITY_DN32175_c0_g1_i10	Chloride intracellular channel protein 4	-11,12
TRINITY_DN32735_c0_g1_i8	Activating molecule in BECN1-regulated autophagy protein 1	-11,11
TRINITY_DN33029_c2_g1_i7	Dihydropyrimidine dehydrogenase [NADP(+)]	-11,11
TRINITY_DN33103_c2_g1_i11	Leukocyte receptor cluster member 8 homolog	-11,10
TRINITY_DN29580_c0_g1_i6	Ubiquitin carboxyl-terminal hydrolase 5	-11,10
TRINITY_DN33069_c3_g1_i12	Ribonucleoside-diphosphate reductase subunit M2 B	-10,97
TRINITY_DN29333_c2_g1_i2	Calcium-binding protein 39-like	-10,96
TRINITY_DN31061_c0_g1_i5	von Willebrand factor A domain-containing protein 8	-10,95
TRINITY_DN27385_c5_g2_i7	Uridine phosphorylase 2	-10,93
TRINITY_DN33024_c5_g3_i4	Cell death activator CIDE-3	-10,92
TRINITY_DN32371_c0_g1_i3	Eukaryotic translation initiation factor 2-alpha kinase 3	-10,92
TRINITY_DN32783_c3_g2_i2	Integrin beta-1	-10,92
TRINITY_DN32245_c0_g1_i10	Thioredoxin-interacting protein	-10,90
TRINITY_DN31579_c0_g2_i8	APOBEC1 complementation factor	-10,88
TRINITY_DN25606_c0_g6_i3	Class I histocompatibility antigen, F10 alpha chain	-10,88
TRINITY_DN30365_c1_g1_i2	DNA transposase THAP9	-10,83
TRINITY_DN30946_c0_g5_i1	Perforin-1	-10,82
TRINITY_DN29852_c3_g2_i1	Arylsulfatase B	-10,81
TRINITY_DN32693_c7_g1_i24	Interleukin-1 receptor type 2	-10,80
TRINITY_DN30314_c0_g2_i11	Heat shock factor protein 1	-10,79
	v -	-10,79
1KINI1 1 DN 52809 C1 21 11	A-kınase anchor protein 13	-10,/2
TRINITY_DN32809_c1_g1_i1 TRINITY_DN32503_c1_g1_i4	A-kinase anchor protein 13 Membrane-associated guanylate kinase, WW and PDZ domain-	
TRINITY_DN32809_C1_g1_i1 TRINITY_DN32503_c1_g1_i4 TRINITY_DN31202_c0_g1_i19	A-kinase anchor protein 13 Membrane-associated guanylate kinase, WW and PDZ domain- 1-phosphatidylinositol 3-phosphate 5-kinase	-10,79 -10,77 -10,76

TRINITY_DN32567_c0_g3_i8	Foulthand have nuctain O.1. A	-10,75
TRINITY_DN32298_c3_g1_i11	Forkhead box protein O1-A Unconventional myosin-IXb	-10,75
TRINITY_DN28036_c1_g3_i2	T-complex protein 1 subunit gamma	-10,75
TRINITY_DN33306_c3_g1_i6	Coiled-coil domain-containing protein 93	-10,75
TRINITY_DN32587_c1_g2_i4	Homeodomain-interacting protein kinase 2	-10,73
TRINITY_DN32529_c0_g1_i4	Monocarboxylate transporter 13	-10,73
TRINITY_DN31649_c0_g1_i9	Sorting nexin-14	-10,73
TRINITY_DN32378_c0_g1_i6	4-aminobutyrate aminotransferase, mitochondrial	-10,73
TRINITY_DN27660_c2_g1_i4	Protein phosphatase 1 regulatory subunit 14B	-10,70
TRINITY_DN33069_c3_g1_i9	Ribonucleoside-diphosphate reductase subunit M2 B	-10,70
TRINITY_DN27341_c0_g1_i2	F-box only protein 33	-10,68
TRINITY_DN32236_c0_g1_i3	Beta-glucuronidase	-10,68
TRINITY_DN32245_c0_g1_i12	Thioredoxin-interacting protein	-10,62
TRINITY_DN30049_c1_g1_i10	Cbp/p300-interacting transactivator 3	-10,62
TRINITY_DN31001_c0_g1_i1	Protein strawberry notch homolog 1	-10,62
TRINITY_DN32882_c1_g1_i11	Inositol 1,4,5-trisphosphate receptor type 1	-10,62
TRINITY_DN32844_c3_g1_i3	5'-nucleotidase domain-containing protein 2	-10,60
TRINITY_DN28782_c0_g1_i14	H-2 class II histocompatibility antigen, E-U alpha chain	-10,60
TRINITY_DN28355_c3_g2_i3	Dual specificity tyrosine-phosphorylation-regulated kinase 1A	-10,59
TRINITY_DN26699_c1_g2_i10	Exocyst complex component 6B	-10,59
TRINITY_DN32882_c1_g1_i9	Inositol 1,4,5-trisphosphate receptor type 1	-10,56
TRINITY_DN30270_c3_g1_i6	Lysocardiolipin acyltransferase 1	-10,56
TRINITY_DN29412_c0_g1_i1	KAT8 regulatory NSL complex subunit 3	-10,55
TRINITY_DN28574_c0_g1_i1	Myocilin	-10,54
TRINITY_DN26523_c0_g1_i11	Insulin receptor substrate 1-B	-10,52
TRINITY_DN26891_c0_g1_i6	Lipoprotein lipase	-10,52
TRINITY_DN26806_c0_g1_i3	E3 ubiquitin-protein ligase ARIH2	-10,51
TRINITY_DN31471_c1_g2_i14	Protein KIAA0100	-10,50
TRINITY_DN31027_c2_g2_i9	Erlin-2	-10,49
TRINITY_DN25391_c0_g1_i1	Glutamine synthetase	-10,48
TRINITY_DN32808_c2_g1_i12	Cell death activator CIDE-A	-10,47
TRINITY_DN32906_c2_g1_i8	Apoptotic chromatin condensation inducer in the nucleus	-10,47
TRINITY_DN31365_c0_g3_i5	Lysosomal acid phosphatase	-10,45
TRINITY_DN28797_c0_g1_i17	Serine/threonine-protein kinase ppk4	-10,43
TRINITY_DN31847_c1_g1_i12	Tyrosine-protein kinase JAK1	-10,41
TRINITY_DN25366_c2_g2_i20	5-hydroxyisourate hydrolase	-10,41
TRINITY_DN30393_c0_g1_i8	RNA helicase aquarius	-10,41
TRINITY_DN26279_c1_g1_i4	Putative hexokinase HKDC1	-10,41
TRINITY_DN33132_c1_g1_i10	Peptidyl-prolyl cis-trans isomerase FKBP5	-10,40
TRINITY_DN30940_c2_g1_i5	Elongin-B	-10,40
TRINITY_DN32707_c1_g1_i2	Coagulation factor VIII	-10,38
TRINITY_DN29887_c3_g1_i15	ERO1-like protein beta	-10,37
TRINITY_DN28498_c1_g1_i11	Kynureninase	-10,36
TRINITY_DN33282_c5_g1_i7	Regulator of microtubule dynamics protein 3	-10,33
TRINITY_DN30475_c0_g2_i8	Lysine-specific demethylase 6B	-10,33
TRINITY_DN32928_c2_g1_i12	Interleukin-6 receptor subunit alpha	-10,33
TRINITY_DN27817_c0_g1_i9	Heparin cofactor 2	-10,32
TRINITY_DN32759_c2_g1_i4	Hyaluronan-binding protein 2	-10,32
TRINITY_DN31472_c0_g1_i11	Autism susceptibility gene 2 protein	-10,32
TRINITY_DN26722_c2_g2_i2	Paired amphipathic helix protein Sin3a	-10,31
TRINITY_DN33122_c1_g2_i2	Autophagy-related protein 2 homolog A	-10,31
TRINITY_DN23495_c0_g1_i12	Fatty acid-binding protein, heart	-10,30
TRINITY_DN32372_c2_g1_i2	Protein Shroom1	-10,29
TRINITY_DN31085_c1_g1_i2	Selenoprotein O	-10,29
TRINITY_DN32685_c2_g1_i11	Erbin	-10,28
TRINITY_DN32356_c3_g1_i5	Striatin Many and alignorable and 1.2 alpha many acidese IC	-10,28
TRINITY_DN28570_c2_g1_i18	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IC	-10,28
TRINITY_DN26823_c0_g1_i5 TRINITY_DN32863_c0_g1_i12	Fatty acid desaturase 6 Nischarin	-10,27 -10,27
1Kiivii 1_Div32003_00_g1_i12	11toChail III	-10,47

TRINITY_DN30752_c0_g1_i3	Repressor of RNA polymerase III transcription MAF1 homolog	-10,24
TRINITY_DN32182_c0_g2_i5	Ras-related protein Rab-21	-10,24
TRINITY_DN31679_c0_g1_i4	Microfibrillar-associated protein 3-like	-10,24
TRINITY_DN27367_c1_g1_i4	Proprotein convertase subtilisin/kexin type 5	-10,24
TRINITY_DN27857_c0_g1_i7	Anaphase-promoting complex subunit 4	-10,24
TRINITY_DN32882_c1_g1_i28	Inositol 1,4,5-trisphosphate receptor type 1	-10,24
TRINITY_DN24650_c1_g1_i2	Nipped-B-like protein B	-10,22
TRINITY_DN32937_c1_g1_i11	Hypoxia-inducible factor 1-alpha	-10,22
TRINITY_DN30475_c0_g2_i11	Lysine-specific demethylase 6B	-10,20
TRINITY_DN24905_c0_g1_i1	Glycine amidinotransferase, mitochondrial	-10,20
TRINITY_DN32812_c2_g1_i1	Glycerol kinase	-10,20
	· ·	-10,20
TRINITY_DN28435_c1_g1_i2	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	
TRINITY_DN31838_c0_g1_i13	Focal adhesion kinase 1	-10,20
TRINITY_DN26852_c1_g1_i3	Zinc finger protein 286A	-10,19
TRINITY_DN32601_c0_g2_i6	Sarcosine dehydrogenase, mitochondrial	-10,19
TRINITY_DN32975_c0_g2_i2	E1A-binding protein p400	-10,17
TRINITY_DN28905_c0_g2_i2	Transcription factor Sox-9-B	-10,14
TRINITY_DN30541_c0_g1_i9	5'-AMP-activated protein kinase subunit beta-1	-10,14
TRINITY_DN31099_c0_g3_i2	Msx2-interacting protein	-10,13
TRINITY_DN28551_c1_g1_i4	Nuclear receptor coactivator 6	-10,13
TRINITY_DN29516_c2_g1_i17	Complement factor H	-10,13
TRINITY_DN33104_c6_g1_i31	Histidine ammonia-lyase	-10,13
TRINITY_DN26993_c2_g1_i15	MAU2 chromatid cohesion factor homolog	-10,13
TRINITY_DN25394_c0_g1_i8	HLA class I histocompatibility antigen, B-47 alpha chain	-10,13
TRINITY_DN33028_c2_g1_i6	Methylenetetrahydrofolate reductase	-10,12
TRINITY_DN30326_c0_g1_i5	Protein spinster homolog 1	-10,12
TRINITY_DN31950_c0_g1_i6	Tight junction protein ZO-1	-10,11
TRINITY_DN28925_c2_g2_i16		-10,11
	[Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme	
TRINITY_DN31936_c0_g2_i5	1-acyl-sn-glycerol-3-phosphate acyltransferase gamma	-10,10
TRINITY_DN28914_c0_g1_i5	Matrix metalloproteinase-14	-10,10
TRINITY_DN27398_c0_g1_i7	Pre-mRNA-splicing factor SLU7	-10,10
TRINITY_DN32665_c0_g1_i7	Immunoglobulin superfamily member 8	-10,08
TRINITY_DN31769_c2_g1_i5	GRAM domain-containing protein 2B	-10,08
TRINITY_DN27367_c1_g1_i14	Proprotein convertase subtilisin/kexin type 5	-10,08
TRINITY_DN30722_c1_g1_i11	Receptor-type tyrosine-protein phosphatase gamma	-10,07
TRINITY_DN30464_c0_g1_i6	Low affinity vacuolar monovalent cation/H(+) antiporter	-10,06
TRINITY_DN33078_c0_g1_i6	Serine/threonine-protein kinase MRCK beta	-10,06
TRINITY_DN26439_c0_g1_i4	Ankyrin repeat domain-containing protein 13A	-10,06
TRINITY_DN32200_c2_g1_i2	Forkhead box protein O4	-10,06
TRINITY_DN27875_c0_g1_i4	Hydroxyacylglutathione hydrolase, mitochondrial	-10,06
TRINITY_DN32228_c0_g1_i1	Translation initiation factor eIF-2B subunit epsilon	-10,04
TRINITY_DN32494_c0_g1_i11	Mitogen-activated protein kinase kinase 5	-10,03
TRINITY_DN32865_c1_g1_i21	RT1 class I histocompatibility antigen, AA alpha chain	-10,02
TRINITY_DN32224_c5_g1_i9	Mitochondrial glycine transporter A	-10,01
TRINITY_DN32587_c1_g2_i10	Homeodomain-interacting protein kinase 2	-9,98
TRINITY_DN31318_c3_g2_i9	Poly(ADP-ribose) glycohydrolase	-9,98
TRINITY_DN28637_c0_g1_i1	Solute carrier family 43 member 3	-9,97
TRINITY_DN29783_c1_g2_i4	Protein disulfide-isomerase A6	-9,96
TRINITY_DN30752_c1_g1_i10	Signal recognition particle subunit SRP68	-9,96
TRINITY_DN29927_c2_g1_i7	Protein FAM135A	-9,95
TRINITY_DN28692_c0_g1_i6	Patatin-like phospholipase domain-containing protein 2	-9,95
TRINITY_DN30147_c1_g1_i9	Amyloid-beta A4 protein	-9,95
TRINITY_DN31186_c0_g1_i6	Ubiquitin carboxyl-terminal hydrolase 36	-9,93
TRINITY_DN30511_c2_g1_i1	Calcium-binding mitochondrial carrier protein Aralar1	-9,93
TRINITY_DN27995_c0_g1_i16	Nuclear respiratory factor 1	-9,93
TRINITY_DN33000_c2_g1_i10	Clustered mitochondria protein homolog	-9,92
TRINITY_DN25394_c0_g1_i5	Class I histocompatibility antigen, F10 alpha chain	-9,92
TRINITY_DN31838_c0_g1_i18		
	Focal adhesion kinase 1	-9,91
TRINITY_DN24100_c2_g5_i9	Focal adhesion kinase 1 Vesicle-associated membrane protein 2	-9,91 -9,90

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TRINITY_DN23418_c0_g1_i1	COP9 signalosome complex subunit 5	-9,89
TRINITY_DN28309_c0_g1_i5	Intersectin-2	-9,89
TRINITY_DN25974_c0_g1_i7	Activating transcription factor 7-interacting protein 1	-9,87
TRINITY_DN31888_c2_g2_i2	Very-long-chain enoyl-CoA reductase	-9,87
TRINITY_DN31272_c0_g1_i18	Ubiquitin-protein ligase E3B	-9,86
TRINITY_DN25520_c0_g1_i3	E3 ubiquitin-protein ligase SMURF2	-9,85
TRINITY_DN25461_c3_g2_i11	Polyadenylate-binding protein-interacting protein 2B	-9,85
TRINITY_DN33321_c5_g1_i3	Microtubule-actin cross-linking factor 1	-9,85
TRINITY_DN29440_c0_g1_i1	Glutamine and serine-rich protein 1	-9,83
TRINITY_DN26016_c2_g1_i13	Carbohydrate sulfotransferase 10	-9,82
TRINITY_DN28306_c1_g1_i16	Flavin reductase (NADPH)	-9,82
TRINITY_DN32568_c2_g1_i14	SAFB-like transcription modulator	-9,82
TRINITY_DN32053_c2_g1_i1	Golgin subfamily A member 3	-9,81
TRINITY_DN32694_c1_g1_i3	Adenylate cyclase type 6	-9,81
TRINITY_DN31661_c0_g1_i1	High affinity cationic amino acid transporter 1	-9,80
TRINITY_DN31250_c0_g1_i11	D-dopachrome decarboxylase-A	-9,80
TRINITY_DN28724_c2_g1_i2	RNA 3'-terminal phosphate cyclase-like protein	-9,80
TRINITY_DN25606_c0_g5_i3	Major histocompatibility complex class I-related gene protein	-9,80
TRINITY_DN29051_c2_g5_i14	Ras-related protein Rab-9A	-9,78
TRINITY_DN29194_c0_g1_i13	Kalirin	-9,78
TRINITY_DN26560_c1_g1_i2	ATP-dependent RNA helicase DDX39A	-9,77
TRINITY_DN26312_c1_g4_i1	Serine/threonine-protein phosphatase PP1-alpha catalytic subu	-9,77
TRINITY_DN28143_c0_g1_i5		-9,77 -9,77
	Arginase-2, mitochondrial	
TRINITY_DN30490_c1_g3_i10	ATP-dependent Clp protease ATP-binding subunit clpX-like, mi	-9,75
TRINITY_DN30200_c3_g2_i6	Signal-induced proliferation-associated 1-like protein 2	-9,75
TRINITY_DN30321_c0_g1_i21	Nuclear transcription factor Y subunit gamma	-9,74
TRINITY_DN32671_c1_g1_i4	Zinc finger protein 706	-9,74
TRINITY_DN32298_c3_g1_i14	Unconventional myosin-IXb	-9,73
TRINITY_DN32080_c0_g1_i2	Histone-lysine N-methyltransferase SETD2	-9,71
TRINITY_DN22552_c0_g1_i3	Transcription factor BTF3 homolog 4	-9,71
TRINITY_DN26272_c0_g1_i1	Gamma-secretase subunit Aph-1b	-9,71
TRINITY_DN31748_c0_g1_i11	Acetyl-CoA acetyltransferase, mitochondrial	-9,70
TRINITY_DN24405_c0_g1_i2	Gelsolin	-9,70
TRINITY_DN30824_c0_g4_i1	Pyruvate kinase PKM	-9,70
TRINITY_DN28882_c0_g2_i7	Nuclear pore complex protein Nup155	-9,70
TRINITY_DN28233_c1_g1_i8	Glycogen synthase kinase-3 beta	-9,68
TRINITY_DN33299_c10_g1_i14	Tyrosine aminotransferase	-9,68
TRINITY_DN29800_c1_g4_i4	Aspartate aminotransferase, cytoplasmic	-9,68
TRINITY_DN28343_c1_g1_i1	Synaptotagmin-like protein 1	-9,68
TRINITY_DN30909_c0_g2_i4	DNA-binding protein RFX5	-9,67
TRINITY_DN31290_c1_g1_i4	CTD small phosphatase-like protein	-9,66
TRINITY_DN27188_c0_g1_i4	Cohesin subunit SA-2	-9,66
TRINITY_DN30984_c0_g1_i8	Ephexin-1	-9,66
TRINITY_DN32951_c2_g1_i2	Bone morphogenetic protein 1	-9,66
TRINITY_DN28263_c0_g1_i3	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	-9,66
TRINITY_DN29751_c0_g1_i5	Autophagy-related protein 16	-9,66
TRINITY_DN30748_c0_g1_i5	Zinc transporter ZIP9	-9,64
TRINITY_DN26436_c0_g1_i11	Histone H1.0	-9,63
TRINITY_DN30944_c2_g1_i7	Cysteine and histidine-rich domain-containing protein 1	-9,63
TRINITY_DN32440_c1_g2_i8	NAD kinase	-9,63
TRINITY_DN24384_c0_g1_i3		-9,63 -9,63
TRINITY_DN32195_c2_g2_i9	Serine/threonine-protein kinase 17A GTPase activating protein and VPSO domain containing protein	-9,63 -9,63
	GTPase-activating protein and VPS9 domain-containing protein	
TRINITY_DN28808_c0_g1_i1	Phosphatidate phosphatase LPIN2	-9,61
TRINITY_DN26538_c0_g1_i9	Synaptotagmin-like protein 4	-9,61
TRINITY_DN27007_c5_g1_i12	Limb region 1 homolog-like protein	-9,60
TRINITY_DN28924_c0_g1_i13	RNA-binding protein 20	-9,60
TRINITY_DN24124_c0_g1_i3	Endothelial lipase	-9,58
TRINITY_DN31057_c0_g1_i2	Endoribonuclease Dicer	-9,58
TRINITY_DN31365_c0_g3_i3	Lysosomal acid phosphatase	-9,58

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TRINITY_DN29898_c0_g2_i3	Pleckstrin homology domain-containing family G member 5	-9,58
TRINITY_DN30067_c1_g1_i4	Ubiquitin carboxyl-terminal hydrolase 42	-9,56
TRINITY_DN32216_c0_g1_i17	Ataxin-7	-9,56
TRINITY_DN28659_c3_g1_i6	SEC14-like protein 1	-9,56
TRINITY_DN28797_c0_g1_i21	Serine/threonine-protein kinase ppk4	-9,56
TRINITY_DN28483_c0_g1_i1	Ubiquitin conjugation factor E4 A	-9,55
TRINITY_DN25520_c0_g1_i1	E3 ubiquitin-protein ligase SMURF2	-9,55
TRINITY_DN26696_c1_g2_i5	SLIT-ROBO Rho GTPase-activating protein 3	-9,55
TRINITY_DN23800_c0_g1_i3	Collagen alpha-3(VI) chain	-9,55
TRINITY_DN30626_c1_g1_i11	CD9 antigen	-9,53
TRINITY_DN32486_c0_g1_i7	Serine/threonine-protein kinase Nek9	-9,53
TRINITY_DN27072_c0_g1_i2	Histone deacetylase 7	-9,52
TRINITY_DN32372_c2_g1_i11	Protein Shroom1	-9,52 -9,52
TRINITY_DN32978_c0_g1_i9	Protein ABHD16A	-9,52 -9,52
TRINITY_DN23495_c0_g1_i9	Fatty acid-binding protein, heart	-9,50
TRINITY_DN30569_c0_g1_i4	Leptin receptor gene-related protein	-9,50
TRINITY_DN26239_c0_g1_i7	Adipocyte enhancer-binding protein 1	-9,50
TRINITY_DN31549_c0_g1_i5	Golgin subfamily A member 4	-9,48
TRINITY_DN25715_c4_g1_i12	Importin subunit alpha-3	-9,48
TRINITY_DN24124_c0_g1_i5	Endothelial lipase	-9,47
TRINITY_DN29194_c0_g1_i14	Kalirin	-9,47
TRINITY_DN28933_c0_g1_i8	Sodium-independent sulfate anion transporter	-9,47
TRINITY_DN30759_c0_g2_i9	Malate synthase	-9,47
TRINITY_DN30748_c0_g1_i4	Zinc transporter ZIP9	-9,45
TRINITY_DN32641_c1_g2_i3	Glucose-6-phosphate exchanger SLC37A4	-9,45
TRINITY_DN30550_c0_g1_i10	Regulator of G-protein signaling 12	-9,45
TRINITY_DN33224_c10_g1_i6	Equilibrative nucleoside transporter 1	-9,45
TRINITY_DN29121_c0_g1_i6	Solute carrier family 35 member F6	-9,45
TRINITY_DN31914_c0_g1_i17	Meprin A subunit beta	-9,43
TRINITY_DN31272_c0_g1_i9	Ubiquitin-protein ligase E3B	-9,43
TRINITY_DN29809_c0_g1_i4	Pogo transposable element with ZNF domain	-9, 4 3
	Zinc finger FYVE domain-containing protein 1	-9,43 -9,43
TRINITY_DN32459_c2_g1_i10		
TRINITY_DN29855_c0_g2_i2	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyli	-9,43
TRINITY_DN27189_c1_g1_i4	Lysine-specific demethylase 5A	-9,43
TRINITY_DN30445_c1_g1_i8	Selenocysteine insertion sequence-binding protein 2	-9,42
TRINITY_DN31923_c0_g1_i5	AT-rich interactive domain-containing protein 1B	-9,42
TRINITY_DN32660_c0_g1_i9	Protein scribble homolog	-9,42
TRINITY_DN25126_c0_g1_i9	Exportin-4	-9,42
TRINITY_DN31316_c0_g1_i11	Tyrosine-protein kinase STYK1	-9,42
TRINITY_DN31778_c0_g1_i11	Patatin-like phospholipase domain-containing protein 7	-9,40
TRINITY_DN26288_c0_g1_i5	Chromatin target of PRMT1 protein	-9,40
TRINITY_DN29495_c2_g1_i8	Ubiquitin carboxyl-terminal hydrolase CYLD	-9,40
TRINITY_DN32053_c2_g1_i8	Golgin subfamily A member 3	-9,40
TRINITY_DN32155_c1_g2_i14	Putative helicase mov-10-B.1	-9,40
TRINITY_DN30038_c2_g2_i1	Transcription factor HIVEP2	-9,40
TRINITY_DN25606_c0_g3_i1	Class I histocompatibility antigen, F10 alpha chain	-9,40
TRINITY_DN33279_c1_g1_i2	RNA-binding protein 4	-9,40
TRINITY_DN29617_c3_g1_i5	Coagulation factor IX	-9,38
TRINITY_DN25394_c0_g1_i12	HLA class I histocompatibility antigen, B-52 alpha chain	-9,38
TRINITY_DN32452_c0_g1_i1	Ubiquitin carboxyl-terminal hydrolase 32	-9,38
TRINITY_DN21579_c0_g1_i2	Peptidyl-prolyl cis-trans isomerase FKBP8	-9,38 -9,38
TRINITY_DN28029_c1_g3_i2	Neurogenic locus notch homolog protein 2	-9,38
TRINITY_DN26658_c4_g1_i5	1-aminocyclopropane-1-carboxylate synthase-like protein 1	-9,38
TRINITY_DN30999_c0_g1_i1	Cullin-5	-9,38
TRINITY_DN31516_c0_g1_i5	Sister chromatid cohesion protein PDS5 homolog A	-9,36
TRINITY_DN28570_c2_g1_i21	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IC	-9,36
TRINITY_DN25038_c0_g1_i2	Growth factor receptor-bound protein 7	-9,36
TRINITY_DN22840_c0_g1_i2	Growth arrest and DNA damage-inducible protein GADD45 ga	-9,36
TRINITY_DN27613_c1_g2_i11	Calcium uptake protein 2, mitochondrial	-9,36

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TRINITY_DN29593_c1_g1_i4	Ubiquitin-conjugating enzyme E2 J2	-9,36
TRINITY_DN32855_c1_g1_i9	Ras-related protein Rab-23	-9,36
TRINITY_DN29884_c0_g1_i5	Inhibitor of nuclear factor kappa-B kinase subunit beta	-9,34
TRINITY_DN30722_c1_g1_i10	Receptor-type tyrosine-protein phosphatase gamma	-9,34
TRINITY_DN31623_c0_g1_i1	Retrovirus-related Pol polyprotein from type-1 retrotransposabi	-9,34
TRINITY_DN22732_c0_g1_i9	NPC1-like intracellular cholesterol transporter 1	-9,34
TRINITY_DN32850_c3_g1_i1	Arrestin domain-containing protein 3	-9,34
TRINITY_DN29966_c4_g1_i7	Mitochondrial glutamate carrier 1	-9,34
TRINITY_DN33042_c3_g1_i6	Guanylate kinase	-9,34
TRINITY_DN28375_c1_g1_i9	Trafficking kinesin-binding protein 1	-9,34
TRINITY_DN24890_c3_g1_i9	Beta-microseminoprotein	-9,34
TRINITY_DN32161_c1_g1_i1	Patched domain-containing protein 3	-9,34
TRINITY_DN32685_c2_g1_i10	Erbin	-9,34
TRINITY_DN31044_c1_g3_i1	Pyruvate dehydrogenase E1 component subunit beta, mitochonc	-9,34
TRINITY_DN33003_c1_g1_i5	Choline kinase alpha	-9,34 -9,34
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TRINITY_DN28808_c0_g1_i2	Phosphatidate phosphatase LPIN2	-9,33
TRINITY_DN26622_c0_g1_i1	Sorting nexin-29	-9,33
TRINITY_DN27015_c0_g3_i2	Membrane-associated phosphatidylinositol transfer protein 2	-9,33
TRINITY_DN32693_c7_g1_i26	Interleukin-1 receptor type 2	-9,33
TRINITY_DN29113_c0_g1_i3	Mitogen-activated protein kinase kinase kinase 10	-9,31
TRINITY_DN26205_c2_g2_i5	Palmitoyltransferase ZDHHC18	-9,31
TRINITY_DN25356_c0_g1_i4	Neurocan core protein	-9,31
TRINITY_DN28256_c0_g1_i5	Transposon Ty3-G Gag-Pol polyprotein	-9,31
TRINITY_DN29489_c0_g1_i13	Elongin-A	-9,29
TRINITY_DN22527_c0_g1_i2	Acidic fibroblast growth factor intracellular-binding protein B	-9,29
TRINITY_DN29714_c0_g1_i8	Serine/threonine-protein kinase Nek7	-9,29
TRINITY_DN29999_c0_g1_i3	RNA-binding protein 27	-9,29
TRINITY_DN27972_c1_g1_i7	Disintegrin and metalloproteinase domain-containing protein 1	-9,29
TRINITY_DN30108_c1_g1_i4	Septin-9	-9,29
TRINITY_DN30462_c0_g1_i20	Excitatory amino acid transporter 1	-9,27
TRINITY_DN32055_c1_g1_i4	Kinesin-like protein KIF21A	-9,27
TRINITY_DN31399_c0_g1_i4	Tyrosine-protein phosphatase non-receptor type 23	-9,27
TRINITY_DN24563_c0_g1_i5	Diphosphomevalonate decarboxylase	-9,27
TRINITY_DN32131_c4_g1_i3	Max dimerization protein 1	-9,27 -9,27
TRINITY_DN22433_c0_g1_i2	WD repeat domain phosphoinositide-interacting protein 4	-9,27 -9,25
TRINITY_DN27070_c0_g1_i19	Glutamine amidotransferase-like class 1 domain-containing pro	-9,25
TRINITY_DN32380_c0_g1_i4	Putative ATP-dependent RNA helicase an3	-9,25
TRINITY_DN32783_c3_g2_i3	Integrin beta-1	-9,25
TRINITY_DN30132_c0_g1_i7	L-fucose kinase	-9,25
TRINITY_DN32978_c0_g1_i14	ValinetRNA ligase	-9,23
TRINITY_DN28091_c1_g3_i10	Alpha-1,2-mannosyltransferase ALG9	-9,23
TRINITY_DN24465_c5_g1_i2	Cytochrome b561	-9,23
TRINITY_DN31810_c0_g1_i12	Protein tweety homolog 3	-9,23
TRINITY_DN26528_c1_g3_i2	Mitogen-activated protein kinase kinase kinase 3	-9,23
TRINITY_DN25727_c0_g2_i5	Inorganic pyrophosphatase	-9,23
TRINITY_DN28866_c0_g1_i6	Solute carrier family 41 member 1	-9,21
TRINITY_DN27323_c0_g2_i3	5-aminolevulinate synthase, erythroid-specific, mitochondrial	-9,21
TRINITY_DN25970_c3_g3_i17	Protein ABHD4	-9,21
TRINITY_DN32865_c1_g1_i8	Major histocompatibility complex class I-related gene protein	-9,21
TRINITY_DN23198_c0_g1_i4	Protein tyrosine phosphatase type IVA 2	-9,21
TRINITY_DN27975_c0_g1_i8	Ubiquitin thioesterase zranb1-B	-9,21
TRINITY_DN28574_c0_g1_i8	Myocilin	-9,21
TRINITY_DN28592_c0_g1_i1	Serine/threonine-protein phosphatase 6 regulatory ankyrin repe	-9,21 -9,21
TRINITY_DN29832_c3_g1_i4	Rho guanine nucleotide exchange factor 6	-9,21 -9,21
TRINITY_DN31735_c1_g2_i2		
_	T-complex protein 1 subunit epsilon	-9,19
TRINITY_DN27544_c0_g1_i14	Dynamin-2	-9,19
TRINITY_DN26192_c1_g1_i6	Transmembrane protein 26	-9,19
TRINITY_DN33088_c3_g1_i20	Inter-alpha-trypsin inhibitor heavy chain H3	-9,19
TRINITY_DN30348_c2_g1_i3	Keratinocyte-associated transmembrane protein 2	-9,17

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TRINITY_DN27266_c1_g1_i6	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransferas	-9,17
TRINITY_DN26488_c0_g1_i6	Rho guanine nucleotide exchange factor 18	-9,17
TRINITY_DN27557_c2_g4_i1	Serine/threonine-protein kinase TAO1	-9,17
TRINITY_DN27160_c12_g1_i2	Vitamin D-binding protein	-9,17
TRINITY_DN31884_c0_g1_i5	DNA repair protein REV1	-9,17
TRINITY_DN32666_c1_g1_i8	Plexin-B1	-9,17
TRINITY_DN25515_c1_g2_i7	Monocarboxylate transporter 2	-9,17
TRINITY_DN32449_c1_g1_i6	Activin receptor type-2A	-9,17
TRINITY_DN32302_c1_g1_i9	ATP-dependent RNA helicase DDX42	-9,17
TRINITY_DN29813_c0_g1_i11	Ribonuclease T2	-9,17
TRINITY_DN28540_c1_g1_i4	HCLS1-associated protein X-1	-9,17
TRINITY_DN31196_c0_g1_i2	Solute carrier family 12 member 4	-9,17
TRINITY_DN27514_c0_g2_i1	Pleckstrin homology domain-containing family O member 2	-9,17 -9,17
_		-9,17 -9,15
TRINITY_DN28651_c0_g2_i11	NLR family member X1	
TRINITY_DN29135_c1_g2_i8	WD repeat-containing protein 26	-9,15
TRINITY_DN24153_c0_g1_i3	Serine/threonine-protein kinase PAK 2	-9,15
TRINITY_DN28793_c0_g1_i3	SWI/SNF-related matrix-associated actin-dependent regulator ϵ	-9,15
TRINITY_DN27148_c1_g3_i12	Coiled-coil domain-containing protein 50	-9,15
TRINITY_DN31822_c0_g1_i20	N-lysine methyltransferase KMT5A	-9,13
TRINITY_DN33118_c2_g1_i3	Protein PRRC2A	-9,13
TRINITY_DN32863_c0_g1_i2	Nischarin	-9,13
TRINITY_DN32024_c1_g1_i12	Chromodomain-helicase-DNA-binding protein 1	-9,13
TRINITY_DN33001_c0_g1_i6	G protein-coupled receptor kinase 5	-9,13
TRINITY_DN24365_c0_g1_i2	Low-density lipoprotein receptor-related protein 6	-9,13
TRINITY_DN26877_c1_g1_i2	Catechol O-methyltransferase A	-9,13
TRINITY_DN29776_c0_g1_i6	Trinucleotide repeat-containing gene 6C protein	-9,13
TRINITY_DN27789_c2_g2_i2	Annexin A6	-9,13
TRINITY_DN29518_c0_g2_i2	Fragile X mental retardation syndrome-related protein 2	-9,13
TRINITY_DN26220_c0_g1_i4	Transcription factor SPT20 homolog	-9,13
TRINITY_DN28360_c1_g1_i3	Murinoglobulin-2	-9,13
TRINITY_DN31903_c0_g1_i19	Serine/threonine-protein kinase ULK3	-9,11
TRINITY_DN27034_c0_g1_i8	WD repeat-containing protein 48	-9,11
TRINITY_DN30600_c3_g1_i13	Insulin-like growth factor 2 mRNA-binding protein 3	-9,11
TRINITY_DN26119_c1_g2_i2	Protein SDA1 homolog	-9,11 -9,11
_	Tetratricopeptide repeat protein 36	
TRINITY_DN26415_c0_g1_i5		-9,11
TRINITY_DN31065_c3_g1_i3	Cytoskeleton-associated protein 4	-9,11
TRINITY_DN28480_c0_g1_i2	Signal recognition particle receptor subunit beta	-9,11
TRINITY_DN27293_c5_g1_i10	Phospholipase ABHD3	-9,11
TRINITY_DN30797_c1_g1_i2	Coenzyme Q-binding protein COQ10 homolog, mitochondrial	-9,09
TRINITY_DN25694_c1_g1_i4	Ubiquitin-conjugating enzyme E2 N	-9,09
TRINITY_DN29516_c2_g1_i13	Complement factor H-related protein 1	-9,09
TRINITY_DN33069_c3_g1_i5	Ribonucleoside-diphosphate reductase subunit M2 B	-9,09
TRINITY_DN28936_c0_g1_i4	Phosphatidate cytidylyltransferase 2	-9,09
TRINITY_DN28923_c0_g1_i3	TATA-binding protein-associated factor 172	-9,09
TRINITY_DN31065_c3_g1_i13	Cytoskeleton-associated protein 4	-9,09
TRINITY_DN21807_c0_g1_i8	Dihydroorotate dehydrogenase (quinone), mitochondrial	-9,09
TRINITY_DN32106_c0_g2_i8	Epidermal growth factor receptor kinase substrate 8	-9,09
TRINITY_DN28066_c0_g1_i1	Mothers against decapentaplegic homolog 9	-9,09
TRINITY_DN26970_c1_g1_i5	Phosphorylase b kinase gamma catalytic chain, skeletal muscle	-9,09
TRINITY_DN32072_c0_g2_i3	Cyclic AMP receptor-like protein A	-9,09
TRINITY_DN24769_c1_g1_i6	Forkhead box protein N4	-9,06
TRINITY_DN32660_c0_g1_i30	Protein scribble homolog	-9,06
TRINITY_DN30783_c0_g1_i4	Probable helicase senataxin	-9,06 -9,06
_		-9,06 -9,06
TRINITY_DN26517_c1_g2_i1	Interferon alpha/beta receptor 2 Powhesia receptor activated protein Coorf80 hamalas	
TRINITY_DN29397_c1_g1_i3	Bombesin receptor-activated protein C6orf89 homolog	-9,06
TRINITY_DN31193_c0_g1_i27	Guanine nucleotide-binding protein G(i) subunit alpha-2	-9,06
TRINITY_DN30680_c4_g1_i7	Interferon-induced very large GTPase 1	-9,06
TRINITY_DN26252_c1_g1_i13	Focal adhesion kinase 1	-9,06
TRINITY_DN32948_c1_g1_i1	Mitogen-activated protein kinase kinase kinase 7	-9,06

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TRINITY_DN31166_c0_g1_i7	Protein polybromo-1	-9,06
TRINITY_DN30711_c0_g1_i8	Ubiquitin carboxyl-terminal hydrolase 12	-9,06
TRINITY_DN26431_c0_g1_i7	Putative adenosylhomocysteinase 3	-9,06
TRINITY_DN30831_c2_g1_i15	Substance-P receptor	-9,06
TRINITY_DN21748_c0_g2_i3	Ictacalcin	-9,04
TRINITY_DN31035_c0_g1_i2	E3 ubiquitin-protein ligase UBR5	-9,04
TRINITY_DN29680_c0_g1_i2	Ornithine decarboxylase antizyme 2	-9,04
TRINITY_DN29633_c0_g1_i11	Ribonuclease 3	-9,04
TRINITY_DN30273_c0_g1_i9	Rab11 family-interacting protein 4B	-9,04
TRINITY_DN31679_c0_g1_i1	Microfibrillar-associated protein 3-like	-9,04
TRINITY_DN30467_c0_g1_i7	Cysteine-rich with EGF-like domain protein 2	-9,04
TRINITY_DN27700_c2_g2_i1	Phosphatidylinositol 5-phosphate 4-kinase type-2 alpha	-9,04
TRINITY_DN32193_c1_g2_i12	Sodium-coupled neutral amino acid transporter 2	-9,04
TRINITY_DN32809_c1_g1_i6	A-kinase anchor protein 13	-9,04
TRINITY_DN32978_c0_g1_i17	Protein ABHD16A	-9,04
TRINITY_DN32060_c4_g1_i6	Multidrug and toxin extrusion protein 1	-9,04
TRINITY_DN25853_c0_g1_i4	Ubiquitin thioesterase otulin	-9,02
TRINITY_DN32093_c3_g2_i21	Cytohesin-2	-9,02
TRINITY_DN30069_c0_g3_i19	Glycogen [starch] synthase, muscle	-9,02
TRINITY_DN22968_c0_g1_i2	Uncharacterized protein C1orf21 homolog	-9,02 -9,02
TRINITY_DN29171_c2_g1_i9	U2 small nuclear ribonucleoprotein A'	-9,02
TRINITY_DN32920_c2_g1_i10	Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondria	-9,02
TRINITY_DN25913_c6_g1_i1	Transposable element Tcb1 transposase	-9,02
TRINITY_DN30357_c0_g1_i4	Chromodomain-helicase-DNA-binding protein 4	-9,02
TRINITY_DN23935_c4_g1_i2	GTPase IMAP family member 5	-9,02
TRINITY_DN29341_c1_g1_i13	RNA-binding Raly-like protein	-9,02
TRINITY_DN29456_c0_g2_i1	Beta-secretase 2	-9,00
TRINITY_DN30610_c0_g1_i2	Mitochondrial fission regulator 1	-9,00
TRINITY_DN30626_c1_g1_i12	CD9 antigen	-9,00
TRINITY_DN28942_c2_g1_i1	Signal peptide peptidase-like 3	-9,00
TRINITY_DN27039_c0_g1_i3	Protein FAM234A	-9,00
TRINITY_DN32020_c4_g1_i12	Junction-mediating and -regulatory protein	-9,00
TRINITY_DN31493_c2_g1_i5	Tensin-1	-9,00
TRINITY_DN30012_c0_g1_i15	Probable RNA-directed DNA polymerase from transposon X-ele	-9,00
TRINITY_DN31256_c1_g2_i20	E3 ubiquitin-protein ligase HECTD1	-9,00
TRINITY_DN32633_c0_g1_i14	A disintegrin and metalloproteinase with thrombospondin motif.	-9,00
TRINITY_DN30862_c0_g1_i18	ATP-dependent RNA helicase DDX19A	-9,00
TRINITY_DN27403_c0_g1_i11	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial	-8,97
TRINITY_DN22214_c0_g1_i3	Male-specific lethal 3 homolog	-8,97
TRINITY_DN27640_c0_g1_i9	Small VCP/p97-interacting protein	-8,97
TRINITY_DN32691_c2_g1_i5	Palmitoyltransferase ZDHHC5	-8,97
TRINITY_DN28236_c0_g1_i4	Arrestin domain-containing protein 2	-8,97
TRINITY_DN26570_c0_g2_i9	Plastin-3	-8,97
TRINITY_DN32383_c0_g1_i8	ATP-dependent DNA helicase Q5	-8,97
TRINITY_DN28450_c1_g1_i3	Choline transporter-like protein 1	-8,97
TRINITY_DN32511_c1_g1_i4	Protein transport protein Sec31A	-8,97
TRINITY_DN28187_c0_g2_i9	PHD finger protein 13	-8,97
TRINITY_DN32069_c0_g1_i6	* * *	-8,97
_	Heterogeneous nuclear ribonucleoprotein D-like	
TRINITY_DN32627_c0_g1_i3	Inhibitor of Bruton tyrosine kinase	-8,97
TRINITY_DN29734_c0_g1_i5	Mucin-5AC	-8,97
TRINITY_DN26717_c0_g2_i8	High mobility group protein B2	-8,97
TRINITY_DN31255_c2_g1_i1	b(0,+)-type amino acid transporter 1	-8,97
TRINITY_DN24860_c0_g1_i12	Pyruvate dehydrogenase E1 component subunit alpha, somatic	-8,97
TRINITY_DN24125_c0_g1_i5	Annexin A1	-8,95
TRINITY_DN31374_c0_g2_i6	2-oxoglutarate dehydrogenase, mitochondrial	-8,95
TRINITY_DN29113_c0_g1_i7	Mitogen-activated protein kinase kinase kinase 10	-8,95
TRINITY_DN26830_c2_g2_i7	Tetratricopeptide repeat protein 17	-8,95
TRINITY_DN28032_c1_g1_i2	Retrovirus-related Pol polyprotein from transposon 297	-8,95
TRINITY_DN29377_c2_g1_i5	Apolipoprotein B-100	-8,95

TRINITY DN22261 a4 a1 :5	HEAT non-cat containing mastein 54	9.05
TRINITY_DN33261_c4_g1_i5	HEAT repeat-containing protein 5A	-8,95
TRINITY_DN27322_c2_g1_i2	Secernin-3	-8,95
TRINITY_DN29940_c2_g1_i3	Specifically androgen-regulated gene protein	-8,95
TRINITY_DN28187_c0_g2_i6	PHD finger protein 13	-8,95
TRINITY_DN26848_c0_g1_i9	Peroxisome proliferator-activated receptor delta	-8,92
TRINITY_DN29790_c2_g1_i1	Ubiquitin-like protein 3	-8,92
TRINITY_DN29922_c2_g1_i24	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	-8,92
TRINITY_DN32459_c2_g1_i15	Zinc finger FYVE domain-containing protein 1	-8,92
TRINITY_DN29302_c1_g2_i7	FGGY carbohydrate kinase domain-containing protein	-8,92
TRINITY_DN32582_c0_g1_i15	RING finger protein 10	-8,92
TRINITY_DN26302_c0_g1_i1	UDP-N-acetylglucosamine transporter	-8,92
TRINITY_DN33003_c1_g1_i18	Choline kinase alpha	-8,92
TRINITY_DN23195_c0_g1_i2	Phosphoinositide-3-kinase-interacting protein 1	-8,90
TRINITY_DN27429_c2_g1_i6	Cytosolic non-specific dipeptidase	-8,90
TRINITY_DN29009_c1_g2_i3	Serine/threonine-protein kinase DCLK3	-8,90
TRINITY_DN26063_c1_g1_i2	SH2B adapter protein 3	-8,90
TRINITY_DN28052_c2_g1_i8	Glucoside xylosyltransferase 1	-8,90
TRINITY_DN30241_c0_g1_i9	Nucleolar transcription factor 1	-8,90
TRINITY_DN31085_c1_g1_i5	Selenoprotein O	-8,90
TRINITY_DN29553_c0_g1_i11	Interferon-induced GTP-binding protein Mx	-8,90
TRINITY_DN26547_c1_g3_i1	Cell cycle control protein 50A	-8,90
TRINITY_DN27382_c0_g1_i3	Protein NEDD1	-8,90
TRINITY_DN24822_c2_g1_i11	Vitelline membrane outer layer protein 1 homolog	-8,90
TRINITY_DN27501_c2_g1_i4	Polyadenylate-binding protein 2	-8,90
TRINITY_DN32708_c2_g1_i3	Period circadian protein homolog 2	-8,90
TRINITY_DN32920_c1_g1_i3	Nuclear receptor coactivator 6	-8,90
TRINITY_DN28386_c1_g1_i5	Mitogen-activated protein kinase kinase kinase 10	-8,87
TRINITY_DN29575_c2_g1_i8	Rap guanine nucleotide exchange factor 6	-8,87
TRINITY_DN27685_c0_g1_i14	Prolyl endopeptidase-like	-8,87
TRINITY_DN30517_c0_g1_i6	Inhibitor of nuclear factor kappa-B kinase subunit alpha	-8,87
TRINITY_DN25778_c0_g4_i4	Inositol-pentakisphosphate 2-kinase	-8,87
TRINITY_DN30983_c0_g1_i4	Interleukin-17 receptor C	-8,87 -8,87
		-8,87 -8,87
TRINITY_DN29936_c0_g3_i2	Muscleblind-like protein 3	
TRINITY_DN29884_c0_g1_i9	Inhibitor of nuclear factor kappa-B kinase subunit beta	-8,87
TRINITY_DN26681_c0_g1_i6	Carbohydrate sulfotransferase 15	-8,87
TRINITY_DN27936_c0_g1_i16	Protein phosphatase 1 regulatory subunit 1B	-8,87
TRINITY_DN31776_c1_g1_i4	Ral GTPase-activating protein subunit beta	-8,87
TRINITY_DN32309_c1_g1_i5	Serum paraoxonase/arylesterase 2	-8,87
TRINITY_DN30067_c1_g1_i1	Ubiquitin carboxyl-terminal hydrolase 42	-8,87
TRINITY_DN29018_c0_g2_i3	Cytosolic purine 5'-nucleotidase	-8,87
TRINITY_DN32861_c1_g1_i2	E3 ubiquitin-protein ligase MARCH6	-8,87
TRINITY_DN30419_c0_g1_i14	Urotensin-2 receptor	-8,87
TRINITY_DN33003_c1_g1_i4	Choline kinase alpha	-8,87
TRINITY_DN29377_c1_g1_i6	KN motif and ankyrin repeat domain-containing protein 1	-8,85
TRINITY_DN29314_c1_g1_i4	Transcription elongation regulator 1	-8,85
TRINITY_DN31230_c0_g1_i2	Acyl-coenzyme A thioesterase 4	-8,85
TRINITY_DN26786_c0_g3_i1	Serine/threonine-protein phosphatase 6 regulatory subunit 3	-8,85
TRINITY_DN32128_c1_g1_i1	Peptidyl-glycine alpha-amidating monooxygenase A	-8,85
TRINITY_DN24860_c0_g1_i11	Pyruvate dehydrogenase E1 component subunit alpha, somatic	-8,85
TRINITY_DN30441_c1_g1_i7	V-set domain-containing T-cell activation inhibitor 1	-8,85
TRINITY_DN32068_c1_g2_i3	Nuclear receptor subfamily 1 group I member 2	-8,85
TRINITY_DN26288_c0_g1_i11	Chromatin target of PRMT1 protein	-8,85
TRINITY_DN25967_c3_g4_i1	LINE-1 reverse transcriptase homolog	-8,85
TRINITY_DN31105_c0_g1_i6	Transcription factor GATA-6	-8,85
TRINITY_DN28753_c0_g1_i9	E3 ubiquitin-protein ligase RNF34	-8,85
TRINITY_DN31398_c0_g1_i2	Phospholipase D2	-8,85
TRINITY_DN27654_c1_g2_i4	Mitochondrial carnitine/acylcarnitine carrier protein	-8,85
TRINITY_DN29503_c2_g1_i5	General transcription and DNA repair factor IIH helicase subu	-8,85
TRINITY_DN26959_c1_g1_i1	NADPHcytochrome P450 reductase	-8,82

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TRINITY_DN32705_c2_g1_i6	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase epsi	-8,82
TRINITY_DN25394_c0_g1_i37	Class I histocompatibility antigen, F10 alpha chain	-8,82
TRINITY_DN25126_c0_g1_i12	Exportin-4	-8,82
TRINITY_DN32965_c0_g1_i10	N-lysine methyltransferase setd6	-8,82
TRINITY_DN25684_c0_g2_i7	Protein Jumonji	-8,82
TRINITY_DN26445_c2_g1_i22	Uncharacterized protein KIAA0930	-8,82
TRINITY_DN27007_c5_g1_i7	Limb region 1 homolog-like protein	-8,82
TRINITY_DN25731_c4_g1_i15	Phosphatidylinositol 3,4,5-trisphosphate 3-phosphatase and du	-8,82
TRINITY_DN25126_c0_g1_i6	Exportin-4	-8,82
TRINITY_DN27891_c2_g1_i3	26S proteasome non-ATPase regulatory subunit 2	-8,82
TRINITY_DN29427_c1_g1_i11	Serine/arginine-rich splicing factor 2	-8,82
TRINITY_DN31480_c7_g1_i3	Anoctamin-1	-8,82
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TRINITY_DN25480_c0_g1_i8	Synaptic vesicle glycoprotein 2B	-8,82
TRINITY_DN29993_c0_g1_i2	Guanine nucleotide-binding protein-like 1	-8,82
TRINITY_DN24421_c2_g3_i11	40S ribosomal protein S25	-8,80
TRINITY_DN27314_c1_g1_i6	Splicing factor 1	-8,80
TRINITY_DN28357_c2_g3_i1	Zinc finger protein 711	-8,80
TRINITY_DN23995_c0_g1_i3	Beta-catenin-interacting protein 1	-8,80
TRINITY_DN22563_c0_g1_i4	Folate receptor alpha	-8,80
TRINITY_DN28600_c0_g1_i3	Hepatocyte nuclear factor 1-beta-A	-8,80
TRINITY_DN22878_c0_g1_i5	PDZ and LIM domain protein 4	-8,80
TRINITY_DN32408_c0_g1_i4	Protein strawberry notch homolog 2	-8,80
TRINITY_DN25654_c6_g1_i9	Ceruloplasmin	-8,80
TRINITY_DN27840_c0_g1_i16	Cell division cycle protein 27 homolog	-8,80
TRINITY_DN28242_c1_g1_i17		-8,80
	Inactive tyrosine-protein kinase PEAK1	
TRINITY_DN28864_c2_g1_i2	CCR4-NOT transcription complex subunit 1	-8,80
TRINITY_DN26059_c0_g1_i2	Sodium/potassium-transporting ATPase subunit beta-233	-8,80
TRINITY_DN30447_c1_g1_i1	60S ribosomal protein L27a	-8,80
TRINITY_DN30073_c1_g1_i10	Adenylate cyclase type 9	-8,80
TRINITY_DN29768_c2_g1_i3	Uncharacterized protein C18orf19 homolog A	-8,80
TRINITY_DN29503_c2_g3_i3	U3 small nucleolar RNA-associated protein 14 homolog A	-8,80
TRINITY_DN32402_c2_g1_i1	E3 ubiquitin-protein ligase HUWE1	-8,77
TRINITY_DN28809_c0_g2_i11	Ras-related protein Rap-2b	-8,77
TRINITY_DN31231_c0_g1_i10	DNA-binding death effector domain-containing protein 2	-8,77
TRINITY_DN30220_c3_g1_i7	KH homology domain-containing protein 4	-8,77
TRINITY_DN28341_c0_g1_i7	SEC23-interacting protein	-8,77
TRINITY_DN29921_c0_g1_i8	Partitioning defective 3 homolog	-8,77
TRINITY_DN29762_c0_g1_i2	E3 ubiquitin-protein ligase RNF38	-8,77
TRINITY_DN32138_c0_g1_i5	Signal transducing adapter molecule 2	-8,77
TRINITY_DN29932_c1_g2_i2	Serine/threonine-protein kinase Sgk1	-8,77
TRINITY_DN31636_c1_g1_i5	Upstream-binding protein 1	-8,77
TRINITY_DN28745_c0_g1_i4	Centrosomal protein of 85 kDa	-8,77
TRINITY_DN30958_c0_g1_i5		-8,77
	Cytochrome c-type heme lyase	
TRINITY_DN26453_c0_g1_i13	Multidrug resistance-associated protein 5	-8,77
TRINITY_DN23772_c0_g1_i1	Carbohydrate sulfotransferase 14	-8,77
TRINITY_DN32056_c0_g1_i8	Interferon-induced very large GTPase 1	-8,77
TRINITY_DN32824_c0_g1_i1	Regulator of nonsense transcripts 1	-8,74
TRINITY_DN31694_c0_g3_i1	Histone acetyltransferase p300	-8,74
TRINITY_DN32159_c0_g1_i15	CSC1-like protein 2	-8,74
TRINITY_DN28542_c1_g1_i3	E3 ubiquitin-protein ligase rnf146	-8,74
TRINITY_DN26279_c1_g1_i2	Putative hexokinase HKDC1	-8,74
TRINITY_DN30429_c0_g1_i18	Coiled-coil domain-containing protein 77	-8,74
TRINITY_DN29098_c1_g1_i4	WD repeat-containing protein 37	-8,74
TRINITY_DN28883_c3_g2_i3	Insulin-like growth factor I, adult form	-8,74
TRINITY_DN30247_c3_g1_i4	Tyrosine 3-monooxygenase	-8,74
TRINITY_DN24822_c2_g1_i19	Vitelline membrane outer layer protein 1 homolog	-8,74
TRINITY_DN25385_c3_g1_i1	Alanine aminotransferase 2, mitochondrial	-8,74 -8,74
	Protein MB21D2	-8,74 -8,74
TRINITY_DN28315_c0_g1_i14		
TRINITY_DN29921_c0_g1_i6	Partitioning defective 3 homolog	-8,74

TRINITY DN21101 .0 .1 .7	7' C' DATA I' I'	0.74
TRINITY_DN31181_c0_g1_i6	Zinc finger RNA-binding protein	-8,74
TRINITY_DN28903_c0_g1_i15	Eukaryotic translation initiation factor 3 subunit K	-8,74
TRINITY_DN24804_c0_g1_i1	DNA-directed RNA polymerase I subunit RPA2	-8,74
TRINITY_DN28433_c0_g1_i5	Transcriptional protein SWT1	-8,74
TRINITY_DN28144_c0_g1_i5	Sodium/potassium/calcium exchanger 3	-8,74
TRINITY_DN23695_c0_g1_i1	TNF receptor-associated factor 2	-8,74
TRINITY_DN27654_c1_g2_i14	Mitochondrial carnitine/acylcarnitine carrier protein	-8,74
TRINITY_DN29338_c2_g1_i1	Small G protein signaling modulator 1	-8,74
TRINITY_DN33098_c1_g2_i8	Probable fructose-2,6-bisphosphatase TIGAR A	-8,71
TRINITY_DN25632_c0_g1_i12	Growth arrest and DNA damage-inducible protein GADD45 be	-8,71
TRINITY_DN29768_c2_g1_i1	Uncharacterized protein C18orf19 homolog A	-8,71
TRINITY_DN31535_c2_g1_i10	Signal transducer and activator of transcription 5B	-8,71
TRINITY_DN28274_c0_g1_i3	NAD-dependent protein deacetylase sirtuin-3, mitochondrial	-8,71
TRINITY_DN29976_c1_g1_i18	Glycylpeptide N-tetradecanoyltransferase 2	-8,71
TRINITY_DN29643_c1_g2_i2	Arylsulfatase D	-8,71
TRINITY_DN26567_c0_g1_i6	Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX10	-8,71
TRINITY_DN32660_c0_g1_i18	Protein scribble homolog	-8,71
TRINITY_DN25366_c2_g2_i23	5-hydroxyisourate hydrolase	-8,71
TRINITY_DN32690_c1_g1_i12	SH3 and PX domain-containing protein 2B	-8,71
TRINITY_DN24842_c0_g1_i9	Vacuolar protein sorting-associated protein 28 homolog	-8,71
TRINITY_DN32966_c3_g1_i6	FERM, ARHGEF and pleckstrin domain-containing protein 1	-8,71
TRINITY_DN31527_c0_g1_i10	Rho GTPase-activating protein SYDE2	-8,71
TRINITY_DN30365_c1_g1_i8	DNA transposase THAP9	-8,71
TRINITY_DN26365_c5_g1_i5	Abl interactor 2	-8,69
TRINITY_DN30928_c0_g1_i13	Glycogen phosphorylase, liver form	-8,69
TRINITY_DN32053_c2_g1_i9	Golgin subfamily A member 3	-8,69
TRINITY_DN23157_c0_g1_i5	Contactin-associated protein like 5-2	-8,69
TRINITY_DN30998_c0_g1_i10	Putative homeodomain transcription factor 2	-8,69
TRINITY_DN25632_c0_g1_i3	Growth arrest and DNA damage-inducible protein GADD45 be	-8,69
TRINITY_DN24929_c0_g1_i5	Bone morphogenetic protein 1	-8,69
TRINITY_DN30554_c1_g1_i3	CLOCK-interacting pacemaker	-8,69
TRINITY_DN30927_c1_g3_i2	Protein LYRIC	-8,69
TRINITY_DN33116_c2_g1_i4	Helicase with zinc finger domain 2	-8,69
TRINITY_DN25771_c0_g1_i12	Protein Smaug homolog 1	-8,69
TRINITY_DN23660_c2_g2_i2	F-actin-capping protein subunit alpha-1	-8,69
TRINITY_DN30789_c0_g1_i5	Presenilins-associated rhomboid-like protein, mitochondrial	-8,69
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TRINITY_DN32426_c1_g1_i18	Protein transport protein Sec24C	-8,66
TRINITY_DN30655_c2_g1_i25	Pseudouridylate synthase 7 homolog	-8,66
TRINITY_DN30856_c3_g1_i3	Electron transfer flavoprotein subunit alpha, mitochondrial	-8,66
TRINITY_DN32300_c1_g1_i3	Nuclear pore complex protein Nup153	-8,66
TRINITY_DN31134_c0_g1_i9	Drebrin	-8,66
TRINITY_DN29611_c0_g1_i3	Ras-related protein Rab-24	-8,66
TRINITY_DN30655_c2_g1_i23	Pseudouridylate synthase 7 homolog	-8,66
TRINITY_DN25616_c0_g1_i3	Ependymin	-8,66
TRINITY_DN32395_c1_g2_i8	Biogenesis of lysosome-related organelles complex 1 subunit 6	-8,66
TRINITY_DN20717_c0_g1_i1	Proteasome subunit beta type-8	-8,66
TRINITY_DN30544_c0_g1_i15	Junctional adhesion molecule A	-8,66
TRINITY_DN27548_c0_g1_i7	Transcription initiation factor TFIID subunit 10	-8,66
TRINITY_DN31299_c0_g1_i5	Deoxynucleoside triphosphate triphosphohydrolase SAMHD1	-8,66
TRINITY_DN23637_c0_g1_i2	ATPase family AAA domain-containing protein 2	-8,66
TRINITY_DN32525_c0_g1_i4	Transcriptional regulator ATRX	-8,66
TRINITY_DN30318_c0_g2_i13	Pregnancy zone protein	-8,66
TRINITY_DN24224_c0_g2_i9	PAX-interacting protein 1	-8,66
TRINITY_DN25843_c1_g3_i10	Cell division cycle-associated protein 4	-8,66
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TRINITY_DN31448_c1_g1_i13	Mitogen-activated protein kinase kinase kinase kinase 5	-8,66 8,66
TRINITY_DN31195_c0_g1_i11	Spectrin alpha chain, non-erythrocytic 1	-8,66
TRINITY_DN22380_c0_g1_i4	Sulfate anion transporter 1	-8,63
TRINITY_DN31604_c0_g1_i2	Insulin-degrading enzyme	-8,63
TRINITY_DN24553_c0_g1_i14	Polyhomeotic-like protein 2	-8,63

TRINITY DN22029 22 21 :7	Mathulan atatuahu duafalata na duataga	0 62
TRINITY_DN33028_c2_g1_i7	Methylenetetrahydrofolate reductase	-8,63
TRINITY_DN29332_c0_g1_i14	Chemokine-like receptor 1	-8,63
TRINITY_DN25384_c0_g1_i1	Synembryn-B	-8,63
TRINITY_DN28375_c1_g1_i1	Trafficking kinesin-binding protein 1	-8,63
TRINITY_DN25663_c0_g4_i5	Ran guanine nucleotide release factor	-8,63
TRINITY_DN27917_c0_g1_i2	Ubiquitin-fold modifier 1	-8,63
TRINITY_DN31804_c1_g1_i3	TBC1 domain family member 5	-8,63
TRINITY_DN28341_c0_g1_i10	SEC23-interacting protein	-8,63
TRINITY_DN26133_c1_g2_i4	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subun	-8,63
TRINITY_DN28419_c1_g2_i3	AN1-type zinc finger protein 5	-8,63
TRINITY_DN25085_c0_g1_i1	Transposon Tf2-9 polyprotein	-8,63
TRINITY_DN25008_c0_g1_i9	Cytoplasmic polyadenylation element-binding protein 4	-8,63
TRINITY_DN32863_c0_g1_i7	Nischarin	-8,63
TRINITY_DN29764_c1_g3_i10	ATP-binding cassette sub-family G member 2	-8,63
TRINITY_DN28155_c0_g1_i3	Vacuolar protein sorting-associated protein 8 homolog	-8,63
TRINITY_DN31258_c1_g2_i1	Serine/threonine-protein phosphatase 6 regulatory subunit 2	-8,63
TRINITY_DN32714_c0_g1_i7	Sortilin	-8,60
TRINITY_DN26718_c2_g1_i7	A disintegrin and metalloproteinase with thrombospondin motif.	-8,60
TRINITY_DN27326_c3_g1_i7	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	-8,60
TRINITY_DN28760_c2_g1_i6	Protein ABHD18	-8,60
TRINITY_DN31945_c1_g1_i12	Protein diaphanous homolog 2	-8,60
TRINITY_DN28763_c0_g2_i4	Cullin-1	-8,60
TRINITY_DN31483_c0_g2_i4	Ras GTPase-activating protein 1	-8,60
TRINITY_DN26848_c0_g1_i5	Peroxisome proliferator-activated receptor delta	-8,60
TRINITY_DN32059_c1_g3_i14	Complement component C7	-8,60
TRINITY_DN28060_c1_g1_i19	AKT-interacting protein	-8,60
TRINITY_DN21550_c0_g1_i1	Complement C1q tumor necrosis factor-related protein 3	-8,60
TRINITY_DN26221_c0_g1_i3	Bifunctional 3'-phosphoadenosine 5'-phosphosulfate synthase 2	-8,60
TRINITY_DN26970_c1_g1_i10	Coiled-coil-helix-coiled-coil-helix domain-containing protein 2	-8,60
TRINITY_DN24501_c0_g1_i1	Serine/arginine-rich splicing factor 1A	-8,60
TRINITY_DN29303_c0_g1_i8	E3 ubiquitin-protein ligase MYCBP2	-8,60
TRINITY_DN32206_c0_g1_i11	Molybdenum cofactor sulfurase	-8,60
TRINITY_DN25686_c0_g1_i4	BRD4-interacting chromatin-remodeling complex-associated pr	-8,60
TRINITY_DN25554_c0_g1_i14	Serine/threonine-protein phosphatase PP1-gamma catalytic suk	-8,60
TRINITY_DN30433_c0_g3_i3	F-box/LRR-repeat protein 17	-8,60
TRINITY_DN28489_c2_g2_i6	Ubiquitin carboxyl-terminal hydrolase 25	-8,60
TRINITY_DN28763_c0_g3_i5	Cullin-1	-8,60
TRINITY_DN28940_c0_g1_i1	Glyoxylate reductase/hydroxypyruvate reductase	-8,60
TRINITY_DN26809_c0_g1_i5	Protein kinase C iota type	-8,60
TRINITY_DN31402_c3_g1_i22	Glucose-6-phosphatase (Fragment)	-8,60
TRINITY_DN32475_c5_g1_i24	UTPglucose-1-phosphate uridylyltransferase	-8,57
TRINITY_DN23829_c0_g1_i3	Nucleoside diphosphate-linked moiety X motif 17	-8,57
TRINITY_DN24529_c0_g1_i3	CD9 antigen	-8,57 -8,57
TRINITY_DN27560_c0_g2_i1	ETS domain-containing protein Elk-1	-8,57 -8,57
TRINITY_DN32459_c2_g1_i4		
TRINITY_DN32207_c0_g1_i8	Zinc finger FYVE domain-containing protein 1	-8,57
	Pseudouridine-5'-phosphate glycosidase	-8,57
TRINITY_DN26285_c2_g1_i2	U6 small nuclear RNA (adenine-(43)-N(6))-methyltransferase	-8,57
TRINITY_DN25285_c4_g1_i5	Transmembrane protease serine 6	-8,57
TRINITY_DN30044_c3_g1_i1	Elongator complex protein 2	-8,57
TRINITY_DN29134_c0_g2_i4	Protein NipSnap homolog 2	-8,57
TRINITY_DN27420_c1_g3_i3	Translocon-associated protein subunit beta	-8,57
TRINITY_DN28362_c1_g2_i2	E3 ubiquitin-protein ligase RNF130	-8,57
TRINITY_DN27816_c1_g1_i4	Vascular endothelial growth factor A-A	-8,57
TRINITY_DN30751_c0_g2_i4	F-box/LRR-repeat protein 20	-8,57
TRINITY_DN28082_c0_g3_i4	Hemicentin-1	-8,57
TRINITY_DN28040_c1_g1_i7	Putative sodium-coupled neutral amino acid transporter 7	-8,57
TRINITY_DN32396_c0_g1_i8	Torsin-1A	-8,57
TRINITY_DN28263_c0_g1_i13	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	-8,57
TRINITY_DN22777_c0_g1_i10	Complement C3 (Fragment)	-8,57

TDDDUTEN DN21000 1 2 7		0.55
TRINITY_DN31998_c1_g3_i7	Zinc finger Y-chromosomal protein 1	-8,57
TRINITY_DN31022_c2_g1_i2	Glucuronokinase 1	-8,57
TRINITY_DN31914_c0_g1_i21	Meprin A subunit beta	-8,57
TRINITY_DN24443_c3_g2_i3	OTU domain-containing protein 5-A	-8,57
TRINITY_DN29349_c1_g1_i1	ATP-dependent RNA helicase DDX19A	-8,53
TRINITY_DN25897_c0_g1_i13	Copine-4	-8,53
TRINITY_DN33110_c2_g2_i7	Myomegalin	-8,53
TRINITY_DN24161_c3_g2_i4	Kelch-like protein 28	-8,53
TRINITY_DN29699_c0_g1_i5	A-kinase anchor protein 8-like	-8,53
TRINITY_DN24747_c4_g1_i3	Ras-related protein Rab-10	-8,53
TRINITY_DN27318_c1_g1_i11	Leucine-rich repeat-containing protein 57	-8,53
TRINITY_DN32643_c1_g1_i3	Max dimerization protein 4	-8,53
TRINITY_DN29121_c0_g1_i4	Solute carrier family 35 member F6	-8,53
TRINITY_DN33241_c2_g1_i7	ATP-binding cassette sub-family G member 8	-8,53
TRINITY_DN28749_c3_g1_i7	CCR4-NOT transcription complex subunit 3	-8,53
TRINITY_DN30160_c1_g2_i5	Growth factor receptor-bound protein 2	-8,53
TRINITY_DN27557_c2_g1_i3	Serine/threonine-protein kinase TAO1	-8,53
TRINITY_DN32785_c3_g1_i1	Heme-binding protein 2	-8,53
TRINITY_DN25021_c0_g1_i10	AP-4 complex subunit beta-1	-8,53
TRINITY_DN26858_c0_g1_i4	Calsyntenin-3	-8,53
TRINITY_DN30949_c0_g1_i5	BUB3-interacting and GLEBS motif-containing protein ZNF207	-8,53
TRINITY_DN32423_c1_g2_i20	Endosomal/lysosomal potassium channel TMEM175	-8,53
TRINITY_DN32792_c1_g1_i14	Alkaline ceramidase 3	-8,53
TRINITY_DN26002_c0_g1_i8	Tudor domain-containing protein 3	-8,53
TRINITY_DN23933_c2_g2_i12	Zinc finger protein 318	-8,53
TRINITY_DN26845_c2_g1_i1	Coagulation factor IX	-8,50
TRINITY_DN29264_c4_g2_i2	Zinc transporter 6	-8,50
TRINITY_DN24839_c4_g2_i13	TIP41-like protein	-8,50
TRINITY_DN27789_c2_g2_i14	Annexin A6	-8,50 -8,50
TRINITY_DN30662_c0_g2_i2	Villin-1	-8,50 -8,50
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TRINITY_DN29313_c4_g2_i8	Zinc finger protein ubi-d4	-8,50
TRINITY_DN25286_c0_g2_i5	Host cell factor 1	-8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1	-8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6	-8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4	-8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase	-8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN29787_c2_g2_i3	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN3162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN2946_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN3162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN28802_c1_g1_i3	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN328287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8	-8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN29046_c0_g1_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN28802_c1_g1_i3	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7	-8,50 -8,50
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN328287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN3162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN31226_c4_g1_i4 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN32433_c1_g1_i3 TRINITY_DN24500_c0_g1_i7	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog	-8,50 -8,47 -8,47
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN328287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN26977_c4_g1_i6 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32433_c1_g1_i3 TRINITY_DN24500_c0_g1_i7 TRINITY_DN32933_c0_g1_i2	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog GRB10-interacting GYF protein 2	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,47 -8,47 -8,47
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN32440_c1_g2_i17 TRINITY_DN30159_c0_g1_i3 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN33162_c1_g1_i7 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32933_c0_g1_i2 TRINITY_DN32933_c0_g1_i2 TRINITY_DN32933_c0_g1_i2 TRINITY_DN33063_c4_g3_i10	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog GRB10-interacting GYF protein 2 Bone morphogenetic protein 1	-8,50 -8,47 -8,47 -8,47
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN328287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32433_c1_g1_i3 TRINITY_DN324500_c0_g1_i7 TRINITY_DN32933_c0_g1_i2 TRINITY_DN33063_c4_g3_i10 TRINITY_DN27843_c0_g1_i13	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog GRB10-interacting GYF protein 2 Bone morphogenetic protein 1 Protein PRR14L	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,47 -8,47 -8,47 -8,47
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN23787_c2_g2_i3 TRINITY_DN3162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN31226_c4_g1_i4 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25086_c3_g1_i15 TRINITY_DN26977_c4_g1_i6 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN24500_c0_g1_i7 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32933_c0_g1_i2 TRINITY_DN32933_c0_g1_i2 TRINITY_DN32063_c4_g3_i10 TRINITY_DN27843_c0_g1_i13 TRINITY_DN27843_c0_g1_i13	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog GRB10-interacting GYF protein 2 Bone morphogenetic protein 1 Protein PRR14L Elongation of very long chain fatty acids protein 1	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,47 -8,47 -8,47 -8,47 -8,47
TRINITY_DN25286_c0_g2_i5 TRINITY_DN30541_c0_g1_i12 TRINITY_DN30541_c0_g1_i12 TRINITY_DN32395_c1_g2_i4 TRINITY_DN32039_c1_g1_i9 TRINITY_DN32440_c1_g2_i17 TRINITY_DN28287_c0_g2_i6 TRINITY_DN30159_c0_g1_i3 TRINITY_DN32965_c0_g1_i12 TRINITY_DN32945_c2_g2_i13 TRINITY_DN32945_c2_g2_i13 TRINITY_DN23787_c2_g2_i3 TRINITY_DN33162_c1_g1_i7 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31547_c5_g3_i5 TRINITY_DN31226_c4_g1_i4 TRINITY_DN28369_c1_g1_i6 TRINITY_DN25086_c3_g1_i14 TRINITY_DN25781_c0_g1_i2 TRINITY_DN26977_c4_g1_i6 TRINITY_DN28802_c1_g1_i8 TRINITY_DN28802_c1_g1_i8 TRINITY_DN24500_c0_g1_i7 TRINITY_DN32433_c1_g1_i3 TRINITY_DN32433_c0_g1_i2 TRINITY_DN32933_c0_g1_i2 TRINITY_DN326467_c0_g1_i13 TRINITY_DN27843_c0_g1_i13 TRINITY_DN27848_c0_g1_i5	Host cell factor 1 5'-AMP-activated protein kinase subunit beta-1 Biogenesis of lysosome-related organelles complex 1 subunit 6 Sodium-coupled neutral amino acid transporter 4 NAD kinase G/T mismatch-specific thymine DNA glycosylase Myotubularin-related protein 14 N-lysine methyltransferase SETD6 Tricarboxylate transport protein, mitochondrial Nucleoporin NDC1 RING finger protein 215 Transmembrane protein KIAA1109 homolog EH domain-containing protein 3 Zinc finger and BTB domain-containing protein 5 Hippocampus abundant transcript 1 protein ATP-binding cassette sub-family G member 2 Chromodomain-helicase-DNA-binding protein 4 Far upstream element-binding protein 1 Solute carrier family 12 member 8 Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 Exonuclease mut-7 homolog GRB10-interacting GYF protein 2 Bone morphogenetic protein 1 Protein PRR14L Elongation of very long chain fatty acids protein 1 Cohesin subunit SA-2	-8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,50 -8,47 -8,47 -8,47 -8,47 -8,47 -8,47

TDINITY DN21707 -4 -2 :5	CIDI Jamain and sining made in 1	0.47
TRINITY_DN31787_c4_g2_i5	SUN domain-containing protein 1	-8,47
TRINITY_DN30989_c0_g1_i8	Translocation protein SEC62	-8,47
TRINITY_DN32671_c0_g1_i3	3-phosphoinositide-dependent protein kinase 1	-8,47
TRINITY_DN28293_c0_g1_i10	Copine-8	-8,47
TRINITY_DN26361_c0_g1_i4	Protein ABHD11	-8,47
TRINITY_DN32377_c1_g1_i7	Glycerol-3-phosphate acyltransferase 1, mitochondrial	-8,47
TRINITY_DN27629_c0_g1_i14	Peroxidasin	-8,47
TRINITY_DN29299_c0_g1_i1	Protein disulfide-isomerase TMX3	-8,47
TRINITY_DN26063_c0_g3_i7	FERM, ARHGEF and pleckstrin domain-containing protein 1	-8,47
TRINITY_DN32392_c0_g1_i7	E3 ubiquitin-protein ligase ubr3	-8,47
TRINITY_DN24769_c1_g1_i2	Forkhead box protein N4	-8,47
TRINITY_DN31661_c0_g1_i12	High affinity cationic amino acid transporter 1	-8,47
TRINITY_DN25399_c2_g2_i3	Ephrin type-B receptor 4	-8,47
TRINITY_DN28833_c1_g1_i9	Casein kinase I isoform alpha	-8,47
TRINITY_DN32484_c0_g1_i3	Apolipoprotein L6	-8,47
TRINITY_DN29677_c1_g1_i16	Nectin-4	-8,47
TRINITY_DN27905_c0_g1_i4	Granulins	-8,44
TRINITY_DN26252_c1_g1_i9	Focal adhesion kinase 1	-8,44
TRINITY_DN32053_c2_g1_i2	Golgin subfamily A member 3	-8,44
TRINITY_DN31202_c0_g1_i2	1-phosphatidylinositol 3-phosphate 5-kinase	-8,44
TRINITY_DN26864_c1_g1_i4	Coiled-coil domain-containing protein 134	-8,44
TRINITY_DN28693_c0_g1_i1	WD repeat-containing protein 41	-8,44
TRINITY_DN31350_c0_g1_i9	Tumor susceptibility gene 101 protein	-8,44 -8,44
_	Cell death activator CIDE-3	
TRINITY_DN33024_c5_g3_i6		-8,44
TRINITY_DN31275_c1_g1_i15	Glucosidase 2 subunit beta	-8,44
TRINITY_DN32045_c0_g1_i7	Scaffold attachment factor B1	-8,44
TRINITY_DN26636_c0_g1_i2	Serine/threonine-protein kinase STK11	-8,44
TRINITY_DN24099_c0_g1_i7	Uncharacterized protein KIAA1211-like	-8,44
TRINITY_DN32582_c0_g1_i2	RING finger protein 10	-8,44
TRINITY_DN27070_c0_g1_i4	Glutamine amidotransferase-like class 1 domain-containing pro	-8,44
TRINITY_DN27843_c0_g1_i17	Protein PRR14L	-8,44
TRINITY_DN32367_c2_g1_i13	ERBB receptor feedback inhibitor 1	-8,44
TRINITY_DN29884_c0_g3_i2	Microtubule-associated protein RP/EB family member 3	-8,44
TRINITY_DN28925_c2_g1_i2	Homeobox-containing protein 1	-8,44
TRINITY_DN31157_c1_g1_i2	Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondr	-8,44
TRINITY_DN31513_c0_g1_i12	Peroxisomal targeting signal 1 receptor	-8,44
TRINITY_DN32405_c0_g1_i6	CDK5 regulatory subunit-associated protein 2	-8,44
TRINITY_DN25201_c0_g1_i7	Cleavage stimulation factor subunit 1	-8,44
TRINITY_DN23933_c2_g2_i6	Surface protein	-8,44
TRINITY_DN28061_c0_g1_i1	Leucine-rich repeat-containing G-protein coupled receptor 4	-8,44
TRINITY_DN30062_c1_g3_i2	Disheveled-associated activator of morphogenesis 2	-8,44
TRINITY_DN22060_c0_g2_i7	Transgelin-3	-8,44
TRINITY_DN22913_c0_g1_i8	Armadillo repeat-containing protein 8	-8,44
TRINITY_DN26717_c0_g1_i4	High mobility group-T protein	-8,44
TRINITY_DN23962_c0_g1_i4	ATP-dependent RNA helicase DHX33	-8,44
TRINITY_DN28317_c2_g1_i8	Beclin-1	-8,44
TRINITY_DN29564_c0_g1_i9	Interferon-induced protein with tetratricopeptide repeats 1	-8,44 -8,44
TRINITY_DN28638_c0_g1_i1		-8,44 -8,44
_	R3H domain-containing protein 2	
TRINITY_DN26253_c0_g1_i4	Centrosomal protein of 164 kDa	-8,40
TRINITY_DN28344_c2_g1_i1	Myotubularin-related protein 3	-8,40
TRINITY_DN27639_c1_g3_i3	Nibrin	-8,40
TRINITY_DN29676_c1_g3_i2	ATP-dependent 6-phosphofructokinase, platelet type	-8,40
TRINITY_DN32331_c1_g1_i13	Hepatocyte nuclear factor 1-alpha	-8,40
TRINITY_DN26313_c4_g1_i4	Tudor and KH domain-containing protein	-8,40
TRINITY_DN33079_c1_g1_i12	Solute carrier family 41 member 1	-8,40
TRINITY_DN26295_c0_g1_i6	Kelch repeat and BTB domain-containing protein 2	-8,40
TRINITY_DN26161_c0_g1_i11	Protein argonaute-3	-8,40
TRINITY_DN28648_c1_g1_i5	Insulin-like growth factor 2 mRNA-binding protein 2	-8,40
TRINITY_DN28053_c0_g2_i14	Heat shock cognate 71 kDa protein	-8,40

TRINITY_DN32777_c0_g1_i3 Histone-lysine N-methyltransferase 2A -8, TRINITY_DN31677_c2_g1_i6 Solute carrier family 25 member 47-B -8, TRINITY_DN29726_c3_g1_i6 Aldehyde dehydrogenase, mitochondrial -8, TRINITY_DN28782_c0_g1_i18 H-2 class II histocompatibility antigen, E-U alpha chain -8, TRINITY_DN28793_c0_g1_i5 SWI/SNF-related matrix-associated actin-dependent regulator ε -8, TRINITY_DN31495_c0_g1_i5 Cation-dependent mannose-6-phosphate receptor -8,	,40 ,40 ,40 ,40 ,40 ,40 ,40 ,40
TRINITY_DN31677_c2_g1_i6 Solute carrier family 25 member 47-B -8, TRINITY_DN29726_c3_g1_i6 Aldehyde dehydrogenase, mitochondrial -8, TRINITY_DN28782_c0_g1_i18 H-2 class II histocompatibility antigen, E-U alpha chain -8, TRINITY_DN28793_c0_g1_i5 SWI/SNF-related matrix-associated actin-dependent regulator c -8, TRINITY_DN31495_c0_g1_i5 Cation-dependent mannose-6-phosphate receptor -8,	,40 ,40 ,40 ,40 ,40
TRINITY_DN29726_c3_g1_i6 Aldehyde dehydrogenase, mitochondrial -8, TRINITY_DN28782_c0_g1_i18 H-2 class II histocompatibility antigen, E-U alpha chain -8, TRINITY_DN28793_c0_g1_i5 SWI/SNF-related matrix-associated actin-dependent regulator c -8, TRINITY_DN31495_c0_g1_i5 Cation-dependent mannose-6-phosphate receptor -8,	,40 ,40 ,40 ,40
TRINITY_DN28782_c0_g1_i18	,40 ,40 ,40
TRINITY_DN28793_c0_g1_i5	,40 ,40
TRINITY_DN31495_c0_g1_i5 Cation-dependent mannose-6-phosphate receptor -8,	,40
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TRINITY_DN25448_c1_g2_i4 Forkhead box protein P1-B -8,	,36
TRINITY_DN23469_c0_g1_i10 4-hydroxy-2-oxoglutarate aldolase, mitochondrial -8,	,36
TRINITY_DN29130_c0_g3_i8 Kinase suppressor of Ras 1 -8,	,36
TRINITY_DN28903_c0_g1_i1 Eukaryotic translation initiation factor 3 subunit K -8,	,36
TRINITY_DN30921_c1_g2_i25 Unconventional myosin-X -8,	,36
TRINITY_DN22826_c0_g1_i3 NF-kappa-B inhibitor alpha -8,	,36
TRINITY_DN24824_c0_g3_i12 Repressor of RNA polymerase III transcription MAF1 homolog -8,	,36
TRINITY_DN24810_c2_g1_i19 Heterogeneous nuclear ribonucleoprotein H3 -8,	,36
TRINITY_DN30813_c0_g1_i2 Protein phosphatase 1 regulatory subunit 3B -8,	,36
TRINITY_DN25934_c2_g2_i5 Ras-related protein Rap-1b -8,	,36
TRINITY_DN23294_c0_g1_i3 Proline-rich transmembrane protein 4 -8,	,36
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TRINITY_DN21963_c0_g1_i2 Sn1-specific diacylglycerol lipase alpha -8,	,36
TRINITY_DN30766_c2_g2_i3 Protein asteroid homolog 1 -8,	,36
TRINITY_DN31181_c0_g1_i8 Zinc finger RNA-binding protein -8,	,36
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TRINITY_DN23954_c1_g2_i14 Ciliogenesis and planar polarity effector 1 -8,	,36
TRINITY_DN26762_c0_g1_i1 Extended synaptotagmin-2 -8,	,36
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TRINITY_DN30378_c3_g1_i9	Zinc finger MYM-type protein 1	-8,33
TRINITY_DN31437_c1_g1_i1	Estrogen receptor beta-2	-8,33
TRINITY_DN30419_c0_g1_i18	Urotensin-2 receptor	-8,33
TRINITY_DN22730_c0_g1_i8	UBX domain-containing protein 7	-8,33
TRINITY_DN28784_c0_g1_i11		-8,33
	Arf-GAP domain and FG repeat-containing protein 1	
TRINITY_DN33168_c3_g2_i7	Proteasome activator complex subunit 4A	-8,33
TRINITY_DN29922_c2_g1_i13	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	-8,33
TRINITY_DN29397_c1_g1_i2	Bombesin receptor-activated protein C6orf89 homolog	-8,33
TRINITY_DN25690_c0_g1_i8	Haloacid dehalogenase-like hydrolase domain-containing 5	-8,33
TRINITY_DN28592_c0_g1_i2	Serine/threonine-protein phosphatase 6 regulatory ankyrin repe	-8,33
TRINITY_DN29362_c1_g1_i2	Protein FAM214B	-8,33
TRINITY_DN27152_c0_g2_i6	Synaptotagmin-like protein 2	-8,33
TRINITY_DN26697_c5_g1_i5	FSD1-like protein	-8,33
TRINITY_DN25525_c0_g1_i9	Centrosomal protein of 85 kDa-like	-8,33
TRINITY_DN31630_c1_g1_i6	Integrin alpha-9	-8,33
TRINITY_DN33240_c9_g1_i3	Complement C3 (Fragment)	-8,33
TRINITY_DN25005_c4_g1_i2	Vacuolar protein sorting-associated protein 37A	-8,33
TRINITY_DN28890_c4_g1_i11	Rhodopsin, GQ-coupled	-8,33
TRINITY_DN32994_c2_g1_i2	Protein O-GlcNAcase	-8,33
TRINITY_DN30833_c2_g1_i1	Stathmin-2	-8,33
TRINITY_DN25948_c1_g1_i4	Lysyl oxidase homolog 2B	-8,33
TRINITY_DN28595_c1_g1_i10	DNA polymerase lambda	-8,29
TRINITY_DN26891_c0_g1_i10	Lipoprotein lipase	-8,29
TRINITY_DN28394_c0_g1_i4	RNA 3'-terminal phosphate cyclase	-8,29
TRINITY_DN30249_c1_g1_i19	Tight junction protein ZO-2	-8,29
TRINITY_DN24672_c4_g2_i1	Rhophilin-1	-8,29
TRINITY_DN32162_c0_g2_i7	FYVE, RhoGEF and PH domain-containing protein 5	-8,29
TRINITY_DN25731_c4_g1_i1	Phosphatidylinositol 3,4,5-trisphosphate 3-phosphatase and du	-8,29
TRINITY_DN32497_c1_g1_i7	Tuberin	-8,29
TRINITY_DN31625_c2_g2_i3	Probable imidazolonepropionase	-8,29
TRINITY_DN33009_c1_g1_i7	Zinc finger protein 616	-8,29
TRINITY_DN22700_c0_g1_i2	Queuosine salvage protein	-8,29
TRINITY_DN31079_c0_g1_i5		
	Stimulated by retinoic acid gene 6 protein-like	-8,29
TRINITY_DN29833_c1_g1_i2	CTP synthase 1	-8,29
TRINITY_DN27629_c0_g1_i21	Peroxidasin	-8,29
TRINITY_DN27072_c0_g1_i11	Histone deacetylase 7	-8,29
TRINITY_DN31335_c0_g1_i2	Transitional endoplasmic reticulum ATPase	-8,29
TRINITY_DN26587_c0_g1_i1	H/ACA ribonucleoprotein complex subunit 2-like protein	-8,29
TRINITY_DN27546_c0_g1_i9	GEM-interacting protein	-8,29
TRINITY_DN31044_c1_g3_i2	Pyruvate dehydrogenase E1 component subunit beta, mitochonc	-8,29
TRINITY_DN28461_c0_g1_i15	HMG domain-containing protein 3	-8,29
TRINITY_DN31517_c0_g1_i6	Oxysterol-binding protein-related protein 9	-8,29
TRINITY_DN32079_c1_g2_i1	Cation-independent mannose-6-phosphate receptor	-8,29
TRINITY_DN32734_c1_g2_i2	Myosin-9	-8,29
TRINITY_DN30429_c0_g1_i14	Coiled-coil domain-containing protein 77	-8,29
TRINITY_DN29261_c0_g1_i7	DNA excision repair protein ERCC-6	-8,29
TRINITY_DN30235_c0_g1_i6	Diacylglycerol kinase delta	-8,29
TRINITY_DN26196_c0_g1_i5	Bromodomain-containing protein 3	-8,29
TRINITY_DN25821_c0_g2_i1	Dual specificity protein phosphatase 6	-8,29
TRINITY_DN27256_c0_g1_i2	GPI mannosyltransferase 3	-8,29
TRINITY_DN26707_c1_g1_i8	Neuropathy target esterase sws	-8,29
TRINITY_DN26218_c0_g1_i4	Lysine-specific demethylase 6B	-8,25
TRINITY_DN33172_c0_g1_i1	Elongation of very long chain fatty acids protein 2	-8,25
TRINITY_DN28592_c0_g1_i7	Serine/threonine-protein phosphatase 6 regulatory ankyrin repe	-8,25
TRINITY_DN31302_c1_g1_i9	Long-chain-fatty-acidCoA ligase 4	-8,25
TRINITY_DN27458_c0_g1_i9	EF-hand calcium-binding domain-containing protein 4A	-8,25
TRINITY_DN22840_c0_g1_i9	Growth arrest and DNA damage-inducible protein GADD45 ga	-8,25
TRINITY_DN32166_c4_g1_i1	NADPH oxidase 5	-8,25
TRINITY_DN29235_c0_g1_i2	Coiled-coil domain-containing protein 97	-8,25
111111111111111111111111111111111111111	concer con domain containing protein >1	0,23

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TRINITY_DN27367_c1_g1_i13	Proprotein convertase subtilisin/kexin type 5	-8,25
TRINITY_DN32104_c0_g2_i10	Sequestosome-1	-8,25
TRINITY_DN26718_c2_g1_i10	A disintegrin and metalloproteinase with thrombospondin motif.	-8,25
TRINITY_DN30084_c0_g1_i7	Zinc fingers and homeoboxes protein 3	-8,25
TRINITY_DN29131_c0_g2_i7	Nuclear cap-binding protein subunit 3	-8,25
TRINITY_DN29349_c1_g1_i2	ATP-dependent RNA helicase DDX19A	-8,25
TRINITY_DN32698_c1_g1_i12	Rab-like protein 6	-8,25
TRINITY_DN28187_c0_g2_i5	PHD finger protein 13	-8,25
TRINITY_DN26457_c3_g1_i2	Retrovirus-related Pol polyprotein from transposon 17.6	-8,25
TRINITY_DN28399_c0_g1_i3	Telomere length regulation protein TEL2 homolog	-8,25
TRINITY_DN29929_c1_g2_i19	Sushi, nidogen and EGF-like domain-containing protein 1	-8,25
TRINITY_DN31416_c0_g1_i3	Glucocorticoid receptor	-8,25
TRINITY_DN29307_c0_g1_i10	LYR motif-containing protein 4	-8,25
TRINITY_DN25854_c0_g1_i1	Zinc finger protein 516	-8,25
TRINITY_DN31173_c0_g2_i15	Ensconsin	-8,25 -8,25
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TRINITY_DN32393_c3_g1_i2	2-hydroxyacyl-CoA lyase 1	-8,25
TRINITY_DN32852_c1_g1_i1	Selenoprotein Pb	-8,25
TRINITY_DN30010_c0_g2_i6	Lipoamide acyltransferase component of branched-chain alpha	-8,25
TRINITY_DN30081_c0_g1_i8	GPI ethanolamine phosphate transferase 1	-8,25
TRINITY_DN30195_c1_g2_i5	Sphingomyelin phosphodiesterase 4	-8,25
TRINITY_DN31052_c0_g1_i29	Transposon Ty3-G Gag-Pol polyprotein	-8,25
TRINITY_DN28289_c0_g1_i8	P2X purinoceptor 5	-8,25
TRINITY_DN32025_c2_g1_i1	Immunoglobulin lambda-1 light chain	-8,25
TRINITY_DN25126_c0_g1_i2	Exportin-4	-8,25
TRINITY_DN23801_c0_g1_i1	ATP-sensitive inward rectifier potassium channel 10	-8,25
TRINITY_DN23371_c0_g1_i1	Occludin	-8,25
TRINITY_DN29548_c0_g1_i2	SWI/SNF-related matrix-associated actin-dependent regulator c	-8,25
TRINITY_DN28943_c2_g1_i4	Nuclear pore complex protein Nup54	-8,21
TRINITY_DN32037_c0_g1_i15	Rho GTPase-activating protein 27	-8,21
TRINITY_DN31280_c1_g1_i2	Phosphatidylinositol-binding clathrin assembly protein	-8,21
TRINITY_DN27203_c2_g1_i8	3'(2'),5'-bisphosphate nucleotidase 1	-8,21
TRINITY_DN32370_c2_g1_i8	Trafficking protein particle complex subunit 6B	-8,21
TRINITY_DN31052_c0_g1_i3	Transposon Ty3-I Gag-Pol polyprotein	-8,21
TRINITY_DN24163_c2_g1_i15	von Willebrand factor A domain-containing protein 7	-8,21
TRINITY_DN25027_c0_g1_i1	R3H domain-containing protein 2	-8,21 -8,21
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TRINITY_DN29583_c0_g1_i4	Serine/threonine-protein kinase SIK3 homolog	-8,21
TRINITY_DN26721_c3_g1_i6	Probable C->U-editing enzyme APOBEC-2	-8,21
TRINITY_DN32084_c1_g1_i11	Zinc finger MYM-type protein 4	-8,21
TRINITY_DN27494_c2_g2_i2	UPF0472 protein C16orf72 homolog	-8,21
TRINITY_DN24739_c1_g1_i9	Transforming protein p29	-8,21
TRINITY_DN31471_c1_g2_i3	Protein KIAA0100	-8,21
TRINITY_DN28544_c0_g1_i2	2',5'-phosphodiesterase 12	-8,21
TRINITY_DN32426_c1_g1_i15	Protein transport protein Sec24C	-8,21
TRINITY_DN28568_c0_g2_i6	Calcium-activated potassium channel subunit alpha-1	-8,21
TRINITY_DN33137_c5_g1_i11	Dimethylglycine dehydrogenase, mitochondrial	-8,21
TRINITY_DN26794_c1_g2_i1	Epidermal growth factor-like protein 7	-8,21
TRINITY_DN28876_c0_g1_i5	Protein phosphatase inhibitor 2	-8,21
TRINITY_DN25106_c1_g1_i4	11-cis retinol dehydrogenase	-8,21
TRINITY_DN32715_c0_g2_i9	Receptor-type tyrosine-protein phosphatase F	-8,21
TRINITY_DN33182_c5_g2_i11	Fructose-1,6-bisphosphatase 1	-8,21
TRINITY_DN31483_c0_g2_i7	Ras GTPase-activating protein 1	-8,21
TRINITY_DN32781_c0_g1_i25	SUN domain-containing	-8,21
TRINITY_DN33202_c5_g1_i1	Cytochrome P450 3A56	-8,21 -8,21
TRINITY_DN25204_c2_g1_i15		-8,21 -8,21
	Serine/threonine-protein kinase MARK2	
TRINITY_DN29667_c0_g1_i1	Iron-responsive element-binding protein 2	-8,21
TRINITY_DN23441_c0_g2_i5	Transformer-2 protein homolog beta	-8,21
TRINITY_DN31797_c0_g1_i5	Phosphatase and actin regulator 4A	-8,21
TRINITY_DN30839_c0_g1_i3	E3 ubiquitin-protein ligase pellino homolog 1	-8,21
TRINITY_DN25915_c0_g1_i5	V-type proton ATPase catalytic subunit A	-8,21

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TRINITY_DN29855_c0_g2_i8	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyli	-8,21
TRINITY_DN30921_c1_g2_i9	Unconventional myosin-X	-8,21
TRINITY_DN29930_c2_g1_i12	Complement C1q-like protein 3	-8,21
TRINITY_DN31567_c1_g2_i12	CREB-regulated transcription coactivator 3	-8,21
TRINITY_DN31174_c0_g2_i6	UDP-glucuronosyltransferase 2A1	-8,21
TRINITY_DN23444_c0_g1_i7	Cerebellin-3	-8,17
TRINITY_DN29313_c4_g3_i7	Zinc finger protein ubi-d4	-8,17
TRINITY_DN24857_c6_g4_i1	60S ribosomal protein L28	-8,17
TRINITY_DN32209_c0_g1_i1		
_	Hepatocyte growth factor receptor	-8,17
TRINITY_DN26418_c1_g1_i3	Mitogen-activated protein kinase 14A	-8,17
TRINITY_DN22976_c0_g1_i3	Probable asparaginetRNA ligase, mitochondrial	-8,17
TRINITY_DN25075_c3_g1_i1	E3 ubiquitin-protein ligase RNF185	-8,17
TRINITY_DN25498_c0_g1_i9	F-box only protein 32	-8,17
TRINITY_DN25045_c0_g1_i1	Mitochondrial glycine transporter B	-8,17
TRINITY_DN26954_c4_g1_i13	RUS1 family protein C16orf58 homolog	-8,17
TRINITY_DN26862_c4_g1_i15	Transportin-1	-8,17
TRINITY_DN31052_c0_g1_i19	Transposon Ty3-G Gag-Pol polyprotein	-8,17
TRINITY_DN31914_c0_g1_i4	Meprin A subunit beta	-8,17
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TRINITY_DN32263_c0_g1_i5	Dual specificity tyrosine-phosphorylation-regulated kinase 1A	-8,17
TRINITY_DN31881_c1_g1_i3	ATP-citrate synthase	-8,17
TRINITY_DN26550_c0_g1_i1	Complement factor B	-8,17
TRINITY_DN28568_c0_g2_i13	Calcium-activated potassium channel subunit alpha-1	-8,17
TRINITY_DN29063_c0_g1_i9	Nucleosome assembly protein 1-like 4	-8,17
TRINITY_DN30327_c2_g1_i2	Beta-galactosidase-1-like protein 2	-8,17
TRINITY_DN29031_c2_g3_i6	Coiled-coil domain-containing protein 148	-8,17
TRINITY_DN31158_c0_g1_i9	Testis-specific Y-encoded-like protein 1	-8,17
TRINITY_DN29322_c0_g1_i2	Two pore calcium channel protein 2	-8,17
TRINITY_DN26976_c0_g2_i1	Prostaglandin E synthase 3	-8,17
TRINITY_DN30797_c1_g1_i6	Coenzyme Q-binding protein COQ10 homolog, mitochondrial	-8,17
TRINITY_DN26988_c0_g1_i3	Girdin	-8,17
TRINITY_DN30318_c0_g2_i15	Alpha-2-macroglobulin-like protein 1	-8,17
TRINITY_DN31786_c0_g2_i8	Ligand of Numb protein X 2	-8,17
TRINITY_DN32693_c7_g1_i2	Interleukin-1 receptor type 2	-8,17
TRINITY_DN28311_c0_g1_i1	Integral membrane protein 2B	-8,17
TRINITY_DN26942_c3_g2_i4	Isoaspartyl peptidase/L-asparaginase	-8,17
TRINITY_DN27847_c5_g1_i7	Sorting nexin-12	-8,17
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TRINITY_DN28165_c2_g3_i4	Ribosomal protein S6 kinase alpha-1	-8,17
TRINITY_DN29515_c0_g1_i9	14-3-3 protein theta	-8,13
TRINITY_DN26102_c2_g2_i4	Complement C3 (Fragment)	-8,13
TRINITY_DN30687_c0_g1_i23	Cyclin-G-associated kinase	-8,13
TRINITY_DN28358_c4_g1_i3	THAP domain-containing protein 7	-8,13
TRINITY_DN31997_c0_g1_i1	Eukaryotic translation initiation factor 4 gamma 3	-8,13
TRINITY_DN31110_c1_g1_i9	Histone-lysine N-methyltransferase, H3 lysine-79 specific	-8,13
TRINITY_DN25375_c0_g1_i1	E3 ubiquitin-protein ligase rnf168	-8,13
TRINITY_DN24928_c2_g3_i12	Uronyl 2-sulfotransferase	-8,13
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TRINITY_DN27006_c1_g1_i16	Sodium- and chloride-dependent GABA transporter 2	-8,13
TRINITY_DN31041_c0_g1_i2	Trinucleotide repeat-containing gene 6B protein	-8,13
TRINITY_DN23358_c0_g1_i5	PHD finger protein 14	-8,13
TRINITY_DN28920_c0_g1_i13	Leucine-rich repeat flightless-interacting protein 2	-8,13
TRINITY_DN28419_c1_g2_i7	AN1-type zinc finger protein 5	-8,13
TRINITY_DN30787_c3_g2_i4	Prospero homeobox protein 1	-8,13
TRINITY_DN29230_c2_g2_i5	RNA-binding protein 39	-8,13
TRINITY_DN27837_c1_g2_i7	Transcription initiation factor TFIID subunit 1	-8,13
TRINITY_DN30454_c1_g2_i2	Protein phosphatase Slingshot homolog 2	-8,13
TRINITY_DN31083_c0_g1_i7	Probable JmjC domain-containing histone demethylation protei	-8,13
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TRINITY_DN27453_c0_g1_i8	Poly [ADP-ribose] polymerase 11	-8,13
TRINITY_DN30131_c0_g1_i4	Sentrin-specific protease 2	-8,13
TRINITY_DN30101_c0_g1_i6	NHS-like protein 1	-8,13
TRINITY_DN25894_c0_g1_i6	Noelin	-8,13

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TRINITY_DN30992_c0_g1_i4	Poly [ADP-ribose] polymerase 14	-8,13
TRINITY_DN33095_c2_g1_i7	Mitogen-activated protein kinase kinase kinase kinase 3	-8,13
TRINITY_DN31058_c0_g2_i9	Moesin	-8,13
TRINITY_DN32960_c1_g1_i7	SNF-related serine/threonine-protein kinase	-8,13
TRINITY_DN31314_c1_g1_i5	Retinol dehydrogenase 8	-8,13
TRINITY_DN32719_c0_g2_i9	Proprotein convertase subtilisin/kexin type 5	-8,13
TRINITY_DN25566_c0_g1_i2	Zinc fingers and homeoboxes protein 1	-8,13
TRINITY_DN32311_c1_g1_i12	Histone deacetylase 4	-8,13
TRINITY_DN32781_c0_g1_i8	SUN domain-containing	-8,13
TRINITY_DN30886_c1_g2_i1	PHD finger protein 12	-8,13
TRINITY_DN24327_c0_g1_i11	Transcription factor PU.1	-8,13
TRINITY_DN31769_c2_g1_i12	GRAM domain-containing protein 2B	-8,13
TRINITY_DN25632_c0_g1_i4	Growth arrest and DNA damage-inducible protein GADD45 be	-8,13
TRINITY_DN26369_c0_g2_i3	Max-binding protein MNT	-8,13
TRINITY_DN25432_c1_g2_i11	60S ribosomal protein L8	-8,13
TRINITY_DN32816_c0_g1_i8	NEDD4-like E3 ubiquitin-protein ligase WWP1	-8,13
TRINITY_DN27328_c1_g1_i4	DNA repair protein RAD50	-8,13
TRINITY_DN30696_c1_g1_i1	Zinc finger CCCH domain-containing protein 4	-8,13
TRINITY_DN26679_c1_g1_i5	Leucine-rich repeat protein 1	-8,13
TRINITY_DN31445_c0_g1_i6	Long-chain fatty acid transport protein 1	-8,13
TRINITY_DN30925_c1_g1_i1	Ubiquitin-associated protein 2	-8,13
TRINITY_DN25915_c0_g1_i4	V-type proton ATPase catalytic subunit A	-8,09
TRINITY_DN31856_c1_g2_i2	Xin actin-binding repeat-containing protein 2	-8,09
TRINITY_DN33282_c5_g1_i10	Regulator of microtubule dynamics protein 3	-8,09
TRINITY_DN32838_c0_g1_i5	Forkhead box protein P4	-8,09
TRINITY_DN23352_c0_g2_i2	Aminoacyl tRNA synthase complex-interacting multifunctional p	-8,09
TRINITY_DN26148_c1_g1_i11	LIM domain-binding protein 1	-8,09
TRINITY_DN24930_c2_g5_i14	Leukocyte cell-derived chemotaxin-2	-8,09
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TRINITY_DN29121_c0_g1_i3	Solute carrier family 35 member F6	
TRINITY_DN29585_c0_g1_i6	Nck-associated protein 1	-8,09
TRINITY_DN27337_c1_g1_i4	Ras association domain-containing protein 9	-8,09
TRINITY_DN29750_c2_g2_i3	Glyoxalase domain-containing protein 4	-8,09
TRINITY_DN28157_c1_g1_i11	Golgi SNAP receptor complex member 1	-8,09
TRINITY_DN27688_c0_g1_i4	Matrix metalloproteinase-9	-8,09
TRINITY_DN32128_c1_g1_i7	Peptidyl-glycine alpha-amidating monooxygenase A	-8,09
TRINITY_DN23491_c0_g1_i6	Epidermal growth factor receptor kinase substrate 8-like protein	-8,09
TRINITY_DN32850_c3_g1_i9	Arrestin domain-containing protein 3	-8,09
TRINITY_DN25435_c0_g1_i1	ATP-dependent RNA helicase DHX36	-8,09
TRINITY_DN30309_c1_g1_i17	Acetyl-coenzyme A synthetase, cytoplasmic	-8,09
TRINITY_DN31704_c2_g1_i21	Interferon-induced very large GTPase 1	-8,09
TRINITY_DN25765_c3_g3_i11	Retinol-binding protein 2	-8,09
TRINITY_DN32781_c0_g1_i27	SUN domain-containing	-8,09
TRINITY_DN29514_c0_g1_i9	Serine/threonine-protein kinase Sgk2	-8,09
TRINITY_DN31992_c2_g1_i8	Complement C1q subcomponent subunit C	-8,09
TRINITY_DN28302_c1_g1_i12	General transcription factor IIH subunit 1	-8,09
TRINITY_DN30130_c5_g1_i6	Tetratricopeptide repeat protein 39B	-8,09
TRINITY_DN29150_c1_g1_i3	FYN-binding protein 1	-8,09
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TRINITY_DN31707_c0_g1_i1	Guanine nucleotide-binding protein subunit beta-4	-8,09
TRINITY_DN26712_c0_g2_i5	Transcription factor Sp3	-8,09
TRINITY_DN28110_c2_g1_i1	Phytanoyl-CoA dioxygenase, peroxisomal	-8,09
TRINITY_DN33119_c2_g1_i3	Syndecan-2	-8,09
TRINITY_DN33110_c2_g1_i3	Myomegalin	-8,09
TRINITY_DN32556_c0_g1_i6	Lysine-specific demethylase 6A	-8,09
TRINITY_DN30264_c0_g1_i2	Integrin alpha-3	-8,09
TRINITY_DN31523_c0_g1_i18	Phosphorylase b kinase regulatory subunit alpha, skeletal musc	-8,09
TRINITY_DN31567_c1_g2_i5	CREB-regulated transcription coactivator 3	-8,09
TRINITY_DN28367_c0_g1_i9	Sodium/hydrogen exchanger 6	-8,09
TRINITY_DN32582_c0_g1_i13	RING finger protein 10	-8,09
TRINITY_DN27866_c1_g1_i8	Syntaxin-2	-8,09
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TRINITY_DN28165_c2_g1_i3	Ribosomal protein S6 kinase alpha-3	-8,09
TRINITY_DN27148_c1_g3_i15	Coiled-coil domain-containing protein 50	-8,09
TRINITY_DN26278_c3_g1_i9	Sphingomyelin synthase-related protein 1	-8,09
TRINITY_DN25715_c4_g1_i8	Importin subunit alpha-3	-8,09
TRINITY_DN32079_c1_g2_i10	Cation-independent mannose-6-phosphate receptor	-8,09
TRINITY_DN25654_c7_g2_i1	Phosphatidylinositol-binding clathrin assembly protein	-8,09
TRINITY_DN28386_c0_g1_i3	LINE-1 retrotransposable element ORF2 protein	-8,04
TRINITY_DN25115_c0_g2_i3	MARVEL domain-containing protein 2	-8,04
TRINITY_DN32311_c1_g1_i10	Histone deacetylase 4	-8,04
TRINITY_DN30254_c0_g1_i2	Synaptoporin	-8,04
TRINITY_DN26828_c0_g3_i1	Barrier-to-autointegration factor	-8,04
TRINITY_DN31022_c2_g1_i41	Glucuronokinase 1	-8,04
TRINITY_DN29338_c2_g2_i4	Integrin alpha-4	-8,04
TRINITY_DN29104_c2_g1_i5	Casein kinase I isoform gamma-3	-8,04
TRINITY_DN30890_c1_g1_i4	Endoplasmic reticulum membrane sensor NFE2L1	-8,04
TRINITY_DN29536_c3_g1_i4	BTB/POZ domain-containing protein 2	-8,04
TRINITY_DN24839_c4_g2_i17	TIP41-like protein	-8,04
TRINITY_DN30828_c1_g1_i9	Alpha-catulin	-8,04
TRINITY_DN29787_c1_g1_i3	Tetraspanin-31	-8,04
TRINITY_DN30887_c3_g1_i5	Claudin domain-containing protein 1	-8,04
TRINITY_DN27209_c1_g2_i7	Cyclin-T1	-8,04
TRINITY_DN28321_c3_g1_i9	Mitochondrial peptide methionine sulfoxide reductase	-8,04
TRINITY_DN27631_c0_g1_i3	N-acylethanolamine-hydrolyzing acid amidase	-8,04
_	Histone-lysine N-methyltransferase 2A	-8,04 -8,04
TRINITY_DN32777_c0_g1_i28		
TRINITY_DN26148_c1_g1_i16	LIM domain-binding protein 1	-8,04
TRINITY_DN26956_c3_g1_i1	Transcription elongation factor SPT5	-8,04
TRINITY_DN32311_c1_g1_i5	Histone deacetylase 4	-8,04
TRINITY_DN33190_c2_g1_i35	RNA-directed DNA polymerase from mobile element jockey	-8,04
TRINITY_DN31630_c1_g1_i7	Integrin alpha-9	-8,04
TRINITY_DN24290_c0_g1_i1	GDP-fucose protein O-fucosyltransferase 1	-8,04
TRINITY_DN24473_c0_g1_i2	Integrator complex subunit 2	-8,04
TRINITY_DN31880_c2_g1_i23	Growth factor receptor-bound protein 10	-8,04
TRINITY_DN27318_c1_g1_i10	Leucine-rich repeat-containing protein 57	-8,04
TRINITY_DN30759_c0_g2_i4	Malate synthase 1, glyoxysomal	-8,04
TRINITY_DN28499_c1_g1_i4	ETS translocation variant 4	-8,04
TRINITY_DN29117_c1_g2_i8	Cyclin-K	-8,04
TRINITY_DN27843_c0_g1_i5	Protein PRR14L	-8,04
TRINITY_DN28950_c0_g2_i5	Cyclin-C	-8,04
TRINITY_DN27576_c0_g1_i3	Histone-lysine N-methyltransferase ASH1L	-8,04
TRINITY_DN29353_c0_g1_i3	Signal peptide peptidase-like 2A	-8,04
TRINITY_DN25654_c6_g1_i12	Ceruloplasmin	-8,04
TRINITY_DN32977_c0_g1_i4	Inositol polyphosphate 5-phosphatase OCRL-1	-8,04
TRINITY_DN25926_c2_g2_i9	NLR family CARD domain-containing protein 3	-8,04
TRINITY_DN30037_c0_g1_i5	Engulfment and cell motility protein 1	-8,04
TRINITY_DN30557_c5_g1_i10	Solute carrier family 35 member E2	-8,04
TRINITY_DN32017_c2_g1_i5	E3 ubiquitin-protein ligase TRIM39	-8,04
TRINITY_DN28526_c1_g1_i10	Rab5 GDP/GTP exchange factor	-8,04
TRINITY_DN23491_c0_g1_i4	Epidermal growth factor receptor kinase substrate 8-like protein	-8,04
TRINITY_DN24582_c0_g1_i1	Aromatic-L-amino-acid decarboxylase	-8,04
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TRINITY_DN30844_c2_g1_i9	40S ribosomal protein S6	
TRINITY_DN31333_c0_g1_i4	Centrosomal protein of 295 kDa	-8,04
TRINITY_DN27125_c1_g1_i3	Myosin-11	-8,00
TRINITY_DN28199_c1_g1_i5	Epsin-1	-8,00
TRINITY_DN32144_c1_g1_i9	Fermitin family homolog 2	-8,00
TRINITY_DN28750_c3_g1_i11	CD276 antigen	-8,00
TRINITY_DN29961_c1_g3_i14	Acyl-coenzyme A thioesterase 4	-8,00
TRINITY_DN32903_c1_g2_i6	UPF0183 protein C16orf70	-8,00
TRINITY_DN33331_c1_g1_i5	Tryptophan 2,3-dioxygenase B	-8,00
TRINITY_DN33192_c2_g1_i4	Transcription factor 20	-8,00

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TRINITY_DN20691_c0_g1_i2	Sialoadhesin	-8,00
TRINITY_DN30598_c2_g1_i3	Homer protein homolog 2	-8,00
TRINITY_DN31547_c5_g3_i3	EH domain-containing protein 3	-8,00
TRINITY_DN22791_c0_g1_i3	FGFR1 oncogene partner 2 homolog	-8,00
TRINITY_DN26449_c1_g1_i6	CCR4-NOT transcription complex subunit 4	-8,00
TRINITY_DN31368_c1_g3_i2	MAP/microtubule affinity-regulating kinase 3	-8,00
TRINITY_DN31321_c0_g1_i2	[F-actin]-monooxygenase MICAL2	-8,00
TRINITY_DN29098_c1_g1_i9	WD repeat-containing protein 37	-8,00
TRINITY_DN30722_c1_g1_i12	Receptor-type tyrosine-protein phosphatase gamma	-8,00
TRINITY_DN23873_c0_g1_i3	Tomoregulin-2	-8,00
TRINITY_DN27007_c5_g1_i6	Limb region 1 homolog-like protein	-8,00
TRINITY_DN25163_c0_g1_i4	DNA polymerase kappa	-8,00
TRINITY_DN28415_c0_g1_i9	Small leucine-rich protein 1	-8,00
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TRINITY_DN25778_c0_g4_i7	Inositol-pentakisphosphate 2-kinase	-8,00
TRINITY_DN29757_c1_g1_i8	Sorbin and SH3 domain-containing protein 2	-8,00
TRINITY_DN30540_c2_g1_i4	Vitamin K-dependent protein C	-8,00
TRINITY_DN24603_c0_g3_i21	Zinc finger protein 430	-8,00
TRINITY_DN31834_c0_g1_i4	Interleukin-10 receptor subunit beta	-8,00
TRINITY_DN28915_c2_g1_i11	Kynurenine formamidase	-8,00
TRINITY_DN30792_c1_g1_i7	Transcription factor SOX-4	-8,00
TRINITY_DN30484_c0_g2_i4	Scavenger receptor class B member 1	-8,00
TRINITY_DN24207_c1_g1_i10	Nucleolar transcription factor 1-B	-8,00
TRINITY_DN29249_c0_g1_i8	CD276 antigen	-8,00
TRINITY_DN29171_c2_g1_i5	U2 small nuclear ribonucleoprotein A'	-8,00
TRINITY_DN30638_c2_g1_i13	Receptor tyrosine-protein kinase erbB-3	-8,00
TRINITY_DN24701_c0_g1_i2	Ubiquitin carboxyl-terminal hydrolase MINDY-2	-8,00
TRINITY_DN22937_c0_g1_i4	Interleukin-17 receptor E	-8,00
TRINITY_DN29999_c0_g2_i15	RNA-binding protein 26	-8,00
TRINITY_DN30432_c1_g2_i12	Ovostatin homolog	-8,00
TRINITY_DN27980_c0_g1_i18	DNA polymerase beta	-8,00
TRINITY_DN30323_c2_g2_i2	DNA damage-binding protein 1	-8,00
TRINITY_DN30018_c0_g1_i18	Zinc finger E-box-binding homeobox 2	-8,00
TRINITY_DN31079_c0_g1_i14	Stimulated by retinoic acid gene 6 protein-like	-8,00
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TRINITY_DN31000_c3_g1_i11	Drebrin-like protein	-8,00
TRINITY_DN25270_c1_g1_i9	Dynactin subunit 4	-8,00
TRINITY_DN26102_c2_g1_i3	Leucine-zipper-like transcriptional regulator 1	-8,00
TRINITY_DN32990_c1_g2_i6	SPRY domain-containing SOCS box protein 3	-8,00
TRINITY_DN28638_c0_g1_i13	R3H domain-containing protein 2	-8,00
TRINITY_DN27481_c2_g2_i6	DNA repair protein SWI5 homolog	-8,00
TRINITY_DN25986_c2_g3_i1	Retinal-specific ATP-binding cassette transporter	-8,00
TRINITY_DN29880_c1_g1_i4	TIMELESS-interacting protein	-8,00
TRINITY_DN25162_c0_g1_i10	Geranylgeranyl transferase type-1 subunit beta	-8,00
TRINITY_DN23110_c3_g5_i6	Pappalysin-2	-8,00
TRINITY_DN28826_c3_g2_i4	Pre-mRNA-processing factor 39	-7,95
TRINITY_DN29501_c0_g1_i20	Interferon-induced protein 44-like	-7,95
TRINITY_DN31501_c1_g1_i14	WAS/WASL-interacting protein family member 2	-7,95
TRINITY_DN32250_c0_g2_i4	Zinc finger protein 185	-7,95
TRINITY_DN29035_c0_g1_i1	A-kinase anchor protein 9	-7,95
TRINITY_DN28116_c1_g1_i4	Cingulin-like protein 1	-7,95
TRINITY_DN29757_c1_g1_i17	Sorbin and SH3 domain-containing protein 2	-7,95
TRINITY_DN31058_c0_g2_i2	Moesin	-7,95
TRINITY_DN27103_c0_g1_i3	Reticulon-4-interacting protein 1 homolog, mitochondrial	-7,95
TRINITY_DN31694_c0_g3_i2	Histone acetyltransferase p300	-7,95 -7,95
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TRINITY_DN29127_c4_g2_i11	WD repeat domain phosphoinositide-interacting protein 2	-7,95
TRINITY_DN29467_c3_g1_i5	Protein FAM78A	-7,95
TRINITY_DN31507_c1_g1_i5	Multidrug resistance-associated protein 5	-7,95
TRINITY_DN23241_c3_g1_i1	Cytoplasmic protein NCK2	-7,95
TRINITY_DN32459_c1_g1_i5	Phosphatidylinositol N-acetylglucosaminyltransferase subunit I	-7,95
TRINITY_DN30038_c2_g2_i2	Transcription factor HIVEP2	-7,95

TRINITY_DN30107_c0_g1_i6	Signal transducer and activator of transcription 1	-7,95
TRINITY_DN32424_c1_g1_i4	Pericentriolar material 1 protein	-7,95
TRINITY_DN31635_c1_g2_i1	Retinoic acid receptor RXR-gamma-B	-7,95
TRINITY_DN30714_c2_g1_i2	Very low-density lipoprotein receptor	-7,95
TRINITY_DN26968_c0_g1_i8	Type I inositol 3,4-bisphosphate 4-phosphatase	-7,95
TRINITY_DN32408_c0_g1_i8	Protein strawberry notch homolog 2	-7,95
TRINITY_DN31407_c1_g1_i3	Peroxisomal sarcosine oxidase	-7,95
TRINITY_DN27857_c0_g2_i2	Anaphase-promoting complex subunit 4	-7,95
TRINITY_DN28026_c1_g1_i3	Non-specific lipid-transfer protein (Fragment)	-7,95
TRINITY_DN24309_c0_g1_i8	GATOR complex protein NPRL2	-7,95
TRINITY_DN22313_c0_g1_i4	Asialoglycoprotein receptor 1	-7,95
TRINITY_DN32582_c0_g1_i12	RING finger protein 10	-7,95
TRINITY_DN33061_c1_g1_i8	Low-density lipoprotein receptor-related protein 12 (Fragment)	-7,95
TRINITY_DN25020_c0_g1_i8	Transcription factor HIVEP2	-7,95
TRINITY_DN25032_c1_g1_i5	Activating molecule in BECN1-regulated autophagy protein 1	-7,95
TRINITY_DN31875_c0_g2_i1	Retrovirus-related Pol polyprotein from transposon 17.6	-7,95
TRINITY_DN26550_c0_g1_i6	Complement factor B	-7,95
TRINITY_DN29005_c1_g3_i9	FYVE, RhoGEF and PH domain-containing protein 4	-7,95
TRINITY_DN26924_c3_g1_i9	Keratinocyte-associated protein 2	-7,95
TRINITY_DN23051_c0_g1_i5	N-lysine methyltransferase SMYD2-A	-7,95
TRINITY_DN25363_c0_g1_i2	ETS-related transcription factor Elf-4	-7,95
TRINITY_DN25743_c0_g1_i5	Interferon regulatory factor 1	-7,95
TRINITY_DN25781_c1_g2_i2	Casein kinase II subunit alpha'	-7,95 -7,95
		-7,95 -7,95
TRINITY_DN28475_c0_g2_i6	V-type proton ATPase 116 kDa subunit a isoform 2	
TRINITY_DN33141_c3_g1_i5	Uncharacterized protein KIAA1522 homolog	-7,95
TRINITY_DN25180_c5_g1_i9	Calcium-dependent secretion activator 2	-7,95
TRINITY_DN19608_c0_g1_i4	Putative UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminylt	-7,95
TRINITY_DN30108_c0_g1_i10	Septin-9	-7,95
TRINITY_DN32850_c3_g1_i7	Arrestin domain-containing protein 3	-7,95
TRINITY_DN28574_c0_g2_i1	Interferon-inducible GTPase 5	-7,95
TRINITY_DN28826_c3_g1_i2	Homeodomain-interacting protein kinase 1	-7,95
TRINITY_DN28352_c0_g1_i1	Espin-like protein	-7,95
TRINITY_DN21921_c0_g1_i1	Paladin	-7,95
TRINITY_DN28968_c1_g2_i1	AsparaginetRNA ligase, cytoplasmic	-7,95
TRINITY_DN28252_c2_g1_i4	Growth factor receptor-bound protein 10	-7,90
TRINITY_DN26313_c4_g1_i9	Tudor and KH domain-containing protein	-7,90
TRINITY_DN26858_c0_g1_i6	Calsyntenin-3	-7,90
TRINITY_DN31989_c1_g1_i8	Protein FAM208B	-7,90
TRINITY_DN25370_c0_g1_i7	Poly(U)-specific endoribonuclease- C	-7,90
TRINITY_DN31079_c0_g1_i8	Stimulated by retinoic acid gene 6 protein-like	-7,90
TRINITY_DN26144_c0_g1_i17	PTB domain-containing engulfment adapter protein 1	-7,90
TRINITY_DN33013_c1_g1_i3	Scavenger receptor class F member 1	-7,90
TRINITY_DN29104_c2_g1_i21	Casein kinase I isoform gamma-3	-7,90
TRINITY_DN27836_c0_g1_i1	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mi	-7,90
TRINITY_DN28034_c2_g1_i13	Guanine nucleotide exchange factor VAV3	-7,90
TRINITY_DN31579_c0_g2_i3	APOBEC1 complementation factor	-7,90
TRINITY_DN25138_c0_g1_i3	Ubiquitin-protein ligase E3C	-7,90
TRINITY_DN31830_c0_g3_i2	WW domain binding protein 1-like	-7,90
TRINITY_DN28611_c2_g2_i13	DnaJ homolog subfamily C member 5	-7,90
TRINITY_DN29161_c0_g1_i1	Cell growth regulator with EF hand domain protein 1	-7,90
TRINITY_DN26028_c0_g3_i3	Neurogenic locus notch homolog protein 1	-7,90
TRINITY_DN29843_c1_g1_i3	Neurabin-1	-7,90
TRINITY_DN30382_c0_g1_i1	Splicing factor 3B subunit 3	-7,90 -7,90
TRINITY_DN27236_c0_g2_i9	POU domain, class 2, transcription factor 1	-7,90 -7,90
TRINITY_DN29732_c0_g1_i10	Phosphatidate phosphatase LPIN2	-7,90 -7,90
TRINITY_DN23127_c0_g1_i3	Heat-stable enterotoxin receptor	-7,90 -7,90
TRINITY_DN31440_c0_g1_i8	Rab3 GTPase-activating protein non-catalytic subunit	-7,90 7,00
TRINITY_DN22865_c0_g1_i3	Eukaryotic translation initiation factor 4E	-7,90 7,00
TRINITY_DN25495_c0_g1_i6	Golgi-associated plant pathogenesis-related protein 1	-7,90

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TRINITY_DN21380_c0_g1_i1	Carnitine O-palmitoyltransferase 1, liver isoform	-7,90
TRINITY_DN26952_c4_g2_i4	Sterol-4-alpha-carboxylate 3-dehydrogenase, decarboxylating	-7,90
TRINITY_DN23355_c0_g1_i4	Proteasome subunit alpha type-2	-7,90
TRINITY_DN30202_c0_g1_i5	Disco-interacting protein 2 homolog B-A	-7,90
TRINITY_DN27496_c1_g1_i14	AH receptor-interacting protein	-7,90
TRINITY_DN32654_c0_g1_i1	GON-4-like protein	-7,90
TRINITY_DN23420_c0_g1_i4	Proliferation marker protein Ki-67	-7,90
TRINITY_DN27181_c1_g1_i1	Aquaporin-11	-7,90
TRINITY_DN28216_c1_g4_i5	Transmembrane 4 L6 family member 1	-7,90
TRINITY_DN28782_c0_g1_i4	H-2 class II histocompatibility antigen, E-K alpha chain	-7,90
TRINITY_DN30863_c1_g1_i1	Inversin	-7,90
TRINITY_DN32056_c0_g1_i3	Interferon-induced very large GTPase 1	-7,90
TRINITY_DN26923_c2_g1_i8	Krueppel-like factor 12	-7,90
TRINITY_DN25204_c2_g1_i28	Serine/threonine-protein kinase MARK2	-7,90
TRINITY_DN31079_c0_g1_i34	Stimulated by retinoic acid gene 6 protein-like	-7,90
TRINITY_DN33010_c1_g2_i3	Xanthine dehydrogenase/oxidase	-7,90
TRINITY_DN29173_c2_g3_i8	1-acylglycerol-3-phosphate O-acyltransferase ABHD5	-7,90
TRINITY_DN26659_c0_g2_i12	SMC5-SMC6 complex localization factor protein 2	-7,90
TRINITY_DN25818_c3_g1_i13	MAP7 domain-containing protein 1	-7,90
TRINITY_DN31172_c0_g1_i25	OTU domain-containing protein 3	-7,90 -7,90
TRINITY_DN27540_c4_g1_i8	Saccharopine dehydrogenase-like oxidoreductase	-7,90 -7,90
TRINITY_DN30321_c0_g1_i20	Nuclear transcription factor Y subunit gamma	-7,85
TRINITY_DN28408_c1_g1_i2	H(+)/Cl(-) exchange transporter 5	-7,85
TRINITY_DN31483_c0_g1_i5	Ras GTPase-activating protein 1	-7,85
TRINITY_DN22660_c0_g1_i2	E3 ubiquitin-protein ligase RNF31	-7,85
TRINITY_DN28419_c1_g4_i3	Myotrophin	-7,85
TRINITY_DN30315_c0_g1_i3	Glutathione S-transferase kappa 1	-7,85
TRINITY_DN22833_c0_g1_i2	Target of rapamycin complex subunit lst8	-7,85
TRINITY_DN29159_c2_g1_i1	Purine-rich element-binding protein gamma	-7,85
TRINITY_DN32725_c1_g3_i1	Actin-binding LIM protein 1	-7,85
TRINITY_DN30036_c0_g1_i4	CREB-binding protein	-7,85
TRINITY_DN22516_c0_g1_i7	Cathepsin S	-7,85
TRINITY_DN30921_c1_g2_i20	Unconventional myosin-X	-7,85
TRINITY_DN30905_c1_g1_i7	Serinepyruvate aminotransferase, mitochondrial	-7,85
TRINITY_DN30247_c3_g1_i3	Tyrosine 3-monooxygenase	-7,85
TRINITY_DN25987_c2_g3_i1	Tetratricopeptide repeat protein 39B	-7,85
TRINITY_DN29355_c0_g1_i3	Partner and localizer of BRCA2	-7,85
TRINITY_DN33254_c8_g1_i6	Ceruloplasmin	-7,85
TRINITY_DN23684_c0_g1_i6	Plexin-B1	-7,85
TRINITY_DN29698_c1_g1_i9	H-2 class II histocompatibility antigen, I-E beta chain	-7,85
TRINITY_DN33317_c6_g1_i8	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	-7,85
TRINITY_DN26987_c0_g1_i1	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	-7,85
TRINITY_DN27877_c0_g1_i2	Histone H4 transcription factor	-7,85
TRINITY_DN31830_c0_g3_i5	WW domain binding protein 1-like	-7,85
TRINITY_DN30717_c0_g3_i4	Serine/threonine-protein kinase 10	-7,85 -7,85
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TRINITY_DN28950_c0_g2_i4	Cyclin-C	-7,85
TRINITY_DN30702_c0_g1_i4	Glutaminase kidney isoform, mitochondrial	-7,85
TRINITY_DN31073_c1_g2_i10	85/88 kDa calcium-independent phospholipase A2	-7,85
TRINITY_DN30473_c3_g1_i11	Ecotropic viral integration site 5 protein homolog	-7,85
TRINITY_DN32623_c6_g1_i13	Atrial natriuretic peptide receptor 2	-7,85
TRINITY_DN28284_c1_g1_i13	Protein jagged-1b	-7,85
TRINITY_DN22829_c0_g1_i2	TNF receptor-associated factor 4	-7,85
TRINITY_DN30303_c1_g2_i1	Diphosphoinositol polyphosphate phosphohydrolase 1	-7,85
TRINITY_DN30526_c0_g1_i1	Diacylglycerol kinase zeta	-7,85
TRINITY_DN31624_c0_g1_i2	Mitochondrial dynamics protein MID51	-7,85
TRINITY_DN28302_c1_g1_i5	General transcription factor IIH subunit 1	-7,85
TRINITY_DN31101_c0_g1_i9	Natural resistance-associated macrophage protein 2	-7,85
TRINITY_DN33318_c5_g2_i4	Methylenetetrahydrofolate reductase	-7,85
TRINITY_DN25554_c0_g1_i2	Serine/threonine-protein phosphatase PP1-gamma catalytic suk	-7,85

TDINITY DN26221 -0 -1 :7	D l l	7.05
TRINITY_DN26231_c0_g1_i7	Probable carboxypeptidase X1	-7,85
TRINITY_DN24772_c4_g3_i8	GPI mannosyltransferase 1	-7,85
TRINITY_DN31066_c0_g1_i5	Spermatogenesis-associated protein 13	-7,85
TRINITY_DN24571_c3_g2_i3	Tyrosine-protein kinase FRK	-7,85
TRINITY_DN32777_c0_g1_i14	Histone-lysine N-methyltransferase 2A	-7,85
TRINITY_DN25683_c4_g1_i6	EF-hand calcium-binding domain-containing protein 6	-7,85
TRINITY_DN30643_c1_g1_i18	Protein lifeguard 1	-7,85
TRINITY_DN30549_c0_g1_i2	Transmembrane protein 245	-7,85
TRINITY_DN21586_c0_g1_i2	Cytochrome c oxidase assembly factor 4 homolog, mitochondric	-7,85
TRINITY_DN27692_c1_g3_i2	Ras GTPase-activating protein nGAP	-7,85
TRINITY_DN24194_c0_g1_i1	40S ribosomal protein S9	-7,85
TRINITY_DN32263_c0_g1_i9	Dual specificity tyrosine-phosphorylation-regulated kinase 1A	-7,85
TRINITY_DN26624_c2_g2_i2	Cytosolic purine 5'-nucleotidase	-7,85
TRINITY_DN30388_c8_g1_i4	Probable RNA-directed DNA polymerase from transposon X-ele	-7,85
TRINITY_DN25337_c0_g1_i6	Ral guanine nucleotide dissociation stimulator	-7,85
TRINITY_DN23316_c0_g1_i4	Serine/threonine-protein kinase 19	-7,85
TRINITY_DN30460_c2_g1_i13	Ras association domain-containing protein 6	-7,85
TRINITY_DN32665_c0_g1_i6	Immunoglobulin superfamily member 8	-7,85
TRINITY_DN26977_c4_g1_i5	Far upstream element-binding protein 1	-7,80
TRINITY_DN31980_c1_g1_i7	Cytochrome P450 2J2	-7,80
TRINITY_DN29068_c1_g1_i6	Kelch-like protein 8	-7,80
TRINITY_DN26574_c2_g1_i3	Macoilin-2	-7,80
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TRINITY_DN24598_c0_g1_i1	ATPase family AAA domain-containing protein 2B	-7,80 7,80
TRINITY_DN27403_c0_g1_i7	Long-chain-fatty-acidCoA ligase ACSBG1	-7,80
TRINITY_DN32919_c4_g2_i3	Glutaminase liver isoform, mitochondrial	-7,80 7 ,80
TRINITY_DN31637_c0_g1_i2	Neurofibromin	-7,80
TRINITY_DN33063_c4_g3_i5	Bone morphogenetic protein 1	-7,80
TRINITY_DN32785_c3_g1_i8	Heme-binding protein 2	-7,80
TRINITY_DN29685_c1_g3_i7	Cip1-interacting zinc finger protein	-7,80
TRINITY_DN30304_c1_g1_i3	Kynurenine 3-monooxygenase	-7,80
TRINITY_DN25474_c0_g2_i6	Sentrin-specific protease 1	-7,80
TRINITY_DN32096_c0_g2_i11	Gap junction beta-6 protein	-7,80
TRINITY_DN30833_c2_g1_i7	Stathmin-2	-7,80
TRINITY_DN26147_c1_g3_i6	Pyridoxal phosphate phosphatase	-7,80
TRINITY_DN24856_c0_g1_i4	Mesoderm induction early response protein 1	-7,80
TRINITY_DN28191_c0_g4_i1	RAC-beta serine/threonine-protein kinase	-7,80
TRINITY_DN25427_c2_g1_i5	TBC1 domain family member 25	-7,80
TRINITY_DN30168_c1_g2_i4	Death-associated protein kinase 1	-7,80
TRINITY_DN28464_c1_g2_i6	Amino-terminal enhancer of split	-7,80
TRINITY_DN30891_c2_g2_i9	Thrombospondin-1	-7,80
TRINITY_DN27467_c4_g3_i10	Mitochondrial sodium/calcium exchanger protein	-7,80
TRINITY_DN24556_c8_g1_i2	General transcription factor II-I repeat domain-containing prot	-7,80
TRINITY_DN31673_c0_g1_i13	Major histocompatibility complex class I-related gene protein	-7,80
_	Tensin-1	
TRINITY_DN31493_c2_g1_i8		-7,80
TRINITY_DN25303_c0_g1_i10	Hippocalcin-like protein 1	-7,80
TRINITY_DN28454_c2_g2_i8	Phospholipase D3	-7,80
TRINITY_DN23223_c2_g1_i1	Protein ENL	-7,80
TRINITY_DN31606_c0_g1_i2	Endoplasmin (Fragment)	-7,80
TRINITY_DN28880_c0_g1_i10	Insulin-induced gene 2 protein	-7,80
TRINITY_DN27484_c0_g1_i9	BTB/POZ domain-containing protein KCTD2	-7,80
TRINITY_DN29819_c3_g1_i16	Leucine-rich repeat and calponin homology domain-containing	-7,80
TRINITY_DN27581_c0_g1_i11	Adiponectin receptor protein 1	-7,80
TRINITY_DN23125_c0_g1_i2	Selenoprotein M	-7,80
TRINITY_DN27840_c0_g1_i2	Cell division cycle protein 27 homolog	-7,80
TRINITY_DN22972_c0_g1_i1	NF-kappa-B inhibitor-interacting Ras-like protein 1	-7,80
TRINITY_DN32269_c3_g2_i3	Myosin light chain kinase family member 4	-7,80
TRINITY_DN29385_c2_g1_i5	Complement component C6	-7,80
TRINITY_DN25350_c2_g1_i8	Mitochondrial intermembrane space import and assembly prote	-7,80
TRINITY_DN32777_c0_g1_i8	Histone-lysine N-methyltransferase 2A	-7,80
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TDINITY DN27126 -2 -1 :1	A domilato binaco iso comuno 1	-7,80
TRINITY_DN27126_c2_g1_i1	Adenylate kinase isoenzyme 1 Glucuronokinase 1	
TRINITY_DN31022_c2_g1_i18		-7,80
TRINITY_DN28356_c0_g2_i14	Ribosome biogenesis protein BRX1 homolog	-7,80
TRINITY_DN27409_c5_g1_i7	CD99 antigen-like protein 2	-7,80
TRINITY_DN31272_c0_g1_i8	Ubiquitin-protein ligase E3B	-7,80
TRINITY_DN29699_c0_g1_i10	A-kinase anchor protein 8-like	-7,80
TRINITY_DN30626_c1_g1_i5	CD9 antigen	-7,80
TRINITY_DN31986_c0_g1_i8	Liprin-beta-1	-7,80
TRINITY_DN27189_c1_g1_i2	Lysine-specific demethylase 5A	-7,80
TRINITY_DN30411_c1_g1_i14	Transposon Ty3-I Gag-Pol polyprotein	-7,80
TRINITY_DN25072_c0_g1_i4	Ankyrin-1	-7,80
TRINITY_DN32463_c0_g1_i3	Replication stress response regulator SDE2	-7,80
TRINITY_DN30179_c0_g1_i3	Epoxide hydrolase 1	-7,80
TRINITY_DN28603_c0_g1_i1	NLR family CARD domain-containing protein 3	-7,80
TRINITY_DN31945_c1_g1_i4	Protein diaphanous homolog 2	-7,80
TRINITY_DN26848_c0_g1_i4	Peroxisome proliferator-activated receptor delta	-7,80
TRINITY_DN30840_c0_g2_i9	Ubiquitin carboxyl-terminal hydrolase 3	-7,80
TRINITY_DN28401_c0_g2_i7	Unconventional myosin-Ie	-7,80
TRINITY_DN24055_c0_g1_i2	5'-3' exoribonuclease 2	-7,80
TRINITY_DN27207_c1_g1_i5	Interferon-induced helicase C domain-containing protein 1	-7,80
TRINITY_DN28658_c0_g1_i3	Insulin-induced gene 1 protein	-7,80
TRINITY_DN31702_c1_g1_i7	Uveal autoantigen with coiled-coil domains and ankyrin repeat:	-7,80
TRINITY_DN29887_c3_g1_i13	ERO1-like protein beta	-7,80 -7,79
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TRINITY_DN26551_c0_g1_i3	Ceramide kinase	-7,75
TRINITY_DN30391_c0_g1_i1	Inositol 1,4,5-trisphosphate receptor type 2	-7,75
TRINITY_DN28095_c1_g1_i2	ORM1-like protein 1	-7,75
TRINITY_DN22234_c0_g1_i1	Helicase with zinc finger domain 2	-7,75
TRINITY_DN28658_c0_g1_i7	Insulin-induced gene 1 protein	-7,75
TRINITY_DN32574_c4_g1_i3	PAX3- and PAX7-binding protein 1	-7,75
TRINITY_DN32096_c0_g2_i12	Gap junction beta-6 protein	-7,75
TRINITY_DN30272_c0_g2_i8	Retinoblastoma-binding protein 5	-7,75
TRINITY_DN25016_c0_g2_i4	Type II inositol 1,4,5-trisphosphate 5-phosphatase	-7,75
TRINITY_DN29536_c3_g1_i14	BTB/POZ domain-containing protein 2	-7,75
TRINITY_DN22978_c0_g1_i4	MIT domain-containing protein 1	-7,75
TRINITY_DN26231_c0_g1_i6	Probable carboxypeptidase X1	-7,75
TRINITY_DN31374_c0_g2_i1	2-oxoglutarate dehydrogenase, mitochondrial	-7,75
TRINITY_DN23591_c0_g1_i3	Ubl carboxyl-terminal hydrolase 18	-7,75
TRINITY_DN27350_c3_g2_i15	GREB1-like protein	-7,75
TRINITY_DN31752_c0_g1_i4	Dynactin subunit 1	-7,75
TRINITY_DN30019_c0_g1_i7	Polypyrimidine tract-binding protein 1	-7,75
TRINITY_DN31986_c0_g1_i16	Liprin-beta-1	-7,75
TRINITY_DN24611_c1_g1_i2	Zinc finger protein 827	-7,75
TRINITY_DN28082_c0_g3_i8	Hemicentin-1	-7,75
TRINITY_DN27189_c0_g1_i6	5'-AMP-activated protein kinase subunit gamma-2	-7,75
TRINITY_DN26655_c0_g2_i5	Growth hormone-inducible transmembrane protein	-7,75
TRINITY_DN27424_c0_g3_i8	Apoptosis regulator BAX	-7,75
TRINITY_DN24858_c0_g1_i18		-7,75 -7,75
	Ninjurin-1	
TRINITY_DN27733_c0_g1_i4	Mitochondrial import inner membrane translocase subunit Tim.	-7,75
TRINITY_DN31804_c0_g1_i4	Golgi integral membrane protein 4	-7,75
TRINITY_DN26549_c5_g1_i1	Polyubiquitin 12	-7,75
TRINITY_DN29201_c0_g2_i5	Purine nucleoside phosphorylase	-7,75
TRINITY_DN23274_c0_g1_i2	Glucose-6-phosphate exchanger SLC37A2	-7,75
TRINITY_DN29065_c0_g1_i6	Phosphatidylcholine-sterol acyltransferase (Fragment)	-7,75
TRINITY_DN30359_c3_g1_i4	RelA-associated inhibitor	-7,75
TRINITY_DN27545_c0_g1_i11	DNA excision repair protein ERCC-8	-7,75
TRINITY_DN30428_c2_g1_i19	Speckle-type POZ protein-like B	-7,75
TRINITY_DN29178_c2_g2_i4	Ataxin-7-like protein 1	-7,75
TRINITY_DN29050_c0_g2_i6	SAM domain-containing protein SAMSN-1	-7,75
TRINITY_DN23355_c0_g1_i1	Proteasome subunit alpha type-2	-7,75

TRINITY_DN31924_c1_g1_i4	Ribose-phosphate pyrophosphokinase 1	-7,75
TRINITY_DN32712_c2_g1_i8	Elongation factor 1-alpha 2	-7,75
TRINITY_DN26696_c1_g2_i4	SLIT-ROBO Rho GTPase-activating protein 3	-7,75
TRINITY_DN33175_c2_g1_i10	Golgi-specific brefeldin A-resistance guanine nucleotide exchan	-7,75
TRINITY_DN28675_c1_g1_i11	Protein FAM117B	-7,75
TRINITY_DN29791_c0_g1_i5	Mediator of RNA polymerase II transcription subunit 13-like	-7,75
TRINITY_DN25702_c3_g1_i1	Telomerase RNA component interacting RNase	-7,75
TRINITY_DN30959_c0_g1_i10	Uncharacterized protein KIAA1671	-7,75
TRINITY_DN27137_c0_g1_i3	DnaJ homolog subfamily B member 1	-7,75
TRINITY_DN26469_c1_g3_i8	Solute carrier organic anion transporter family member 2A1	-7,75
TRINITY_DN32709_c0_g1_i2	RNA-binding protein NOB1	-7,75
TRINITY_DN31294_c0_g1_i19	Cyclin-L1	-7,75
TRINITY_DN29691_c0_g1_i8	·	-7,75 -7,75
	Peripheral plasma membrane protein CASK	
TRINITY_DN28981_c0_g1_i6	Protein CASP	-7,75
TRINITY_DN33155_c2_g1_i8	Probable E3 ubiquitin-protein ligase HERC3	-7,75
TRINITY_DN30988_c6_g1_i3	AF4/FMR2 family member 4	-7,75
TRINITY_DN28333_c1_g1_i11	Patatin-like phospholipase domain-containing protein 2	-7,75
TRINITY_DN25273_c0_g1_i4	Nuclear protein localization protein 4 homolog	-7,75
TRINITY_DN26547_c1_g3_i2	Cell cycle control protein 50A	-7,75
TRINITY_DN28394_c0_g1_i1	RNA 3'-terminal phosphate cyclase	-7,75
TRINITY_DN23679_c7_g2_i10	NXPE family member 3	-7,75
TRINITY_DN25632_c0_g1_i11	Growth arrest and DNA damage-inducible protein GADD45 be	-7,75
TRINITY_DN23920_c0_g1_i1	Retrovirus-related Pol polyprotein from transposon TNT 1-94	-7,75
TRINITY_DN31636_c1_g1_i1	Upstream-binding protein 1	-7,75
TRINITY_DN32816_c0_g1_i13	Regulator of microtubule dynamics protein 1	-7,75
TRINITY_DN30583_c0_g1_i3	Rho GTPase-activating protein 45	-7,75
TRINITY_DN28519_c0_g1_i13	Serine/threonine-protein phosphatase 2A 56 kDa regulatory sul	-7,75
TRINITY_DN28515_c2_g1_i9	Intersectin-1	-7,75
TRINITY_DN29764_c1_g3_i7	ATP-binding cassette sub-family G member 2	-7,75
TRINITY_DN27958_c0_g1_i1	LYR motif-containing protein 5B	-7,75
TRINITY_DN28135_c0_g1_i1	Cell surface hyaluronidase	-7,75
TRINITY_DN26481_c3_g1_i31	Prostaglandin-H2 D-isomerase	-7,75 -7,75
TRINITY_DN31257_c0_g1_i3	Poly(A)-specific ribonuclease PARN	
	* ' ' * * *	-7,75
TRINITY_DN29936_c0_g5_i1	Transposon TX1 uncharacterized 82 kDa protein Plexin-B1	-7,75
TRINITY_DN32666_c1_g1_i5		-7,69
TRINITY_DN30173_c1_g1_i9	60S ribosomal protein L3	-7,69
TRINITY_DN28649_c0_g1_i22	Poly(U)-binding-splicing factor PUF60	-7,69
TRINITY_DN32206_c0_g1_i5	Molybdenum cofactor sulfurase	-7,69
TRINITY_DN23157_c0_g1_i8	Contactin-associated protein-like 4	-7,69
TRINITY_DN30773_c2_g2_i10	Syntaxin-1A	-7,69
TRINITY_DN32736_c0_g1_i9	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	-7,69
TRINITY_DN31523_c0_g1_i7	Phosphorylase b kinase regulatory subunit alpha, skeletal musc	-7,69
TRINITY_DN29969_c1_g1_i10	WD repeat domain phosphoinositide-interacting protein 1	-7,69
TRINITY_DN30890_c1_g1_i10	Endoplasmic reticulum membrane sensor NFE2L1	-7,69
TRINITY_DN26664_c0_g2_i3	28S ribosomal protein S36, mitochondrial	-7,69
TRINITY_DN33158_c0_g1_i2	Midasin	-7,69
TRINITY_DN33124_c1_g1_i2	Plectin (Fragment)	-7,69
TRINITY_DN27900_c1_g1_i1	Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial	-7,69
TRINITY_DN26954_c4_g1_i14	RUS1 family protein C16orf58 homolog	-7,69
TRINITY_DN26222_c1_g1_i7	Dynamin-1-like protein	-7,69
TRINITY_DN23143_c0_g1_i1	GPI transamidase component PIG-T	-7,69
TRINITY_DN27019_c1_g2_i2		
	Transmembrane protein 14C	-7,69
TRINITY_DN32249_c0_g1_i8	Uncharacterized protein KIAA2013 homolog	-7,69
TRINITY_DN30086_c0_g2_i1	Protein sprouty homolog 2	-7,69
TRINITY_DN29490_c0_g2_i5	Methyl-CpG-binding domain protein 5	-7,69
TRINITY_DN31674_c0_g2_i11	Solute carrier family 12 member 2	-7,69
TRINITY_DN22756_c0_g1_i1	Protein NipSnap homolog 1	-7,69
TRINITY_DN30572_c0_g1_i5	Lysosomal alpha-glucosidase	-7,69
TRINITY_DN24394_c0_g1_i5	Mitochondrial fission process protein 1	-7,69

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TRINITY_DN27839_c2_g1_i6	Short coiled-coil protein A	-7,69
TRINITY_DN24890_c3_g1_i7	Beta-microseminoprotein	-7,69
TRINITY_DN31651_c0_g1_i8	Zinc finger and BTB domain-containing protein 10	-7,69
TRINITY_DN25175_c1_g3_i9	Ataxin-7-like protein 2	-7,69
TRINITY_DN32174_c0_g1_i3	Probable E3 ubiquitin-protein ligase HERC3	-7,69
TRINITY_DN28893_c1_g1_i6	Phospholipase A2	-7,69
TRINITY_DN29611_c0_g1_i11	Ras-related protein Rab-24	-7,69
TRINITY_DN23813_c0_g2_i3	Leucine-rich repeat transmembrane protein FLRT3	-7,69
TRINITY_DN29867_c3_g1_i9	Protein rapunzel	-7,69
TRINITY_DN32965_c0_g1_i26	·	
	N-lysine methyltransferase setd6	-7,69
TRINITY_DN27802_c0_g1_i8	Suppression of tumorigenicity 5 protein	-7,69
TRINITY_DN32450_c0_g1_i14	Lysine-specific demethylase 2B	-7,69
TRINITY_DN29914_c0_g4_i8	Uncharacterized protein KIAA1211-like	-7,69
TRINITY_DN29686_c0_g1_i4	Ubiquitin carboxyl-terminal hydrolase 19	-7,69
TRINITY_DN23841_c0_g1_i6	E3 ubiquitin-protein ligase Topors	-7,69
TRINITY_DN29718_c0_g1_i1	Zinc finger protein 280D	-7,69
TRINITY_DN26335_c0_g2_i8	Serine/threonine-protein kinase H1 homolog	-7,69
TRINITY_DN25994_c3_g1_i2	Protein FAM126B	-7,69
TRINITY_DN32979_c2_g3_i12	A-kinase anchor protein 7 isoform alpha	-7,69
TRINITY_DN20854_c0_g1_i1	THAP domain-containing protein 3	-7,69
TRINITY_DN25348_c0_g1_i3	Protein asteroid homolog 1	-7,69
TRINITY_DN25140_c0_g1_i10	Cytoplasmic protein NCK1	-7,69
TRINITY_DN24776_c0_g1_i4	RNA-binding protein 45	-7,69
TRINITY_DN33058_c1_g1_i19	H-2 class II histocompatibility antigen, E-D beta chain	-7,69
TRINITY_DN26669_c0_g1_i5	Protein-glutamine gamma-glutamyltransferase K	-7,69
TRINITY_DN23628_c0_g1_i4	rRNA methyltransferase 1, mitochondrial	-7,69
TRINITY_DN28821_c2_g2_i5	Unconventional myosin-VI	-7,69
TRINITY_DN28554_c4_g1_i8	Kinesin-associated protein 3	-7,69
TRINITY_DN31693_c0_g1_i3	Splicing factor, suppressor of white-apricot homolog	-7,69
TRINITY_DN28236_c0_g1_i9	Arrestin domain-containing protein 2	-7,69
TRINITY_DN25285_c3_g1_i3	S-acyl fatty acid synthase thioesterase, medium chain	-7,69
TRINITY_DN25039_c0_g1_i4		
=	Elongation of very long chain fatty acids protein 4	-7,69
TRINITY_DN23933_c2_g2_i4	Zinc finger protein 318	-7,69
TRINITY_DN26099_c0_g2_i1	Proline-rich protein PRCC	-7,69
TRINITY_DN31924_c1_g1_i27	Ribose-phosphate pyrophosphokinase 1	-7,69
TRINITY_DN25889_c3_g1_i7	Septin-6	-7,69
TRINITY_DN30828_c1_g1_i11	Alpha-catulin	-7,69
TRINITY_DN22056_c0_g1_i4	Glycoprotein hormone beta-5	-7,69
TRINITY_DN31819_c0_g1_i10	UNC93-like protein MFSD11	-7,69
TRINITY_DN26309_c1_g1_i10	Protein quaking-B	-7,69
TRINITY_DN25461_c3_g2_i1	Polyadenylate-binding protein-interacting protein 2B	-7,69
TRINITY_DN30751_c0_g2_i8	F-box/LRR-repeat protein 20	-7,69
TRINITY_DN28238_c0_g2_i3	Importin-7	-7,69
	Cyclin-T1	
TRINITY_DN27209_c1_g2_i18	· ·	-7,69
TRINITY_DN31297_c0_g1_i8	TBC1 domain family member 10B	-7,69
TRINITY_DN29313_c4_g3_i12	Zinc finger protein ubi-d4	-7,69
TRINITY_DN22783_c0_g1_i1	Regulator of G-protein signaling 8	-7,69
TRINITY_DN30792_c1_g1_i9	Transcription factor SOX-4	-7,69
TRINITY_DN29104_c2_g1_i13	Casein kinase I isoform gamma-2	-7,69
TRINITY_DN25394_c0_g1_i9	Class I histocompatibility antigen, F10 alpha chain	-7,63
TRINITY_DN30591_c0_g2_i13	Zinc finger protein 609	-7,63
TRINITY_DN25394_c0_g1_i16	Class I histocompatibility antigen, F10 alpha chain	-7,63
TRINITY_DN27581_c0_g1_i10	Adiponectin receptor protein 1	-7,63
TRINITY_DN32584_c0_g2_i2	Transport and Golgi organization protein 1 homolog	-7,63
TRINITY_DN27034_c1_g2_i15	Sodium-dependent multivitamin transporter	-7,63
TRINITY_DN24188_c0_g1_i5	26S proteasome non-ATPase regulatory subunit 6	-7,63 -7,63
TRINITY_DN31778_c0_g1_i3	Patatin-like phospholipase domain-containing protein 7	-7,63
TRINITY_DN28607_c2_g1_i15	CSC1-like protein 1	-7,63
TRINITY_DN29292_c0_g2_i1	Chemokine-like receptor 1	-7,63

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TRINITY_DN23976_c0_g1_i14	Immunoglobulin kappa light chain	-7,63
TRINITY_DN32267_c4_g1_i20	NACHT, LRR and PYD domains-containing protein 5	-7,63
TRINITY_DN31431_c2_g1_i7	Bromodomain adjacent to zinc finger domain protein 2A	-7,63
TRINITY_DN29726_c3_g2_i6	Aldehyde dehydrogenase, mitochondrial	-7,63
TRINITY_DN26569_c0_g1_i9	RAF proto-oncogene serine/threonine-protein kinase	-7,63
TRINITY_DN27692_c1_g1_i11	NAD-dependent protein deacylase sirtuin-5, mitochondrial	-7,63
TRINITY_DN26391_c0_g1_i6	Embryonal Fyn-associated substrate	-7,63
TRINITY_DN28066_c0_g1_i11	Regulatory factor X-associated protein	-7,63
TRINITY_DN29606_c0_g1_i13	Collagen alpha-1(V) chain	-7,63
TRINITY_DN25482_c2_g1_i1	Bromodomain-containing protein 3	-7,63
TRINITY_DN29047_c0_g2_i3	Matrin-3	-7,63
TRINITY_DN32027_c0_g1_i8	Cytochrome P450 2J2	-7,63
TRINITY_DN29120_c0_g2_i5	FACT complex subunit SSRP1	-7,63
TRINITY_DN31229_c0_g1_i10	Exocyst complex component 4	-7,63
TRINITY_DN32938_c1_g2_i9	Misshapen-like kinase 1	-7,63
TRINITY_DN30958_c0_g2_i1	Nuclear receptor ROR-beta	-7,63
TRINITY_DN32474_c4_g1_i2	Collagen alpha-1(XIII) chain	-7,63
TRINITY_DN28850_c0_g1_i5	Nuclear GTPase SLIP-GC	-7,63
TRINITY_DN26993_c2_g1_i20	MAU2 chromatid cohesion factor homolog	-7,63
TRINITY_DN31779_c0_g2_i6	C-type lectin domain family 4 member E	-7,63
TRINITY_DN26919_c2_g1_i2	Somatomedin-B and thrombospondin type-1 domain-containing	-7,63
TRINITY_DN28157_c1_g1_i2	Golgi SNAP receptor complex member 1	-7,63
TRINITY_DN29191_c2_g1_i10	BRISC and BRCA1-A complex member 2	-7,63
TRINITY_DN24086_c4_g6_i4	Netrin receptor UNC5D	-7,63
TRINITY_DN30305_c0_g1_i5	Cdc42 effector protein 4	-7,63
TRINITY_DN31294_c0_g1_i16	Cyclin-L1	-7,63
TRINITY_DN29519_c2_g1_i7	Splicing regulatory glutamine/lysine-rich protein 1	-7,63
TRINITY_DN26148_c0_g1_i4	GATOR complex protein WDR59	-7,63
TRINITY_DN26505_c0_g2_i19	Eukaryotic translation initiation factor 4E transporter	-7,63
TRINITY_DN29974_c0_g1_i9	Cytoplasmic FMR1-interacting protein 2	-7,63
TRINITY_DN24766_c0_g1_i5	Ellis-van Creveld syndrome protein homolog	-7,63
TRINITY_DN25944_c1_g1_i13	Peroxisome proliferator-activated receptor gamma coactivator-	-7,63
TRINITY_DN21737_c0_g1_i2	Mediator of RNA polymerase II transcription subunit 27	-7,63 -7,63
TRINITY_DN31417_c0_g1_i1	Ras-related protein Rab-31	
	•	-7,63
TRINITY_DN32425_c0_g1_i10	Unconventional myosin-IXa	-7,63
TRINITY_DN28233_c1_g3_i1	Glycogen synthase kinase-3 beta	-7,63
TRINITY_DN27670_c1_g1_i5	Axin interactor, dorsalization-associated protein	-7,63
TRINITY_DN31099_c0_g1_i1	Msx2-interacting protein	-7,63
TRINITY_DN24429_c1_g1_i14	H-2 class II histocompatibility antigen, E-S beta chain (Fragme	-7,63
TRINITY_DN32224_c5_g1_i3	Mitochondrial glycine transporter A	-7,63
TRINITY_DN21895_c0_g1_i3	Cathepsin L	-7,63
TRINITY_DN29338_c2_g1_i4	Small G protein signaling modulator 1	-7,63
TRINITY_DN21144_c0_g1_i4	Transmembrane and TPR repeat-containing protein 4	-7,63
TRINITY_DN28709_c0_g1_i9	Translation initiation factor eIF-2B subunit delta	-7,63
TRINITY_DN25562_c0_g1_i5	Kelch-like protein 15	-7,63
TRINITY_DN31707_c0_g2_i1	Guanine nucleotide-binding protein $G(I)/G(S)/G(T)$ subunit beta	-7,63
TRINITY_DN28614_c1_g1_i5	Inorganic pyrophosphatase	-7,63
TRINITY_DN29931_c1_g2_i12	Cytidine monophosphate-N-acetylneuraminic acid hydroxylase	-7,63
TRINITY_DN29123_c2_g1_i9	LON peptidase N-terminal domain and RING finger protein 2	-7,63
TRINITY_DN32141_c0_g2_i5	SPRY domain-containing SOCS box protein 3	-7,63
TRINITY_DN32120_c2_g1_i11	Protein FAM149B1	-7,63
TRINITY_DN25847_c0_g2_i9	Protein 4.1	-7,63
TRINITY_DN27165_c1_g2_i6	Translation initiation factor eIF-2B subunit alpha	-7,63
TRINITY_DN25418_c3_g1_i13	Transposon Tf2-6 polyprotein	-7,63
TRINITY_DN27293_c5_g1_i6	Phospholipase ABHD3	-7,63
TRINITY_DN30404_c1_g2_i17	ATP-dependent 6-phosphofructokinase, liver type	-7,63
TRINITY_DN32171_c0_g2_i6	E3 ubiquitin-protein ligase ZNF598	-7,63
TRINITY_DN26141_c0_g1_i3	Protein YIPF1	-7,63
TRINITY_DN26195_c0_g1_i1	Radical S-adenosyl methionine domain-containing protein 2	-7,63
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TRINITY DNI22200 .0 .1 '2	P FAMOMA	7.62
TRINITY_DN22200_c0_g1_i2	Protein FAM204A	-7,63
TRINITY_DN30706_c0_g1_i19	Keratin, type I cytoskeletal 13	-7,63
TRINITY_DN30311_c1_g2_i13	CMP-N-acetylneuraminate-beta-1,4-galactoside alpha-2,3-sialy	-7,63
TRINITY_DN28371_c1_g2_i2	Sodium- and chloride-dependent creatine transporter 1	-7,63
TRINITY_DN28512_c0_g1_i6	Class I histocompatibility antigen, Gogo-C*0101/C*0102 alpha	-7,58
TRINITY_DN24890_c3_g1_i3	Beta-microseminoprotein	-7,50 7,00
TRINITY_DN25394_c0_g1_i19	Class I histocompatibility antigen, F10 alpha chain	-7,09
TRINITY_DN24543_c6_g2_i6	Lipocalin	-7,03
TRINITY_DN30749_c0_g2_i5	Poly [ADP-ribose] polymerase 12	-6,87
TRINITY_DN33238_c5_g1_i18	Telomerase protein component 1	-6,47
TRINITY_DN25606_c0_g4_i1	Major histocompatibility complex class I-related gene protein	-6,35
TRINITY_DN25606_c0_g6_i1	Class I histocompatibility antigen, F10 alpha chain	-6,28
TRINITY_DN31401_c0_g2_i12	C4b-binding protein alpha chain	-6,24
TRINITY_DN29065_c0_g1_i5	Phosphatidylcholine-sterol acyltransferase (Fragment)	-6,15
TRINITY_DN29858_c1_g1_i1	Forkhead box protein O3	-6,14
TRINITY_DN25594_c3_g2_i1	Mitogen-activated protein kinase kinase kinase 2	-6,07
TRINITY_DN25059_c0_g1_i4	Zinc finger X-chromosomal protein	-6,05
TRINITY_DN30107_c0_g1_i4	Signal transducer and activator of transcription 1	-6,04
TRINITY_DN32865_c1_g1_i4	Major histocompatibility complex class I-related gene protein	-5,95
TRINITY_DN33320_c4_g1_i7	Protein PRRC2B	-5,93
TRINITY_DN30609_c0_g2_i3	Adhesion G protein-coupled receptor L1	-5,92
TRINITY_DN32424_c1_g1_i10	Pericentriolar material 1 protein	-5,90
TRINITY_DN24377_c1_g1_i3	Pancreatic progenitor cell differentiation and proliferation factor	-5,85
TRINITY_DN32243_c2_g1_i12	Centrosomal protein of 152 kDa	-5,84
TRINITY_DN32309_c1_g1_i1	Serum paraoxonase/arylesterase 2	-5,83
TRINITY_DN28833_c1_g1_i5	Casein kinase I isoform alpha	-5,79
TRINITY_DN24493_c0_g1_i3	Retrovirus-related Pol polyprotein from transposon 17.6	-5,76
TRINITY_DN33013_c1_g1_i11	Scavenger receptor class F member 1	-5,74
TRINITY_DN31365_c0_g1_i9	Lysosomal acid phosphatase	-5,69
TRINITY_DN33058_c1_g1_i14	HLA class II histocompatibility antigen, DRB1-13 beta chain	-5,66
TRINITY_DN28242_c1_g1_i2	Inactive tyrosine-protein kinase PEAK1	-5,61
TRINITY_DN32666_c1_g1_i1	Plexin-B1	-5,61
TRINITY_DN32039_c1_g1_i3	Sodium-coupled neutral amino acid transporter 4	-5,59
TRINITY_DN26920_c0_g1_i3	Katanin p80 WD40 repeat-containing subunit B1	-5,55
TRINITY_DN32641_c1_g1_i4	Trafficking protein particle complex subunit 4	-5,49
TRINITY_DN31587_c0_g1_i16	Nuclear receptor subfamily 1 group D member 1	-5,49
TRINITY_DN26919_c3_g1_i15	Septin-7	-5,43
TRINITY_DN32463_c1_g1_i10	Ras-associated and pleckstrin homology domains-containing pr	-5,40
TRINITY_DN25684_c0_g2_i6	Protein Jumonji	-5,38
TRINITY_DN32309_c1_g1_i6	Serum paraoxonase/arylesterase 2	-5,37
TRINITY_DN27367_c1_g1_i7	Proprotein convertase subtilisin/kexin type 5	-5,33
TRINITY_DN32865_c1_g1_i23	RT1 class I histocompatibility antigen, AA alpha chain	-5,31
TRINITY_DN29221_c1_g1_i22	Cordon-bleu protein-like 1	-5,27
TRINITY_DN29931_c1_g2_i17	Cytidine monophosphate-N-acetylneuraminic acid hydroxylase	-5,27
TRINITY_DN29047_c0_g2_i7	Matrin-3	-5,26
TRINITY_DN32414_c0_g1_i7	Neurabin-2	-5,24
TRINITY_DN25075_c4_g1_i3	60S ribosomal protein L15	-5,23
TRINITY_DN32710_c1_g1_i8	Unconventional myosin-Ib	-5,23
TRINITY_DN27774_c7_g1_i11	Androgen receptor	-5,23
TRINITY_DN25394_c0_g1_i15	BOLA class I histocompatibility antigen, alpha chain BL3-7	-5,21
TRINITY_DN26181_c0_g1_i7	Speckle-type POZ protein	-5,20
TRINITY_DN32122_c2_g2_i9	SHC-transforming protein 2	-5,19
TRINITY_DN29047_c0_g2_i9	Matrin-3	-5,18
TRINITY_DN29711_c3_g2_i6	Protein lifeguard 1	-5,15
TRINITY_DN32307_c1_g1_i8	Cytidine deaminase	-5,14
TRINITY_DN29257_c1_g2_i1	Zinc finger CCCH domain-containing protein 7B	-5,14
TRINITY_DN26960_c1_g2_i1	Proliferation-associated protein 2G4	-5,12
TRINITY_DN31906_c1_g2_i6	Fatty aldehyde dehydrogenase	-5,12
TRINITY_DN26219_c0_g1_i1	5'-nucleotidase domain-containing protein 3	-5,12

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TRINITY_DN24501_c0_g1_i7	Serine/arginine-rich splicing factor 1B	-5,08
TRINITY_DN27774_c7_g1_i6	Androgen receptor (Fragment)	-5,03
TRINITY_DN26112_c0_g1_i10	Tripartite motif-containing protein 16	-5,02
TRINITY_DN29816_c1_g1_i18	E3 ubiquitin-protein ligase NEDD4-like	-5,02
TRINITY_DN27863_c1_g1_i10	SLIT-ROBO Rho GTPase-activating protein 2	-4,99
TRINITY_DN30888_c0_g1_i4	Protein Shroom2	-4,97
TRINITY_DN21386_c0_g1_i9	Butyrophilin subfamily 1 member A1	-4,95
TRINITY_DN29800_c1_g4_i3	Aspartate aminotransferase, cytoplasmic	-4,94
TRINITY_DN29005_c0_g1_i2	FYVE, RhoGEF and PH domain-containing protein 1	-4,92
TRINITY_DN26578_c0_g2_i7	Protein fem-1 homolog A	-4,89
TRINITY_DN30334_c1_g1_i2	X-box-binding protein 1	-4,88
TRINITY_DN31950_c0_g1_i13	Tight junction protein ZO-1	-4,88
TRINITY_DN24124_c0_g1_i1	Endothelial lipase	-4,87
TRINITY_DN33061_c0_g1_i2	Suppressor of cytokine signaling 3	-4,84
TRINITY_DN32797_c0_g1_i6	Fatty acid synthase	-4,84
TRINITY_DN28923_c0_g1_i8	TATA-binding protein-associated factor 172	-4,82
TRINITY_DN25699_c0_g1_i7	Nucleolar protein 58	-4,79
TRINITY_DN32246_c0_g1_i4	Coiled-coil domain-containing protein 186	-4,77
TRINITY_DN24890_c3_g1_i4	Beta-microseminoprotein	-4,69
TRINITY_DN23444_c0_g1_i2	Cerebellin-3	-4,68
TRINITY_DN31356_c0_g1_i5	Solute carrier family 25 member 43	-4,67
TRINITY_DN32800_c0_g4_i1	SWI/SNF-related matrix-associated actin-dependent regulator c	-4,63
TRINITY_DN32463_c1_g1_i2	Ras-associated and pleckstrin homology domains-containing pr	-4,63
TRINITY_DN24993_c0_g1_i1	Protein NLRC3	-4,63
TRINITY_DN29936_c0_g7_i1	Transposon TX1 uncharacterized 149 kDa protein	-4,61
TRINITY_DN24659_c0_g1_i4	Histone-binding protein RBBP7	-4,60
TRINITY_DN25069_c1_g1_i13	Sushi, nidogen and EGF-like domain-containing protein 1	-4,58
TRINITY_DN32380_c0_g1_i8		-4,56 -4,56
	Putative ATP-dependent RNA helicase an3	
TRINITY_DN24917_c0_g1_i3	Solute carrier family 25 member 46	-4,54 4,53
TRINITY_DN30696_c1_g1_i9	Zinc finger CCCH domain-containing protein 4	-4,53
TRINITY_DN32865_c1_g1_i22	H-2 class I histocompatibility antigen, K-K alpha chain	-4,51
TRINITY_DN29922_c2_g1_i16	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	-4,48
TRINITY_DN32293_c1_g2_i6	Retrovirus-related Pol polyprotein from transposon 412	-4,45
TRINITY_DN29621_c0_g1_i12	Ras-related protein Rab-8B	-4,45
TRINITY_DN31021_c1_g1_i6	Bifunctional polynucleotide phosphatase/kinase	-4,43
TRINITY_DN30652_c0_g1_i8	Influenza virus NS1A-binding protein homolog B	-4,42
TRINITY_DN28784_c0_g1_i3	Arf-GAP domain and FG repeat-containing protein 1	-4,41
TRINITY_DN31193_c0_g1_i12	Guanine nucleotide-binding protein $G(i)$ subunit alpha-2	-4,40
TRINITY_DN32622_c1_g1_i12	Alanineglyoxylate aminotransferase 2, mitochondrial	-4,39
TRINITY_DN27326_c3_g1_i1	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	-4,38
TRINITY_DN27053_c1_g2_i2	Ubiquitin-60S ribosomal protein L40	-4,38
TRINITY_DN25765_c3_g3_i10	Retinol-binding protein 2	-4,37
TRINITY_DN32412_c3_g4_i6	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	-4,35
TRINITY_DN30473_c3_g1_i3	Ecotropic viral integration site 5 protein homolog	-4,35
TRINITY_DN31365_c0_g1_i6	Lysosomal acid phosphatase	- 4 ,33
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TRINITY_DN25328_c1_g1_i11	KH homology domain-containing protein 4	-4,32
TRINITY_DN33061_c0_g1_i5	Suppressor of cytokine signaling 3	-4,31
TRINITY_DN27104_c2_g1_i15	Ras-related protein R-Ras	-4,31
TRINITY_DN26336_c7_g2_i4	Chromodomain-helicase-DNA-binding protein 6	-4,31
TRINITY_DN25142_c4_g1_i5	Sorting nexin-10A	-4,30
TRINITY_DN32360_c0_g4_i10	Ankyrin repeat and SAM domain-containing protein 1A	-4,28
TRINITY_DN32023_c3_g2_i2	cAMP-specific 3',5'-cyclic phosphodiesterase 4D	-4,28
TRINITY_DN27959_c0_g1_i7	NEDD4 family-interacting protein 2	-4,28
TRINITY_DN25564_c0_g1_i2	Protein QNR-71	-4,27
TRINITY_DN30445_c1_g1_i4	Selenocysteine insertion sequence-binding protein 2	-4,24
TRINITY_DN31963_c0_g1_i10	Leucine-rich repeat and immunoglobulin-like domain-containin	-4,24
TRINITY_DN33114_c4_g2_i2	Angiotensinogen	-4,24
TRINITY_DN28132_c0_g1_i6	Bridging integrator 2	-4,22
TRINITY_DN30989_c0_g1_i12	Translocation protein SEC62	-4,21
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TRINITY_DN33230_c5_g1_i2	Interferon-induced very large GTPase 1	-4,21
TRINITY_DN26934_c2_g1_i8	Insulin-like growth factor II	-4,20
TRINITY_DN31683_c0_g1_i8	Acidic mammalian chitinase	-4,19
TRINITY_DN17760_c0_g1_i2	Acidic mammalian chitinase	-4,19
TRINITY_DN28114_c0_g3_i3	Kelch-like ECH-associated protein 1	-4,19
TRINITY_DN25394_c0_g1_i22	Class I histocompatibility antigen, F10 alpha chain	-4,16
TRINITY_DN29858_c1_g1_i2	Forkhead box protein O3	-4,16
TRINITY_DN31487_c2_g1_i11	Monocarboxylate transporter 7	-4,15
TRINITY_DN26663_c0_g1_i7	Plasma serine protease inhibitor	-4,13
TRINITY_DN33227_c14_g1_i4	Apolipoprotein B-100	-4,13
TRINITY_DN30671_c4_g1_i5	Protein Jumonji	-4,12
TRINITY_DN25156_c0_g1_i4	Glycerol-3-phosphate acyltransferase 3	-4,11
TRINITY_DN32165_c4_g5_i7	CAP-Gly domain-containing linker protein 2	-4,09
TRINITY_DN31421_c0_g1_i8	Rab-interacting lysosomal protein	-4,04
TRINITY_DN32109_c0_g1_i7	Mitochondrial dicarboxylate carrier	-4,04
TRINITY_DN23398_c1_g1_i1	Redox-regulatory protein FAM213A	-4,02
TRINITY_DN28717_c3_g1_i12	BMP/retinoic acid-inducible neural-specific protein 3	-4,02
TRINITY_DN26467_c0_g1_i5	Elongation of very long chain fatty acids protein 1	-4,01
TRINITY_DN27879_c0_g1_i7	Polypyrimidine tract-binding protein 1	-3,99
TRINITY_DN26991_c1_g1_i4	Low-density lipoprotein receptor-related protein 1	-3,99
TRINITY_DN26970_c1_g1_i4	Coiled-coil-helix-coiled-coil-helix domain-containing protein 2	-3,99
TRINITY_DN28687_c1_g4_i14	Bile acid receptor	-3,97
TRINITY_DN29792_c0_g1_i6	Cyclin-dependent kinase inhibitor 1B	-3,97
TRINITY_DN32964_c0_g2_i18	ATP-binding cassette sub-family A member 1	-3,95
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TRINITY_DN25394_c0_g1_i36	BOLA class I histocompatibility antigen, alpha chain BL3-6	-3,94
TRINITY_DN33333_c4_g1_i7	Centriolin	-3,94
TRINITY_DN22840_c0_g1_i8	Growth arrest and DNA damage-inducible protein GADD45 ga	-3,93
TRINITY_DN28602_c0_g1_i4	Cell surface glycoprotein MUC18	-3,92
TRINITY_DN26980_c0_g1_i5	Pannexin-1	-3,90
TRINITY_DN25110_c0_g1_i9	E3 ubiquitin-protein ligase BRE1B	-3,89
TRINITY_DN28143_c0_g1_i7	Arginase, non-hepatic 2	-3,89
TRINITY_DN31421_c0_g1_i9	RILP-like protein 1	-3,87
TRINITY_DN32165_c4_g5_i9	CAP-Gly domain-containing linker protein 2	-3,85
TRINITY_DN29272_c0_g2_i1	Metastasis-associated in colon cancer protein 1	-3,83
TRINITY_DN29079_c1_g3_i4	SWI/SNF complex subunit SMARCC1	-3,80
TRINITY_DN32901_c3_g2_i15	Low-density lipoprotein receptor	-3,80
TRINITY_DN32820_c0_g2_i8	Phospholipid-transporting ATPase IG	-3,78
TRINITY_DN32080_c0_g1_i5	Histone-lysine N-methyltransferase SETD2	-3,77
TRINITY_DN32251_c1_g2_i6	Protein broad-minded	-3,77
TRINITY_DN28857_c0_g1_i2	TNFAIP3-interacting protein 1	-3,73
TRINITY_DN30475_c0_g2_i3	Lysine-specific demethylase 6B	-3,72
TRINITY_DN30012_c0_g1_i3	Probable RNA-directed DNA polymerase from transposon X-ele	-3,71
TRINITY_DN32720_c2_g3_i2	25-hydroxyvitamin D-1 alpha hydroxylase, mitochondrial	-3,71
TRINITY_DN32898_c1_g1_i7	UDP-xylose and UDP-N-acetylglucosamine transporter	-3,69
TRINITY_DN25970_c3_g3_i16	Protein ABHD4	-3,67
TRINITY_DN30309_c1_g1_i8	Acetyl-coenzyme A synthetase, cytoplasmic	-3,67
TRINITY_DN32729_c0_g1_i20	Aldose 1-epimerase	-3,66
TRINITY_DN28750_c3_g1_i1	CD276 antigen	-3,65
TRINITY_DN29825_c0_g2_i3	Olfactomedin-like protein 2A	-3,63
TRINITY_DN33003_c1_g1_i9	Choline kinase alpha	-3,60
TRINITY_DN26425_c3_g1_i2	Translocation protein SEC63 homolog	-3,59
TRINITY_DN30680_c4_g1_i6	Interferon-induced very large GTPase 1	-3,54
TRINITY_DN30475_c0_g2_i6	Lysine-specific demethylase 6B	-3,54 -3,53
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TRINITY_DN29856_c0_g2_i7	Sodium-coupled neutral amino acid transporter 3	-3,53
TRINITY_DN33104_c6_g1_i27	Histidine ammonia-lyase	-3,53
TRINITY_DN29612_c0_g1_i5	Protein ATP1B4	-3,51
TRINITY_DN30723_c0_g1_i4	Mitogen-activated protein kinase kinase kinase kinase 3	-3,48
TRINITY_DN32368_c0_g2_i10	Oxysterols receptor LXR-alpha	-3,46
TRINITY_DN33029_c2_g1_i16	$Dihydropyrimidine\ dehydrogenase\ [NADP(+)]$	-3,44

TRINITY_DN31912_c0_g1_i15	High affinity copper uptake protein 1	-3,44
TRINITY_DN33029_c1_g1_i1	Protein phosphatase 1 regulatory subunit 15B	-3,43
TRINITY_DN29922_c2_g1_i5	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	-3,42
TRINITY_DN28924_c0_g1_i8	RNA-binding protein 20	-3,42
TRINITY_DN32109_c0_g1_i21	Mitochondrial dicarboxylate carrier	-3,42
TRINITY_DN31335_c0_g1_i1	Transitional endoplasmic reticulum ATPase	-3,42
TRINITY_DN29289_c0_g1_i7	Ubiquitin carboxyl-terminal hydrolase MINDY-1	-3,41
TRINITY_DN30364_c0_g2_i9	Basic leucine zipper and W2 domain-containing protein 1-B	-3,41
TRINITY_DN28335_c0_g2_i1	Staphylococcal nuclease domain-containing protein 1	-3,40
TRINITY_DN31368_c1_g3_i4	MAP/microtubule affinity-regulating kinase 3	-3,40
TRINITY_DN30969_c0_g1_i8	Protein CASC4	-3,39
TRINITY_DN29335_c0_g1_i14	PRELI domain containing protein 3B	-3,36
TRINITY_DN31340_c2_g3_i6	Sterol O-acyltransferase 1	-3,36
TRINITY_DN32236_c0_g1_i9	Beta-glucuronidase	-3,36
TRINITY_DN25632_c0_g1_i9	Growth arrest and DNA damage-inducible protein GADD45 be	-3,34
TRINITY_DN31474_c5_g3_i9	Transmembrane protein 131	-3,34
TRINITY_DN30309_c1_g1_i4	Acetyl-coenzyme A synthetase, cytoplasmic	-3,33
TRINITY_DN29157_c0_g2_i3	RNA-binding motif, single-stranded-interacting protein 2	-3,32
TRINITY_DN25394_c0_g1_i10		-3,29
TRINITY_DN33318_c5_g2_i1	HLA class I histocompatibility antigen, B-40 alpha chain	
	Chloride transport protein 6	-3,28
TRINITY_DN24666_c1_g1_i1	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 3	-3,28
TRINITY_DN31472_c0_g1_i2	Probable fibrosin-1	-3,27
TRINITY_DN27237_c0_g1_i5	Cis-aconitate decarboxylase	-3,27
TRINITY_DN33071_c2_g1_i4	UDP-N-acetylglucosaminepeptide N-acetylglucosaminyltransj	-3,27
TRINITY_DN24221_c0_g1_i5	Nuclear factor interleukin-3-regulated protein	-3,26
TRINITY_DN31172_c0_g1_i13	OTU domain-containing protein 3	-3,23
TRINITY_DN31365_c0_g3_i4	Lysosomal acid phosphatase	-3,23
TRINITY_DN33333_c4_g1_i9	Complement C5	-3,20
TRINITY_DN32865_c1_g1_i30	HLA class I histocompatibility antigen, alpha chain F	-3,18
TRINITY_DN33149_c0_g1_i4	Protein GREB1	-3,17
TRINITY_DN31279_c0_g1_i7	Insulin receptor substrate 1-B	-3,16
TRINITY_DN31365_c0_g1_i8	Lysosomal acid phosphatase	-3,14
TRINITY_DN33322_c1_g2_i2	Sacsin	-3,13
TRINITY_DN32923_c1_g1_i7	F-box only protein 9	-3,12
TRINITY_DN33295_c5_g2_i4	Tight junction protein ZO-1	-3,12
TRINITY_DN32100_c1_g4_i9	Calcium/calmodulin-dependent protein kinase type II subunit ge	-3,08
TRINITY_DN33088_c3_g1_i2	Inter-alpha-trypsin inhibitor heavy chain H3	-3,06
TRINITY_DN20714_c0_g1_i3	Suppressor of cytokine signaling 3	-2,99
TRINITY_DN23839_c0_g1_i2	Cytokine-inducible SH2-containing protein	-2,93
TRINITY_DN25606_c0_g3_i3	Class I histocompatibility antigen, F10 alpha chain	-2,92
TRINITY_DN26189_c0_g1_i10	S-adenosylmethionine synthase isoform type-2	-2,87
TRINITY_DN31599_c0_g1_i2	Retrovirus-related Pol polyprotein from transposon 17.6	-2,79
Transcritos upregulated		<u> </u>
TRINITY_DN31753_c0_g1_i4	Mid1-interacting protein 1-B	2,76
TRINITY_DN31606_c0_g1_i16	Endoplasmin (Fragment)	2,82
TRINITY_DN31157_c1_g1_i20	Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondr	2,85
TRINITY_DN32127_c1_g3_i7	NUAK family SNF1-like kinase 1	2,87
TRINITY_DN31748_c0_g1_i1	Acetyl-CoA acetyltransferase, mitochondrial	2,87
TRINITY_DN26693_c0_g1_i5	Creatine kinase U-type, mitochondrial	2,88
TRINITY_DN32493_c0_g1_i14	Carboxylesterase 5A	2,89
TRINITY_DN29862_c2_g1_i27	Sulfotransferase family cytosolic 2B member 1	2,92
TRINITY_DN30830_c0_g1_i2	Calumenin-A	2,93
TRINITY_DN31363_c1_g1_i10	Pumilio homolog 1	2,95
TRINITY_DN29275_c0_g2_i1	Cysteine dioxygenase type 1	2,95
TRINITY_DN31606_c0_g1_i20	Endoplasmin	2,96
TRINITY_DN28227_c2_g2_i5	Ornithine aminotransferase, mitochondrial	2,90
		2,98
TRINITY_DN33318_c5_g2_i2 TRINITY_DN33002_c1_g2_i5	Chloride transport protein 6	
TRINITY_DN33002_c1_g2_i5	DnaJ homolog subfamily C member 3 Pilosome hinding protein 1	3,00
TRINITY_DN32983_c1_g1_i3	Ribosome-binding protein 1	3,01

TRINITY_DN30476_c0_g1_i3	Apolipoprotein F	3,01
TRINITY_DN32840_c3_g1_i4	Hypoxia up-regulated protein 1	3,01
TRINITY_DN29858_c1_g1_i9	Forkhead box protein O3	3,04
TRINITY_DN25425_c3_g2_i4	•	3,04
_	Heat shock protein HSP 90-alpha	
TRINITY_DN28007_c0_g1_i2	Transmembrane emp24 domain-containing protein 9	3,07
TRINITY_DN29429_c1_g1_i11	Phenylalanine-4-hydroxylase	3,12
TRINITY_DN32360_c0_g4_i13	Ankyrin repeat and SAM domain-containing protein 1A	3,13
TRINITY_DN33112_c5_g1_i2	Carnitine O-palmitoyltransferase 1, liver isoform	3,14
TRINITY_DN28027_c1_g2_i1	Tectonin beta-propeller repeat-containing protein 1	3,15
TRINITY_DN30520_c2_g1_i5	Manganese-transporting ATPase 13A1	3,15
TRINITY_DN28883_c3_g2_i7	Insulin-like growth factor I, adult form	3,17
TRINITY_DN32174_c0_g3_i6	Probable E3 ubiquitin-protein ligase HERC4	3,18
TRINITY_DN25659_c1_g1_i3	Dolichyl-diphosphooligosaccharideprotein glycosyltransferas	3,19
TRINITY_DN31558_c0_g1_i2	Chromodomain-helicase-DNA-binding protein 2	3,20
TRINITY_DN23671_c0_g1_i4	Lipoyl synthase, mitochondrial	3,20
TRINITY_DN31753_c0_g1_i2	Mid1-interacting protein 1-B	3,21
TRINITY_DN32954_c2_g1_i2	Endoplasmic reticulum chaperone BiP	3,22
TRINITY_DN24330_c0_g1_i6	N-lysine methyltransferase KMT5A-A	3,22
TRINITY_DN28800_c1_g1_i2	Heat shock protein HSP 90-alpha	3,22
TRINITY_DN30380_c4_g1_i12	10 kDa heat shock protein, mitochondrial	3,23
TRINITY_DN29375_c1_g1_i8	Metastasis suppressor protein 1	3,24
TRINITY_DN28193_c0_g2_i2	Transmembrane protein 208	3,24
TRINITY_DN32180_c0_g1_i8	Complement C3 (Fragment)	3,25
TRINITY_DN32363_c1_g2_i1	Vigilin	3,27
TRINITY_DN32433_c2_g1_i5	Probable E3 ubiquitin-protein ligase HERC4	3,27
TRINITY_DN24030_c2_g1_i3	Serine/threonine-protein phosphatase 6 regulatory subunit 2	3,28
TRINITY_DN32167_c0_g1_i8	Vitelline membrane outer layer protein 1 homolog	3,28
TRINITY_DN28171_c4_g1_i5	Derlin-2	3,28
TRINITY_DN31551_c1_g1_i6		3,28
	Glutaminyl-peptide cyclotransferase	
TRINITY_DN27777_c1_g1_i14	Protein transport protein Sec31A	3,32
TRINITY_DN31924_c1_g1_i24	Ribose-phosphate pyrophosphokinase 1	3,32
TRINITY_DN30813_c0_g1_i3	Protein phosphatase 1 regulatory subunit 3B	3,34
TRINITY_DN27501_c2_g1_i3	Polyadenylate-binding protein 2	3,34
TRINITY_DN25468_c0_g1_i5	CCA tRNA nucleotidyltransferase 1, mitochondrial	3,35
TRINITY_DN29388_c0_g2_i6	Nuclear factor erythroid 2-related factor 2	3,38
TRINITY_DN31889_c1_g1_i4	A disintegrin and metalloproteinase with thrombospondin motif.	3,39
TRINITY_DN30714_c2_g1_i9	Very low-density lipoprotein receptor	3,40
TRINITY_DN29078_c1_g1_i5	Vitelline membrane outer layer protein 1	3,40
TRINITY_DN27544_c0_g1_i3	Dynamin-2	3,41
TRINITY_DN28223_c1_g1_i1	Hydroxymethylglutaryl-CoA synthase, cytoplasmic	3,42
TRINITY_DN25425_c3_g2_i1	Heat shock protein HSP 90-alpha	3,42
TRINITY_DN27106_c1_g2_i13	Motile sperm domain-containing protein 1	3,43
TRINITY_DN31065_c3_g1_i20	Cytoskeleton-associated protein 4	3,43
TRINITY_DN32307_c1_g1_i20	Cytidine deaminase	3,45
TRINITY_DN33224_c10_g1_i13	Equilibrative nucleoside transporter 1	3,45
TRINITY_DN30380_c4_g1_i16	10 kDa heat shock protein, mitochondrial	3,46
TRINITY_DN29810_c0_g1_i7	Glycosyltransferase 8 domain-containing protein 1	3,46
TRINITY_DN29458_c0_g1_i32	Phosphatidate phosphatase LPIN1	3,46
TRINITY_DN31416_c0_g1_i7	Glucocorticoid receptor	3,47
TRINITY_DN32808_c2_g1_i4	Cell death activator CIDE-A	3,47
TRINITY_DN30035_c2_g1_i6	Group 3 secretory phospholipase A2	3,47
TRINITY_DN31522_c1_g2_i8	Choline-phosphate cytidylyltransferase A	3,49
TRINITY_DN26942_c3_g2_i1	Isoaspartyl peptidase/L-asparaginase	3,50
TRINITY_DN31256_c1_g2_i21	E3 ubiquitin-protein ligase HECTD1	3,50
TRINITY_DN29713_c0_g1_i3	Glutaryl-CoA dehydrogenase, mitochondrial	3,50
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TRINITY_DN30304_c1_g1_i8	Kynurenine 3-monooxygenase Cold shook domain containing protein C2	3,52
TRINITY_DN31990_c1_g1_i18	Cold shock domain-containing protein C2	3,53
TRINITY_DN27472_c0_g1_i5	Glutathione reductase, mitochondrial	3,53
TRINITY_DN21998_c0_g1_i1	Peptidyl-prolyl cis-trans isomerase FKBP4	3,54

TDD WTW DWG 110 0 1 10	GOPO : I I I I I I	2.56
TRINITY_DN23418_c0_g1_i3	COP9 signalosome complex subunit 5	3,56
TRINITY_DN31280_c1_g1_i1	Phosphatidylinositol-binding clathrin assembly protein	3,57
TRINITY_DN31172_c0_g1_i11	OTU domain-containing protein 3	3,60
TRINITY_DN28882_c0_g2_i5	Nuclear pore complex protein Nup155	3,60
TRINITY_DN31902_c0_g1_i5	E3 ubiquitin-protein ligase mib1	3,61
TRINITY_DN33113_c1_g1_i5	Protein LLP homolog	3,63
TRINITY_DN31172_c0_g1_i8	OTU domain-containing protein 3	3,65
TRINITY_DN29416_c2_g4_i8	Translocon-associated protein subunit alpha	3,65
TRINITY_DN33098_c1_g2_i2	Probable fructose-2,6-bisphosphatase TIGAR A	3,66
TRINITY_DN31537_c2_g1_i4	Retinol dehydrogenase 13	3,67
TRINITY_DN26812_c4_g1_i7	Electron transfer flavoprotein subunit beta	3,70
TRINITY_DN27203_c2_g1_i1	3'(2'),5'-bisphosphate nucleotidase 1	3,70
TRINITY_DN26970_c1_g1_i13	Coiled-coil-helix-coiled-coil-helix domain-containing protein 2	3,72
TRINITY_DN28479_c1_g3_i9	Serine/arginine-rich splicing factor 3	3,75
TRINITY_DN30445_c1_g1_i16	Selenocysteine insertion sequence-binding protein 2	3,76
TRINITY_DN31081_c1_g1_i6	Phytanoyl-CoA dioxygenase domain-containing protein 1	3,77
TRINITY_DN27269_c1_g1_i4	E3 ubiquitin-protein ligase MARCH5	3,78
TRINITY_DN28175_c0_g1_i10	Peptidyl-prolyl cis-trans isomerase B	3,78
TRINITY_DN27791_c1_g1_i1	BAG family molecular chaperone regulator 3	3,79
TRINITY_DN28335_c0_g2_i3	Staphylococcal nuclease domain-containing protein 1	3,79
TRINITY_DN28427_c0_g1_i13	Vitellogenin	3,79
TRINITY_DN29755_c2_g1_i4	Transcription factor E3	3,80
TRINITY_DN30863_c1_g1_i4	Inversin	3,80
TRINITY_DN29516_c2_g1_i9	Complement factor H	3,80
TRINITY_DN29508_c0_g1_i3	Prolyl 4-hydroxylase subunit alpha-2	3,80
TRINITY_DN32180_c0_g1_i12	Complement C3 (Fragment)	3,81
TRINITY_DN27420_c1_g3_i1	Translocon-associated protein subunit beta	3,82
TRINITY_DN28649_c0_g1_i3	Poly(U)-binding-splicing factor PUF60	3,84
TRINITY_DN32188_c2_g1_i14	Elongation of very long chain fatty acids protein 6	3,85
TRINITY_DN32927_c1_g1_i5	Protein kinase C and casein kinase II substrate protein 3	3,86
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TRINITY_DN29936_c0_g2_i1	Transposon TX1 uncharacterized 149 kDa protein	3,86
TRINITY_DN30508_c1_g2_i2	60 kDa heat shock protein, mitochondrial	3,86
TRINITY_DN32180_c0_g2_i1	Complement C3	3,87
TRINITY_DN25083_c0_g1_i2	Islet cell autoantigen 1-like protein	3,88
TRINITY_DN29585_c0_g1_i4	Nck-associated protein 1	3,88
TRINITY_DN33332_c2_g2_i5	Solute carrier family 25 member 36-A	3,90
TRINITY_DN32159_c0_g1_i12	CSC1-like protein 2	3,93
TRINITY_DN29263_c0_g3_i2	Nuclear receptor-binding protein 2	3,93
TRINITY_DN32725_c1_g3_i12	Actin-binding LIM protein 1	3,93
TRINITY_DN31173_c0_g2_i3	Ensconsin	3,93
TRINITY_DN31906_c1_g2_i8	Fatty aldehyde dehydrogenase	3,93
TRINITY_DN31875_c1_g1_i1	Protein	3,95
TRINITY_DN28086_c0_g1_i1	Alpha-taxilin	3,96
TRINITY_DN24822_c2_g1_i13	Vitelline membrane outer layer protein 1 homolog	3,96
TRINITY_DN25171_c0_g1_i3	Complement C3 (Fragment)	3,98
TRINITY_DN25755_c0_g1_i3	Hemoglobin subunit alpha	3,99
TRINITY_DN31757_c2_g1_i13	DEP domain-containing mTOR-interacting protein	4,03
TRINITY_DN33062_c3_g1_i3	Rho guanine nucleotide exchange factor 5	4,04
TRINITY_DN29565_c1_g1_i15		4,04
	Mucolipin-1	
TRINITY_DN31928_c3_g1_i8	EMILIN-1	4,04
TRINITY_DN24833_c4_g1_i4	Protein FAM53B	4,04
TRINITY_DN24110_c3_g1_i1	Vasopressin V2 receptor	4,05
TRINITY_DN28086_c0_g1_i12	Alpha-taxilin	4,07
TRINITY_DN31357_c2_g3_i6	Insulin-like growth factor-binding protein 1	4,08
TRINITY_DN33114_c4_g2_i1	Angiotensinogen	4,08
TRINITY_DN30708_c1_g2_i7	MOB kinase activator 2	4,13
TRINITY_DN30110_c0_g1_i5	Ras-related protein Rab-33B	4,17
TRINITY_DN31632_c0_g2_i1	Complement C3	4,17
TRINITY_DN26075_c4_g1_i2	Survival motor neuron protein 1	4,20
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TRINITY_DN28892_c0_g3_i3	Guanine nucleotide-binding protein $G(t)$ subunit alpha-2	4,23
TRINITY_DN25761_c2_g1_i3	Arrestin red cell isoform 3	4,24
TRINITY_DN32161_c1_g1_i4	Patched domain-containing protein 3	4,24
TRINITY_DN32566_c0_g1_i5	Receptor-type tyrosine-protein phosphatase S	4,24
TRINITY_DN26806_c0_g1_i2	E3 ubiquitin-protein ligase ARIH2	4,25
TRINITY_DN30153_c0_g1_i2	Transmembrane protein 39A	4,25
TRINITY_DN32650_c5_g1_i3	Amine sulfotransferase	4,26
TRINITY_DN27453_c0_g2_i5	V-type proton ATPase subunit C 1-B	4,27
TRINITY_DN29535_c1_g1_i4	Nuclear factor interleukin-3-regulated protein	4,28
TRINITY_DN33040_c4_g2_i2	Protein FAM210B, mitochondrial	4,30
TRINITY_DN31632_c0_g2_i3	Complement C3	4,33
TRINITY_DN32710_c1_g1_i9	Unconventional myosin-Ib	4,34
TRINITY_DN28942_c3_g1_i1	Receptor-binding cancer antigen expressed on SiSo cells	4,34
_		4,34
TRINITY_DN28603_c0_g1_i6	NLR family CARD domain-containing protein 3	
TRINITY_DN31990_c1_g1_i6	Calcium-regulated heat-stable protein 1	4,34
TRINITY_DN30131_c0_g1_i5	Sentrin-specific protease 2	4,35
TRINITY_DN32781_c0_g1_i24	SUN domain-containing	4,35
TRINITY_DN31992_c2_g1_i12	Complement C1q subcomponent subunit C	4,39
TRINITY_DN32877_c0_g1_i7	Disabled homolog 1	4,40
TRINITY_DN26086_c0_g1_i3	TyrosinetRNA ligase, cytoplasmic	4,40
TRINITY_DN29932_c1_g1_i6	Serine/threonine-protein kinase Sgk1	4,43
TRINITY_DN32180_c0_g2_i2	Complement C3 (Fragment)	4,43
TRINITY_DN25699_c0_g1_i10	Nucleolar protein 58	4,43
TRINITY_DN31606_c0_g1_i5	Endoplasmin	4,45
TRINITY_DN27403_c0_g1_i14	Long-chain-fatty-acidCoA ligase ACSBG1	4,46
TRINITY_DN29734_c0_g1_i4	Mucin-5AC	4,46
TRINITY_DN30393_c0_g1_i2	RNA helicase aquarius	4,46
TRINITY_DN32965_c1_g1_i1	Cytochrome P450 2K1	4,48
TRINITY_DN29282_c1_g1_i8	Monoacylglycerol lipase ABHD12	4,49
TRINITY_DN32072_c0_g2_i13	Mitochondrial import inner membrane translocase subunit Tim.	4,51
TRINITY_DN31606_c0_g1_i6	Endoplasmin (Fragment)	4,53
TRINITY_DN31743_c2_g3_i9	Calreticulin	4,55
TRINITY_DN25394_c0_g1_i27	Class I histocompatibility antigen, F10 alpha chain	4,57
TRINITY_DN27769_c0_g1_i8	60 kDa SS-A/Ro ribonucleoprotein	4,57
	Coatomer subunit zeta-1	4,57
TRINITY_DN27726_c2_g1_i12	-	
TRINITY_DN29692_c2_g1_i9	60S ribosomal protein L13a	4,60
TRINITY_DN29995_c0_g1_i15	Phospholipase A1 member A	4,61
TRINITY_DN32740_c0_g1_i2	Vitellogenin	4,61
TRINITY_DN22741_c0_g1_i1	3-hydroxyisobutyryl-CoA hydrolase, mitochondrial	4,62
TRINITY_DN32465_c0_g1_i11	Formimidoyltransferase-cyclodeaminase	4,62
TRINITY_DN30690_c2_g1_i2	Lysine-specific demethylase 4B	4,67
TRINITY_DN29084_c6_g1_i2	Folylpolyglutamate synthase, mitochondrial	4,68
TRINITY_DN33224_c10_g1_i4	Equilibrative nucleoside transporter 1	4,68
TRINITY_DN28157_c1_g1_i7	Golgi SNAP receptor complex member 1	4,69
TRINITY_DN25709_c1_g3_i6	Fatty acid-binding protein 10-A, liver basic	4,70
TRINITY_DN31202_c0_g1_i10	1-phosphatidylinositol 3-phosphate 5-kinase	4,72
TRINITY_DN30585_c0_g1_i3	Hemopexin	4,72
TRINITY_DN28856_c2_g1_i10	Plastin-1	4,73
TRINITY_DN32260_c2_g1_i12	Cationic amino acid transporter 2	4,75
TRINITY_DN29531_c2_g1_i25	Nuclear factor 1 X-type	4,81
TRINITY_DN25060_c1_g1_i9	Vasodilator-stimulated phosphoprotein	4,82
TRINITY_DN25838_c2_g3_i3	Alpha-1-inhibitor 3	4,93
TRINITY_DN23797_c1_g1_i16	Methylmalonate-semialdehyde dehydrogenase [acylating], mito	4,93
_		4,95
TRINITY_DN32154_c1_g1_i11	Long-chain-fatty-acid-CoA ligase 1	
TRINITY_DN32154_c1_g1_i9	Long-chain-fatty-acidCoA ligase 1	4,97
TRINITY_DN28586_c0_g1_i3	Mitogen-activated protein kinase kinase kinase 4	4,97
TRINITY_DN30690_c1_g1_i7	Protein FAM50A	4,97
TRINITY_DN32382_c0_g1_i12	Synergin gamma	5,01
TRINITY_DN24375_c0_g1_i3	Mothers against decapentaplegic homolog 3	5,01

TRINITY DNOOCO 12 1 14	D	5.00
TRINITY_DN29962_c2_g1_i4	Protein disulfide-isomerase A4	5,02
TRINITY_DN32980_c3_g5_i16	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	5,03
TRINITY_DN24494_c0_g1_i8	Glutaredoxin 3	5,05
TRINITY_DN32192_c2_g2_i6	Fatty acid desaturase 2	5,08
TRINITY_DN27438_c1_g3_i8	cAMP-specific 3',5'-cyclic phosphodiesterase 4B	5,08
TRINITY_DN22718_c0_g1_i4	CBP80/20-dependent translation initiation factor	5,15
TRINITY_DN29769_c2_g2_i6	Chloride transport protein 6	5,20
TRINITY_DN30452_c0_g2_i14	DnaJ homolog subfamily A member 1	5,22
TRINITY_DN25685_c0_g1_i9	Transmembrane protein 220	5,22
TRINITY_DN30069_c0_g3_i6	Glycogen [starch] synthase, muscle	5,23
TRINITY_DN30002_c2_g1_i11	Major histocompatibility complex class I-related gene protein	5,23
TRINITY_DN25394_c0_g1_i25	DLA class I histocompatibility antigen, A9/A9 alpha chain	5,27
TRINITY_DN26413_c3_g1_i9	Breast cancer metastasis-suppressor 1-like protein-A	5,29
TRINITY_DN26900_c2_g2_i9	60S ribosomal protein L5	5,32
TRINITY_DN32180_c0_g1_i5	Complement C3 (Fragment)	5,35
TRINITY_DN23002_c0_g1_i6	Gem-associated protein 5	5,38
	-	
TRINITY_DN27508_c1_g1_i8	Fibroblast growth factor receptor 2	5,38
TRINITY_DN23290_c0_g2_i8	Protein YIPF6	5,41
TRINITY_DN27019_c1_g1_i5	Complement C3 (Fragment)	5,41
TRINITY_DN28936_c0_g1_i8	Phosphatidate cytidylyltransferase 2	5,43
TRINITY_DN30740_c0_g4_i3	Transmembrane protein 65	5,46
TRINITY_DN31058_c0_g2_i10	Moesin	5,46
TRINITY_DN25427_c2_g1_i8	TBC1 domain family member 25	5,49
TRINITY_DN31587_c0_g1_i11	Nuclear receptor subfamily 1 group D member 1	5,57
TRINITY_DN30964_c0_g1_i11	Hepatocyte nuclear factor 1-beta-B	5,57
TRINITY_DN30585_c0_g1_i2	Hemopexin	5,62
TRINITY_DN26436_c0_g1_i1	Histone H1.0	5,64
TRINITY_DN25974_c0_g1_i4	Activating transcription factor 7-interacting protein 1	5,69
TRINITY_DN28592_c0_g1_i10	Serine/threonine-protein phosphatase 6 regulatory ankyrin repe	5,69
TRINITY_DN31958_c0_g1_i6	Ubiquitin-associated protein 2-like	5,70
TRINITY_DN33002_c1_g2_i6	DnaJ homolog subfamily C member 3	5,71
TRINITY_DN30531_c0_g1_i7	Coiled-coil and C2 domain-containing protein 1A	5,72
TRINITY_DN27116_c0_g1_i7	Autophagy-related protein 101	5,74
TRINITY_DN26453_c0_g1_i2	Multidrug resistance-associated protein 5	5,78
TRINITY_DN25461_c3_g2_i14	Polyadenylate-binding protein-interacting protein 2B	5,78
TRINITY_DN33132_c1_g1_i2	Peptidyl-prolyl cis-trans isomerase FKBP5	5,81
TRINITY_DN26436_c0_g1_i4	Histone H1.0	5,84
TRINITY_DN25394_c0_g1_i1		5,87
_	Patr class I histocompatibility antigen, A-5 alpha chain	
TRINITY_DN32356_c3_g1_i4	Striatin	5,95
TRINITY_DN32682_c1_g1_i22	Hypoxia-inducible factor 1-alpha	5,97
TRINITY_DN26200_c0_g1_i6	Transmembrane protein 185-like	5,97
TRINITY_DN24563_c0_g1_i2	Diphosphomevalonate decarboxylase	6,03
TRINITY_DN33274_c9_g3_i4	MAP7 domain-containing protein 1	6,03
TRINITY_DN27443_c0_g2_i2	Heat shock cognate 71 kDa protein	6,05
TRINITY_DN27116_c0_g1_i5	Autophagy-related protein 101	6,06
TRINITY_DN29260_c0_g1_i2	Cullin-4B	6,06
TRINITY_DN30752_c1_g1_i13	Signal recognition particle subunit SRP68	6,11
TRINITY_DN32865_c1_g1_i20	Major histocompatibility complex class I-related gene protein	6,12
TRINITY_DN28211_c0_g1_i3	Ferritin, middle subunit	6,19
TRINITY_DN30473_c3_g1_i7	Ecotropic viral integration site 5 protein homolog	6,37
TRINITY_DN31085_c1_g1_i7	Selenoprotein O	6,49
TRINITY_DN29483_c2_g2_i3	Cystatin	6,50
TRINITY_DN30467_c0_g1_i8	Cysteine-rich with EGF-like domain protein 2	6,52
TRINITY_DN32247_c1_g1_i5	Tensin-2	6,53
TRINITY_DN32192_c2_g2_i3	Fatty acid desaturase 2	6,77
TRINITY_DN30600_c3_g1_i3	Insulin-like growth factor 2 mRNA-binding protein 3	7,03
TRINITY_DN27228_c0_g1_i6	Diamine acetyltransferase 1	7,09
TRINITY_DN26877_c1_g1_i3	Catechol O-methyltransferase A	7,05
TRINITY_DN30688_c0_g1_i10	Cytochrome P450 2D17	7,67
	System of the BD11	7,07

TDINITY DN21600 .0 .1 :11	D (NI DC)	7.67
TRINITY_DN31680_c0_g1_i11	Protein NLRC3	7,67
TRINITY_DN29750_c2_g1_i6	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	7,67
TRINITY_DN22821_c0_g1_i3	Normal mucosa of esophagus-specific gene 1 protein	7,67
TRINITY_DN31452_c2_g1_i9	Vinexin	7,67
TRINITY_DN33052_c0_g1_i1	Hermansky-Pudlak syndrome 6 protein homolog	7,67
TRINITY_DN26914_c1_g1_i2	Palmitoyltransferase ZDHHC17	7,67
TRINITY_DN31523_c0_g2_i4	Phosphorylase b kinase regulatory subunit alpha, liver isoform	7,67
TRINITY_DN25248_c0_g1_i1	Lysosomal protective protein	7,67
TRINITY_DN28960_c0_g1_i19	Interferon-induced GTP-binding protein Mx	7,67
TRINITY_DN30211_c0_g3_i3	FERM domain-containing protein 4B	7,67
TRINITY_DN22525_c0_g1_i2	Vacuolar protein sorting-associated protein 13C	7,67
TRINITY_DN31602_c2_g1_i8	Partitioning defective 3 homolog	7,67
TRINITY_DN29565_c1_g1_i13	Mucolipin-1	7,67
TRINITY_DN32731_c0_g2_i9	•	7,67
	Heterogeneous nuclear ribonucleoprotein C Annexin A6	
TRINITY_DN27789_c2_g2_i6		7,67
TRINITY_DN31852_c2_g1_i10	Pre-mRNA-splicing factor 18	7,67
TRINITY_DN29601_c2_g1_i7	Claudin-10	7,67
TRINITY_DN30637_c0_g1_i2	Phosphatidylinositol 4-phosphate 5-kinase type-1 gamma	7,67
TRINITY_DN28936_c0_g1_i10	Phosphatidate cytidylyltransferase 2	7,67
TRINITY_DN30964_c0_g1_i7	Hepatocyte nuclear factor 1-beta-B	7,67
TRINITY_DN27588_c0_g1_i10	Nucleolar and coiled-body phosphoprotein 1	7,67
TRINITY_DN28494_c0_g1_i6	Transcription elongation regulator 1	7,67
TRINITY_DN25703_c1_g1_i7	Ankyrin repeat domain-containing protein 39	7,67
TRINITY_DN29037_c0_g1_i1	U2 snRNP-associated SURP motif-containing protein	7,67
TRINITY_DN26746_c0_g2_i3	Receptor tyrosine-protein kinase erbB-3	7,67
TRINITY_DN33063_c4_g3_i11	Bone morphogenetic protein 1	7,67
TRINITY_DN26080_c1_g3_i3	Rho-related GTP-binding protein RhoA-D	7,67
TRINITY_DN32311_c1_g1_i6	Histone deacetylase 4	7,67
TRINITY_DN29896_c0_g1_i4	V-type proton ATPase subunit E 1	7,67
TRINITY_DN29186_c0_g1_i3	TBC1 domain family member 12	7,67
TRINITY_DN27902_c1_g2_i1	Embryonic stem cell-specific 5-hydroxymethylcytosine-binding	7,67
TRINITY_DN31959_c1_g1_i1	Solute carrier organic anion transporter family member 1C1	7,67
	Kelch domain-containing protein 3	7,67
TRINITY_DN27250_c1_g2_i4	~ ·	
TRINITY_DN32620_c0_g2_i16	Lysine-specific demethylase phf2	7,67
TRINITY_DN25899_c0_g1_i2	[Protein ADP-ribosylarginine] hydrolase	7,67
TRINITY_DN32611_c1_g2_i15	Death-inducer obliterator 1	7,67
TRINITY_DN26605_c0_g1_i19	Transcription factor 7-like 2	7,67
TRINITY_DN32450_c0_g1_i7	Lysine-specific demethylase 2B	7,67
TRINITY_DN28889_c0_g1_i1	Nuclear factor NF-kappa-B p100 subunit	7,67
TRINITY_DN26227_c0_g1_i7	Diamine acetyltransferase 2	7,67
TRINITY_DN30540_c2_g1_i11	Vitamin K-dependent protein C	7,67
TRINITY_DN33107_c0_g1_i10	Highly reducing polyketide synthase sdnO	7,67
TRINITY_DN22781_c0_g1_i4	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase	7,67
TRINITY_DN25033_c1_g3_i3	Hepatocyte nuclear factor 6	7,67
TRINITY_DN26648_c0_g1_i10	Guanine nucleotide-binding protein subunit alpha-12	7,67
TRINITY_DN26131_c0_g2_i3	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 4	7,67
TRINITY_DN26630_c0_g2_i11	Heterogeneous nuclear ribonucleoprotein L	7,67
TRINITY_DN26223_c0_g2_i7	Protein max	7,67
TRINITY_DN26026_c1_g1_i2	E3 ubiquitin-protein ligase RNF126	7,67
TRINITY_DN31511_c0_g1_i9	Protein AF-10	7,67
TRINITY_DN29882_c1_g1_i13		7,67
TRINITY_DN30273_c0_g1_i8	ADP-ribosylation factor GTPase-activating protein 3	
	Rab11 family-interacting protein 4B	7,67
TRINITY_DN30359_c3_g1_i2	RelA-associated inhibitor	7,67
TRINITY_DN31025_c0_g2_i2	Casein kinase I isoform delta-B	7,67
TRINITY_DN31231_c0_g1_i5	DNA-binding death effector domain-containing protein 2	7,67
TRINITY_DN33172_c0_g1_i2	Elongation of very long chain fatty acids protein 2	7,67
TRINITY_DN28524_c0_g2_i6	Heterogeneous nuclear ribonucleoprotein K	7,67
TRINITY_DN31677_c2_g1_i18	Solute carrier family 25 member 47-B	7,67
TRINITY_DN32965_c0_g1_i1	N-lysine methyltransferase setd6	7,67

TRINITY DN27884_c g jo	TRINITY_DN27834_c1_g1_i9	Mussin 10	7,67
TRINITY_DN2938_c2_g1_i10			
TRINITY DN2855, c0. g. 1.31 Vacaolar protein sorting-associated protein 18 homolog 7.67 TRINITY DN2865, c0. g. 2.118 Protein rapianzel 7.67 TRINITY DN2856, c0. g. 2.118 Protein rapianzel Protein RRX1 homolog 7.67 TRINITY DN2856, c0. g. 2.118 Protein in-7 homolog 7.67 TRINITY DN29730, c0. g. 1.51 Dilujaritin-associated dehalogenase-like hydrolase domain-containing 5 7.67 TRINITY DN30099, c1. g. 1.14 Dilujaritin-associated domain-containing force in 1 TRINITY DN29090, c2. g. g. 1.14 Dilujaritin-associated domain-containing protein 1 7.67 TRINITY DN29688, c0. g. 1.24 Dilujaritin-associated domain-containing protein 1 7.67 TRINITY DN29384, c0. g. 1.25 DIS3-like exonuclease 2 7.67 DIS3-like exonuclease 3 Transcription initiation factor FFIID submit 1 7.67 TRINITY DN28958, c0. g. 2. 12 TRINITY DN29383, c2. g. 1.25 Calcitoni-1 TRINITY DN29383, c2. g. 1.25 TRINITY DN29383, c2. g. 1.25 Calcitoni-1 TRINITY DN293834, c2. g. 1.25 Calcitoni-1 TRINITY DN293834, c2. g. 1.25 Calcitoni-1 TRINITY DN293	_	•	
TRINITY DN28616_cl_gl_115			
TRINITY DN2986, c3, g1, 118	_		
RINITY_DN29560_c0_g1_15	_	*	
REINITY_DN2590_0_0_gl_i2		<u>*</u>	
TRINITY DN30302	_		
TRINITY_DN30095_c1_g1_i1		•	
TRINITY_DN30995_cl_gl_i4	_	· · · · · · · · · · · · · · · · · · ·	
TRINITY_DN29202_c2_g1_i4	_		
TRINITY_DN27907_cl_gl_i4	•		
TRINITY_DN24384_c0_g1_id		•	
TRINITY_DN2968_c0_g1_i1 DIS3-like exonuclease 2 7,67 TRINITY_DN218417_c0_g2_i4 Aryl hydrocarbon receptor nuclear translocator 7,72 TRINITY_DN22844_c0_g1_i2 Opioid growth factor receptor 7,72 TRINITY_DN29825_c1_g2_j18 [Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme 7,72 TRINITY_DN29383_c2_g1_i5 Carnitine O-acetyltransferase NDUFAF7, mitochondrial 7,72 TRINITY_DN29383_c2_g1_i5 Carnitine O-acetyltransferase 7,72 TRINITY_DN30119_c1_g1_i2 Adenosine 3'-phospho 5'-phosphosulfate transporter 1 7,72 TRINITY_DN32979_c2_g3_i6 A-kinase anchor protein 7 isoform alpha 7,72 TRINITY_DN32979_c2_g3_i6 A-kinase anchor protein 7 isoform alpha 7,72 TRINITY_DN32979_c2_g3_i6 LiVer carboxylesterase 2 7,72 TRINITY_DN32915_c0_g1_i10 LVR moil-containing protein 5B 7,72 TRINITY_DN32013_c1_g2_i2 Liver carboxylesterase 2 7,72 TRINITY_DN3168_c1_g1_i3 Relophillin-1 7,72 TRINITY_DN3208_c1_g1_i4 CD2-associated protein 87A 7,72 TRINITY_DN3208_c1_g1_i4 CD2-associated protein 87A 7,72			
TRINITY_DN23181_cq_2_i2_1	•	•	
TRINITY_DN32412_0_gl_i2	_		
TRINITY_DN28948_c0_g1_i2	_		
TRINITY_DN30752_c0_g1_i2	_		
TRINITY_DN29383_c_g_g_i5	_		
TRINITY_DN29383_c2_g1_i5	_	[Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme	
TRINITY_DN22780_c0_g1_i2	•	·	
TRINITY_DN30119_c1_g1_i2	TRINITY_DN29383_c2_g1_i5	Carnitine O-acetyltransferase	7,72
TRINITY_DN29702_c1_g3_i6	TRINITY_DN22780_c0_g1_i5	Calcitonin-1	7,72
TRINITY_DN32979_c2_g3_i6	TRINITY_DN30119_c1_g1_i2	Adenosine 3'-phospho 5'-phosphosulfate transporter 1	7,72
TRINITY_DN32520_c0_g1_i2 Nipped-B-like protein A 7,72 TRINITY_DN27958_c0_g1_i10 LYR motif-containing protein 5B 7,72 TRINITY_DN30113_c1_g2_i2 Liver carboxylesterase 2 7,72 TRINITY_DN28290_c0_g1_i19 Cytochrome P450_2J2 7,72 TRINITY_DN24672_c4_g2_i4 Rhophilin-1 7,72 TRINITY_DN31686_c1_g1_i4 Transmembrane protein 87A 7,72 TRINITY_DN3389_c0_g1_i21 CD2-associated protein 7,72 TRINITY_DN30685_c0_g1_i4 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN32049_c0_g2_i1 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN32049_c0_g2_i1 Nucleolar pre-ribosomal-associated protein 1 7,72 TRINITY_DN32049_c0_g1_i5 Integrin alpha-E 7,72 TRINITY_DN2499_c1_g1_i9 Retinoic acid receptor RRR-beta-A 7,72 TRINITY_DN2499_c1_g1_i1 Selenoprotein Pb 7,72 TRINITY_DN32852_c1_g1_i1 Selenoprotein Pb 7,72 TRINITY_DN32895_c1_g1_i2 Potassium channel subfamily K member 4 7,72 TRINITY_DN2895_c1_g1_i5 Uncharacterized protein C7orf50	TRINITY_DN29702_c1_g3_i2	DNA/RNA-binding protein KIN17	7,72
TRINITY_DN32958_c0_g1_i10	TRINITY_DN32979_c2_g3_i6	A-kinase anchor protein 7 isoform alpha	7,72
TRINITY_DN30113_cl_g2_i2 Liver carboxylesterase 2 7,72 TRINITY_DN28290_c0_gl_i19 Cytochrome P450_2J2 7,72 TRINITY_DN24672_c4_g2_i4 Rhophilin-1 7,72 TRINITY_DN29154_cl_gl_i3 Sentrin-specific protease 5 7,72 TRINITY_DN31686_cl_gl_i4 Transmembrane protein 87A 7,72 TRINITY_DN23389_c0_gl_i21 CD2-associated protein 7,72 TRINITY_DN2969s_cl_gl_i40 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN30865_c0_gl_i4 Cytochrome P450_ZKI 7,72 TRINITY_DN32049_c0_g2_i1 Nucleolar pre-ribosomal-associated protein I 7,72 TRINITY_DN23701_c0_gl_i5 Integrin alpha-E 7,72 TRINITY_DN24996_cl_gl_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN29488_c0_gl_i2 T-complex protein II-like protein I 7,72 TRINITY_DN32885_cl_gl_i11 Selenoprotein Pb 7,72 TRINITY_DN24895_cl_gl_i8 Cryptochrome-1 7,72 TRINITY_DN22885_cl_gl_i11 Activin receptor type-1 7,72 TRINITY_DN26985_c0_gl_i9 Nuclear receptor type-1 7,72	TRINITY_DN32520_c0_g1_i2	Nipped-B-like protein A	7,72
TRINITY_DN30113_cl_g2_i2 Liver carboxylesterase 2 7,72 TRINITY_DN28290_c0_gl_i19 Cytochrome P450_2J2 7,72 TRINITY_DN24672_c4_g2_i4 Rhophilin-1 7,72 TRINITY_DN29154_cl_gl_i3 Sentrin-specific protease 5 7,72 TRINITY_DN31686_cl_gl_i4 Transmembrane protein 87A 7,72 TRINITY_DN23389_c0_gl_i21 CD2-associated protein 7,72 TRINITY_DN2969s_cl_gl_i40 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN30865_c0_gl_i4 Cytochrome P450_ZKI 7,72 TRINITY_DN32049_c0_g2_i1 Nucleolar pre-ribosomal-associated protein I 7,72 TRINITY_DN23701_c0_gl_i5 Integrin alpha-E 7,72 TRINITY_DN24996_cl_gl_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN29488_c0_gl_i2 T-complex protein II-like protein I 7,72 TRINITY_DN32885_cl_gl_i11 Selenoprotein Pb 7,72 TRINITY_DN24895_cl_gl_i8 Cryptochrome-1 7,72 TRINITY_DN22885_cl_gl_i11 Activin receptor type-1 7,72 TRINITY_DN26985_c0_gl_i9 Nuclear receptor type-1 7,72	TRINITY_DN27958_c0_g1_i10	LYR motif-containing protein 5B	7,72
TRINITY_DN24672_c4_g2_i4 Rhophilin-1 7,72 TRINITY_DN29154_c1_g1_i3 Sentrin-specific protease 5 7,72 TRINITY_DN31686_c1_g1_i4 Transmembrane protein 87A 7,72 TRINITY_DN29698_c1_g1_i40 CD2-associated protein 7,72 TRINITY_DN30685_c0_g1_i4 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN32049_c0_g2_i1 Nucleolar pre-ribosomal-associated protein 1 7,72 TRINITY_DN23701_c0_g1_i5 Integrin alpha-E 7,72 TRINITY_DN23884_c1_g2_i5 Deoxynucleotidyltransferase terminal-interacting protein 1 7,72 TRINITY_DN24996_c1_g1_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN24995_c1_g1_i17 Selenoprotein Pb 7,72 TRINITY_DN24895_c1_g1_i18 Cryptochrome-1 7,72 TRINITY_DN28895_c1_g1_i1 Activin receptor type-1 7,72 TRINITY_DN26985_c0_g1_i9 Nuclear receptor 2C2-associated protein 7,72 TRINITY_DN31594_c0_g2_i2 Signal transducing adapter molecule 1 7,72 TRINITY_DN325375_c0_g1_i6 CD9 antigen 7,72 TRINITY_DN32626_c1_g1_i6 CD9 antigen 7,	TRINITY_DN30113_c1_g2_i2		7,72
TRINITY_DN24672_c4_g2_i4 Rhophilin-1 7,72 TRINITY_DN29154_c1_g1_i3 Sentrin-specific protease 5 7,72 TRINITY_DN31686_c1_g1_i4 Transmembrane protein 87A 7,72 TRINITY_DN29698_c1_g1_i40 CD2-associated protein 7,72 TRINITY_DN30685_c0_g1_i4 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN32049_c0_g2_i1 Nucleolar pre-ribosomal-associated protein 1 7,72 TRINITY_DN23701_c0_g1_i5 Integrin alpha-E 7,72 TRINITY_DN23884_c1_g2_i5 Deoxynucleotidyltransferase terminal-interacting protein 1 7,72 TRINITY_DN24996_c1_g1_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN24995_c1_g1_i17 Selenoprotein Pb 7,72 TRINITY_DN24895_c1_g1_i18 Cryptochrome-1 7,72 TRINITY_DN28895_c1_g1_i1 Activin receptor type-1 7,72 TRINITY_DN26985_c0_g1_i9 Nuclear receptor 2C2-associated protein 7,72 TRINITY_DN31594_c0_g2_i2 Signal transducing adapter molecule 1 7,72 TRINITY_DN325375_c0_g1_i6 CD9 antigen 7,72 TRINITY_DN32626_c1_g1_i6 CD9 antigen 7,	TRINITY_DN28290_c0_g1_i19	Cytochrome P450 2J2	7,72
TRINITY_DN2154_cl_gl_i3 Sentrin-specific protease 5 7,72 TRINITY_DN21686_cl_gl_i4 Transmembrane protein 87A 7,72 TRINITY_DN23389_cl_gl_i21 CD2-associated protein 7,72 TRINITY_DN29698_cl_gl_i40 H-2 class II histocompatibility antigen, I-E beta chain 7,72 TRINITY_DN30685_cl_gl_i4 Cytochrome P450 2K1 7,72 TRINITY_DN23701_cl_gl_i5 Integrin alpha-E 7,72 TRINITY_DN23701_cl_gl_i5 Integrin alpha-E 7,72 TRINITY_DN24996_cl_gl_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN27948_cl_gl_i9 Retinoic acid receptor RXR-beta-A 7,72 TRINITY_DN32852_cl_gl_i17 Selenoprotein II-like protein I 7,72 TRINITY_DN32895_cl_gl_i2 Potassium channel subfamily K member 4 7,72 TRINITY_DN32934_cl_gl_i8 Cryptochrome-1 7,72 TRINITY_DN22895_cl_gl_i11 Activin receptor type-1 7,72 TRINITY_DN2895_cl_gl_i1 Activin receptor type-1 7,72 TRINITY_DN31594_cl_g2_i2 Signal transducing adapter molecule 1 7,72 TRINITY_DN30599_cl_gl_i5 Probable ATT-dependent RNA helicase DHX35	_	Rhophilin-1	7,72
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	1KIN11 1_DIN29/04_CU_g1_13	AMF deaminase 2	1,72

TRINITY_DN26721_c3_g1_i3	Probable C->U-editing enzyme APOBEC-2	7,72
TRINITY_DN30722_c1_g1_i5	Receptor-type tyrosine-protein phosphatase gamma	7,72
TRINITY_DN28821_c1_g1_i2	Unconventional myosin-Vb	7,72
TRINITY_DN23836_c0_g1_i1	TM2 domain-containing protein 2	7,72
TRINITY_DN28788_c2_g2_i1	RCC1-like G exchanging factor-like protein	7,72
TRINITY_DN23146_c0_g1_i7	Protein phosphatase 1 regulatory subunit 12A	7,72
TRINITY_DN25060_c1_g1_i4	Vasodilator-stimulated phosphoprotein	7,72
TRINITY_DN28728_c1_g1_i7	Peptidyl-prolyl cis-trans isomerase FKBP8	7,72
TRINITY_DN24807_c1_g1_i6	Integrator complex subunit 4	7,72
TRINITY_DN29826_c0_g1_i3	Cyclin-I	7,72
TRINITY_DN22358_c0_g1_i3	Corticosteroid 11-beta-dehydrogenase isozyme 2	7,72
TRINITY_DN32861_c1_g1_i4	E3 ubiquitin-protein ligase MARCH6	7,72
TRINITY_DN31931_c1_g1_i16	Guanine nucleotide exchange factor DBS	7,72
TRINITY_DN29515_c0_g1_i4	14-3-3 protein theta	7,72
TRINITY_DN30759_c0_g2_i11	Malate synthase	7,72
TRINITY_DN30396_c0_g1_i2	COMM domain-containing protein 7	7,72
TRINITY_DN31172_c0_g1_i26	OTU domain-containing protein 3	7,72
TRINITY_DN28881_c0_g1_i4	Protein pelota homolog	7,72
TRINITY_DN30800_c1_g4_i8	Telomeric repeat-binding factor 2	7,72
TRINITY_DN30751_c0_g2_i2	F-box/LRR-repeat protein 20	7,72
TRINITY_DN26347_c0_g1_i4	SH3 domain-binding glutamic acid-rich-like protein 3	7,72
TRINITY_DN24691_c3_g2_i3	Septin-7 (Fragment)	7,72
TRINITY_DN30952_c0_g1_i16	Microtubule-associated protein 4	7,72
TRINITY_DN28415_c0_g1_i8	Small leucine-rich protein 1	7,72
TRINITY_DN32572_c1_g1_i15	N-lysine methyltransferase KMT5A	7,72
TRINITY_DN25244_c0_g1_i7	Protein canopy-1	7,72
TRINITY_DN28145_c1_g1_i4	UBX domain-containing protein 6	7,72
TRINITY_DN28242_c1_g1_i14	Inactive tyrosine-protein kinase PEAK1	7,72
TRINITY_DN31369_c0_g1_i4	Pleckstrin homology-like domain family B member 2	7,72
TRINITY_DN29350_c2_g1_i3	Palmitoyltransferase ZDHHC2	7,72
TRINITY_DN32440_c1_g2_i5	NAD kinase	7,72
TRINITY_DN32093_c3_g2_i16	Cytohesin-2	7,72
TRINITY_DN27624_c0_g1_i2	RNA polymerase-associated protein RTF1 homolog	7,72
TRINITY_DN29783_c1_g2_i7	Protein disulfide-isomerase A6	7,78
TRINITY_DN22442_c0_g1_i1	Dysferlin	7,78
TRINITY_DN26861_c0_g1_i5	Methyltransferase-like protein 25	7,78
TRINITY_DN26143_c0_g1_i5	CCAAT/enhancer-binding protein zeta	7,78
TRINITY_DN29906_c0_g2_i8		7,78
_	Uncharacterized protein C11orf24 homolog	
TRINITY_DN23660_c2_g1_i9	F-actin-capping protein subunit alpha-1	7,78
TRINITY_DN25399_c2_g2_i9	Ephrin type-B receptor 4	7,78
TRINITY_DN25147_c0_g1_i5	Transmembrane protein 179	7,78
TRINITY_DN28140_c1_g1_i21	Target of Myb protein 1	7,78
TRINITY_DN25994_c3_g1_i9	Protein FAM126B	7,78
TRINITY_DN27546_c0_g1_i6	GEM-interacting protein	7,78
TRINITY_DN29146_c0_g1_i1	Apoptosis-inducing factor 1, mitochondrial	7,78
TRINITY_DN22430_c0_g1_i4	Protein DBF4 homolog A	7,78
TRINITY_DN27326_c3_g1_i2	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	7,78
TRINITY_DN27949_c1_g1_i2	Calpain-1 catalytic subunit	7,78
TRINITY_DN30751_c0_g2_i6		
	F-box/LRR-repeat protein 20	7,78
TRINITY_DN25507_c0_g1_i5	Uridine diphosphate glucose pyrophosphatase	7,78
TRINITY_DN31743_c2_g2_i3	Latent-transforming growth factor beta-binding protein 1	7,78
TRINITY_DN31437_c1_g6_i2	Estrogen receptor beta-2	7,78
TRINITY_DN29168_c0_g2_i8	Calpain-7	7,78
TRINITY_DN29458_c0_g1_i8	Phosphatidate phosphatase LPIN1	7,78
TRINITY_DN31250_c0_g1_i2	Coiled-coil-helix-coiled-coil-helix domain-containing protein 10	7,78
TRINITY_DN28066_c0_g1_i13	Regulatory factor X-associated protein	7,78
TRINITY_DN28165_c2_g1_i2	Ribosomal protein S6 kinase alpha-3	7,78
TRINITY_DN23410_c0_g1_i1	Biogenesis of lysosome-related organelles complex 1 subunit 2	7,78
TRINITY_DN24598_c0_g1_i14	ATPase family AAA domain-containing protein 2B	7,78
111111111111111111111111111111111111111	one james, that adman community protein 20	,,,,

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TRINITY_DN30416_c0_g1_i8	RNA binding protein fox-1 homolog 1-like	7,78
TRINITY_DN29840_c1_g1_i3	Segment polarity protein dishevelled homolog DVL-3	7,78
TRINITY_DN27636_c0_g1_i4	Programmed cell death protein 4	7,78
TRINITY_DN32152_c0_g2_i5	Microtubule-associated tumor suppressor 1 homolog A	7,78
TRINITY_DN21588_c0_g1_i1	TBC1 domain family member 1	7,78
TRINITY_DN31797_c0_g1_i1	Phosphatase and actin regulator 4A	7,78
TRINITY_DN30195_c1_g2_i2	Sphingomyelin phosphodiesterase 4	7,78
TRINITY_DN31314_c1_g1_i13	Retinol dehydrogenase 8	7,78
TRINITY_DN28763_c0_g3_i16	Cullin-1	7,78
TRINITY_DN26102_c2_g1_i6	Leucine-zipper-like transcriptional regulator 1	7,78
TRINITY_DN33131_c2_g2_i3	Triple functional domain protein	7,78
TRINITY_DN28753_c0_g1_i4	E3 ubiquitin-protein ligase RNF34	7,78
TRINITY_DN33036_c1_g1_i6	2-amino-3-carboxymuconate-6-semialdehyde decarboxylase	7,78
TRINITY_DN23679_c7_g2_i2	NXPE family member 3	7,78
TRINITY_DN30184_c0_g1_i4	Bone morphogenetic protein receptor type-2	7,78
TRINITY_DN28208_c0_g1_i7	RAD50-interacting protein 1	7,78
TRINITY_DN30115_c0_g1_i2	Vacuolar protein sorting-associated protein 41 homolog	7,78
	Kinase suppressor of Ras 1	7,78
TRINITY_DN29130_c0_g3_i4		
TRINITY_DN32840_c3_g1_i1	Hypoxia up-regulated protein 1	7,78
TRINITY_DN28234_c2_g1_i19	Septin-4	7,78
TRINITY_DN29104_c2_g1_i22	Casein kinase I isoform gamma-3	7,78
TRINITY_DN25589_c1_g1_i9	Chitinase domain-containing protein 1	7,78
TRINITY_DN30839_c0_g1_i2	E3 ubiquitin-protein ligase pellino homolog 1	7,78
TRINITY_DN28440_c0_g2_i7	Enhancer of polycomb homolog 1	7,78
TRINITY_DN28754_c0_g2_i7	Aspartyl aminopeptidase	7,78
TRINITY_DN26657_c0_g1_i9	Phosphofurin acidic cluster sorting protein 2	7,78
TRINITY_DN29239_c0_g1_i9	Ubiquitin-conjugating enzyme E2 variant 3	7,78
TRINITY_DN25639_c1_g2_i6	Kazrin	7,78
TRINITY_DN32919_c4_g2_i1	Glutaminase liver isoform, mitochondrial	7,78
TRINITY_DN23479_c0_g1_i3	NADH-cytochrome b5 reductase 1	7,78
TRINITY_DN31281_c0_g1_i12	RNA-directed DNA polymerase from mobile element jockey	7,78
TRINITY_DN26173_c0_g1_i3	Mothers against decapentaplegic homolog 5	7,78
TRINITY_DN31907_c0_g2_i11	Protein kinase C and casein kinase substrate in neurons protein	7,78
TRINITY_DN23810_c0_g1_i11	Magnesium transporter NIPA2	7,78
TRINITY_DN26276_c0_g1_i3	Transcription factor E4F1	7,78
TRINITY_DN27289_c1_g1_i7	Aryl hydrocarbon receptor nuclear translocator-like protein 2	7,78
TRINITY_DN28658_c0_g1_i6	Insulin-induced gene 1 protein	7,78
TRINITY_DN30862_c0_g1_i11	ATP-dependent RNA helicase DDX19A	7,78
TRINITY_DN30365_c1_g2_i10	High mobility group protein 20A	7,78
TRINITY_DN24336_c0_g1_i9	Ras-related protein Rab-1A	7,78
TRINITY_DN31703_c0_g1_i8	3 beta-hydroxysteroid dehydrogenase type 7	7,78
TRINITY_DN31878_c2_g1_i7	HEAT repeat-containing protein 5B	7,78
TRINITY_DN26959_c1_g1_i2	NADPHcytochrome P450 reductase	7,78
TRINITY_DN32195_c2_g2_i5	GTPase-activating protein and VPS9 domain-containing protein	7,78
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TRINITY_DN32435_c3_g1_i27	NADPH:adrenodoxin oxidoreductase, mitochondrial	7,78
TRINITY_DN25395_c0_g1_i2	Catenin beta-1	7,78
TRINITY_DN28524_c0_g2_i9	Heterogeneous nuclear ribonucleoprotein K	7,78
TRINITY_DN31272_c0_g1_i20	Ubiquitin-protein ligase E3B	7,83
TRINITY_DN31527_c0_g1_i9	Rho GTPase-activating protein SYDE2	7,83
TRINITY_DN26119_c1_g2_i3	Protein SDA1 homolog	7,83
TRINITY_DN31523_c0_g1_i8	Phosphorylase b kinase regulatory subunit alpha, skeletal musc	7,83
TRINITY_DN32412_c3_g4_i12	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	7,83
TRINITY_DN28491_c0_g2_i3	Discs large homolog 1-like protein	7,83
TRINITY_DN27540_c4_g1_i9	Saccharopine dehydrogenase-like oxidoreductase	7,83
TRINITY_DN33058_c1_g1_i3	H-2 class II histocompatibility antigen, E-B beta chain	7,83
TRINITY_DN32486_c0_g1_i8	Serine/threonine-protein kinase Nek9	7,83
TRINITY_DN31637_c0_g2_i10	Neurofibromin	7,83
TRINITY_DN27546_c0_g1_i2	GEM-interacting protein	7,83
TRINITY_DN31440_c0_g1_i5	Rab3 GTPase-activating protein non-catalytic subunit	7,83
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TDINITY DN05701 .1 .0 111		7.02
TRINITY_DN25781_c1_g2_i11	Casein kinase II subunit alpha'	7,83
TRINITY_DN22958_c0_g1_i1	Abl interactor 1	7,83
TRINITY_DN24922_c0_g1_i1	Protein crumbs homolog 2	7,83
TRINITY_DN31394_c2_g1_i11	Aftiphilin	7,83
TRINITY_DN28912_c2_g1_i12	26S proteasome non-ATPase regulatory subunit 11B	7,83
TRINITY_DN29665_c1_g4_i2	RanBP-type and C3HC4-type zinc finger-containing protein 1	7,83
TRINITY_DN29955_c0_g1_i9	XK-related protein 8	7,83
TRINITY_DN30117_c0_g1_i16	Syntaxin-5	7,83
TRINITY_DN26307_c2_g1_i4	Atypical kinase COQ8A, mitochondrial	7,83
TRINITY_DN27326_c3_g1_i6	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	7,83
TRINITY_DN29314_c1_g1_i5	Transcription elongation regulator 1	7,83
TRINITY_DN32102_c0_g1_i6	RUN and FYVE domain-containing protein 1	7,83
TRINITY_DN33110_c2_g2_i15	Myomegalin	7,83
TRINITY_DN27019_c1_g1_i13	Complement C3	7,83
TRINITY_DN29226_c0_g1_i6	Membrane-bound transcription factor site-2 protease	7,83
TRINITY_DN33154_c4_g2_i5	Alpha-1,3-mannosyl-glycoprotein 4-beta-N-acetylglucosaminyli	7,83
TRINITY_DN30504_c0_g1_i4	Nuclear pore complex protein Nup98-Nup96	7,83
TRINITY_DN31483_c0_g1_i18	Ras GTPase-activating protein 1	7,83
TRINITY_DN28570_c2_g5_i4	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IC	7,83
TRINITY_DN24281_c0_g1_i1	Spermidine synthase	7,83
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TRINITY_DN32665_c0_g1_i9	Immunoglobulin superfamily member 8	7,83
TRINITY_DN31052_c0_g3_i1	Retrovirus-related Pol polyprotein from transposon 297	7,83
TRINITY_DN28574_c0_g1_i10	Myocilin	7,83
TRINITY_DN29229_c0_g2_i30	Histidine ammonia-lyase	7,83
TRINITY_DN30183_c0_g1_i4	Enhancer of mRNA-decapping protein 3	7,83
TRINITY_DN29179_c2_g1_i5	TGF-beta receptor type-2	7,83
TRINITY_DN27940_c0_g2_i8	Complement C3 (Fragment)	7,83
TRINITY_DN28970_c0_g1_i7	Adiponectin receptor protein 1	7,83
TRINITY_DN30788_c0_g1_i16	Interferon regulatory factor 2-binding protein 2-B	7,83
TRINITY_DN28600_c0_g1_i10	Hepatocyte nuclear factor 1-beta-A	7,83
TRINITY_DN28105_c1_g1_i1	Afadin	7,83
TRINITY_DN22657_c0_g1_i6	Probable palmitoyltransferase ZDHHC4	7,83
TRINITY_DN32740_c0_g1_i5	Vitellogenin-2	7,83
TRINITY_DN31394_c2_g1_i3	Aftiphilin	7,83
TRINITY_DN31608_c1_g1_i4	Differentially expressed in FDCP 6 homolog	7,83
TRINITY_DN27441_c0_g2_i12	ELMO domain-containing protein 2	7,83
TRINITY_DN29261_c0_g1_i8	DNA excision repair protein ERCC-6	7,83
TRINITY_DN27581_c0_g3_i14	Adiponectin receptor protein 2	7,83
TRINITY_DN31787_c4_g2_i1	SUN domain-containing protein 1	7,83
TRINITY_DN28986_c2_g1_i5	Protein Shroom4	7,83
TRINITY_DN29216_c4_g1_i7	Major facilitator superfamily domain-containing protein 9	7,88
TRINITY_DN30547_c0_g1_i7	Nuclear protein 1	7,88
TRINITY_DN30062_c1_g3_i4	•	7,88
	Disheveled-associated activator of morphogenesis 2	
TRINITY_DN33318_c5_g2_i3	Chloride transport protein 6	7,88
TRINITY_DN26712_c0_g1_i1	Transcription factor Sp3	7,88
TRINITY_DN31058_c0_g1_i10	Ezrin	7,88
TRINITY_DN23358_c0_g1_i3	PHD finger protein 14	7,88
TRINITY_DN31694_c0_g3_i8	Histone acetyltransferase p300	7,88
TRINITY_DN31271_c2_g1_i11	Aldehyde dehydrogenase family 8 member A1	7,88
TRINITY_DN31272_c0_g1_i15	Ubiquitin-protein ligase E3B	7,88
TRINITY_DN31710_c0_g1_i6	Ectonucleoside triphosphate diphosphohydrolase 1	7,88
TRINITY_DN26847_c0_g1_i1	E3 ubiquitin-protein ligase synoviolin	7,88
TRINITY_DN30562_c1_g3_i5	Tubby-related protein 3	7,88
TRINITY_DN30247_c3_g1_i29	Tyrosine 3-monooxygenase	7,88
TRINITY_DN28179_c1_g1_i16	Poly(A) polymerase gamma	7,88
TRINITY_DN27727_c0_g1_i1	Mediator of RNA polymerase II transcription subunit 16	7,88
TRINITY_DN32734_c1_g1_i3	Myosin-10	7,88
TRINITY_DN25624_c0_g1_i5	Beta-1,3-N-acetylglucosaminyltransferase radical fringe	7,88
TRINITY_DN27070_c0_g1_i13	Glutamine amidotransferase-like class 1 domain-containing pro	7,88
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TRINITY DNOSES 1 1 1 10		7.00
TRINITY_DN28551_c1_g1_i9	Nuclear receptor coactivator 6	7,88
TRINITY_DN28371_c1_g2_i12	Sodium- and chloride-dependent creatine transporter 1	7,88
TRINITY_DN26864_c1_g1_i3	Coiled-coil domain-containing protein 134	7,88
TRINITY_DN32171_c0_g2_i7	E3 ubiquitin-protein ligase ZNF598	7,88
TRINITY_DN30783_c0_g1_i3	Probable helicase senataxin	7,88
TRINITY_DN28082_c0_g3_i10	Hemicentin-1	7,88
TRINITY_DN30247_c3_g1_i28	Tyrosine 3-monooxygenase	7,88
TRINITY_DN29168_c0_g2_i3	Calpain-7	7,88
TRINITY_DN31424_c0_g1_i7	FAST kinase domain-containing protein 4	7,88
TRINITY_DN25894_c0_g1_i5	Noelin	7,88
TRINITY_DN29383_c2_g1_i4	Carnitine O-acetyltransferase	7,88
TRINITY_DN26900_c2_g3_i1	Transmembrane protein 104	7,88
TRINITY_DN32377_c1_g1_i1	Glycerol-3-phosphate acyltransferase 1, mitochondrial	7,88
TRINITY_DN31874_c0_g1_i14	Peroxiredoxin-5, mitochondrial	7,88
TRINITY_DN29118_c0_g1_i11	Centrosomal protein of 63 kDa	7,88
TRINITY_DN27487_c2_g2_i1	Probable RNA polymerase II nuclear localization protein SLC7.	7,88
TRINITY_DN28320_c3_g1_i5	Ectonucleoside triphosphate diphosphohydrolase 5	7,88
TRINITY_DN29565_c1_g1_i12	Mucolipin-1	7,88
TRINITY_DN31850_c1_g3_i5	Glycerol-3-phosphate acyltransferase 4	7,88
TRINITY_DN33149_c1_g1_i6	Sodium-coupled neutral amino acid transporter 3	7,88
TRINITY_DN26287_c0_g1_i2	Multifunctional protein ADE2	7,88
TRINITY_DN32311_c1_g1_i3	Histone deacetylase 4	7,88
	•	7,88 7,88
TRINITY_DN28370_c2_g1_i4	U3 small nucleolar RNA-associated protein 18 homolog	
TRINITY_DN20691_c0_g1_i1	Sialoadhesin	7,88
TRINITY_DN25663_c0_g4_i1	Ran guanine nucleotide release factor	7,88
TRINITY_DN29027_c1_g2_i16	HIRA-interacting protein 3	7,88
TRINITY_DN27972_c1_g1_i11	Disintegrin and metalloproteinase domain-containing protein 1	7,88
TRINITY_DN31949_c0_g1_i5	Storkhead-box protein 2	7,88
TRINITY_DN30819_c2_g1_i3	Dual specificity mitogen-activated protein kinase kinase 7	7,88
TRINITY_DN29483_c1_g1_i2	Torsin-2A	7,88
TRINITY_DN31719_c0_g2_i3	ADP-ribosylation factor 6	7,88
TRINITY_DN32109_c0_g1_i9	Mitochondrial dicarboxylate carrier	7,88
TRINITY_DN31424_c0_g1_i6	FAST kinase domain-containing protein 4	7,88
TRINITY_DN32300_c1_g1_i7	Nuclear pore complex protein Nup153	7,88
TRINITY_DN26432_c0_g1_i2	DnaJ homolog subfamily C member 3	7,88
TRINITY_DN31080_c0_g1_i5	5'-AMP-activated protein kinase subunit gamma-1	7,88
TRINITY_DN28384_c3_g2_i3	5-formyltetrahydrofolate cyclo-ligase	7,88
TRINITY_DN29778_c1_g1_i3	Zinc finger and SCAN domain-containing protein 20	7,88
TRINITY_DN29726_c3_g1_i4	Aldehyde dehydrogenase, mitochondrial	7,88
TRINITY_DN26383_c1_g1_i16	Complement C3 (Fragment)	7,88
TRINITY_DN24716_c1_g1_i4	Zinc finger protein 622	7,88
TRINITY_DN27998_c0_g1_i15	IGF-like family receptor 1	7,88
TRINITY_DN32681_c1_g1_i10	Amine sulfotransferase	7,88
TRINITY_DN30147_c1_g1_i4	Amyloid-beta A4 protein	7,92
TRINITY_DN27241_c0_g1_i3	Mitochondrial basic amino acids transporter	7,92
TRINITY_DN26926_c3_g1_i11	Afadin- and alpha-actinin-binding protein	7,92
TRINITY_DN31354_c1_g1_i14	Synaptotagmin-like protein 2	7,92
TRINITY_DN25350_c2_g1_i6	Mitochondrial intermembrane space import and assembly prote	7,92
TRINITY_DN22862_c0_g1_i5	SET-binding protein	7,92
TRINITY_DN32611_c1_g2_i3	Death-inducer obliterator 1	7,92
TRINITY_DN31938_c1_g1_i12	Insulin receptor substrate 1-B Partitioning defective 3 homolog	7,92
TRINITY_DN29921_c0_g1_i9	Partitioning defective 3 homolog	7,92
TRINITY_DN29092_c0_g1_i4	F-box/WD repeat-containing protein pof1	7,92
TRINITY_DN24309_c0_g1_i2	GATOR complex protein NPRL2	7,92
TRINITY_DN26853_c1_g1_i10	DNA-directed RNA polymerase III subunit RPC8	7,92
TRINITY_DN31547_c5_g3_i2	EH domain-containing protein 3	7,92
TRINITY_DN30952_c0_g1_i1	Microtubule-associated protein 4	7,92
TRINITY_DN27079_c4_g1_i2	FERM domain-containing protein 6	7,92
TRINITY_DN28528_c0_g1_i11	Ubiquitin carboxyl-terminal hydrolase 45	7,92

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TRINITY_DN30893_c3_g1_i1	Digestive cysteine proteinase 2	7,92
TRINITY_DN30610_c0_g1_i8	Mitochondrial fission regulator 1	7,92
TRINITY_DN31710_c0_g4_i2	Sorting nexin-17	7,92
TRINITY_DN33285_c3_g1_i2	Death-associated protein kinase 3	7,92
TRINITY_DN23774_c0_g1_i7	Stress-associated endoplasmic reticulum protein 1	7,92
TRINITY_DN32533_c0_g2_i1	Nectin-2	7,92
TRINITY_DN33285_c3_g1_i1	Death-associated protein kinase 3	7,92
TRINITY_DN32283_c0_g2_i5	Hemagglutinin/amebocyte aggregation factor	7,92
TRINITY_DN28302_c1_g1_i4	General transcription factor IIH subunit 1	7,92
TRINITY_DN30463_c0_g1_i9	Peroxisome proliferator-activated receptor delta	7,92
TRINITY_DN30982_c0_g1_i25	Acyl-coenzyme A thioesterase 9, mitochondrial	7,92
TRINITY_DN31995_c0_g1_i6	Beta-2-syntrophin	7,92
TRINITY_DN30297_c2_g1_i3	Pterin-4-alpha-carbinolamine dehydratase	7,92
TRINITY_DN30467_c0_g1_i2	Cysteine-rich with EGF-like domain protein 2	7,92
TRINITY_DN25852_c7_g2_i12	Adhesion G-protein coupled receptor F1	7,92
TRINITY_DN29875_c2_g1_i9	Fumarylacetoacetase	7,92
TRINITY_DN22983_c1_g1_i6	Septin-8-A	7,92
TRINITY_DN32493_c0_g1_i24	Carboxylesterase 5A	7,92
TRINITY_DN29496_c2_g2_i4	ADP-ribose pyrophosphatase, mitochondrial	7,92
TRINITY_DN24515_c1_g3_i1	Ubiquitin-conjugating enzyme E2 D4	7,92
TRINITY_DN31510_c0_g1_i7	FYVE, RhoGEF and PH domain-containing protein 5	7,92
TRINITY_DN31695_c0_g1_i9	Vacuolar protein sorting-associated protein 4B	7,92
TRINITY_DN28613_c1_g1_i11	Zinc finger protein 226	7,92
TRINITY_DN25053_c0_g1_i8		7,92
_	SLAIN motif-containing protein 2 Short/household shain graphing applicable to the short share and the share are shared by the shared sh	
TRINITY_DN27961_c1_g1_i5	Short/branched chain specific acyl-CoA dehydrogenase, mitoch	7,92
TRINITY_DN23451_c0_g1_i3	V-type proton ATPase subunit G 1	7,92
TRINITY_DN30663_c3_g1_i12	Serine/threonine-protein kinase RIO1	7,92
TRINITY_DN32347_c3_g1_i9	Transposable element Tcb2 transposase	7,92
TRINITY_DN32087_c1_g1_i5	Early endosome antigen 1	7,92
TRINITY_DN26376_c0_g1_i6	Nucleolar transcription factor 1	7,92
TRINITY_DN29092_c0_g1_i8	Vegetative incompatibility protein HET-E-1	7,92
TRINITY_DN30843_c2_g2_i10	Histone-lysine N-methyltransferase NSD3	7,92
TRINITY_DN28492_c0_g4_i3	AP2-associated protein kinase 1	7,92
TRINITY_DN25197_c3_g1_i1	UDP-glucuronosyltransferase 2B16 (Fragment)	7,97
TRINITY_DN28749_c3_g1_i3	CCR4-NOT transcription complex subunit 3	7,97
TRINITY_DN28394_c0_g1_i6	RNA 3'-terminal phosphate cyclase	7,97
TRINITY_DN30862_c0_g1_i20	ATP-dependent RNA helicase DDX19A	7,97
TRINITY_DN29178_c2_g1_i2	Serine hydroxymethyltransferase, mitochondrial	7,97
TRINITY_DN29875_c2_g1_i11	Fumarylacetoacetase	7,97
TRINITY_DN25203_c2_g2_i3	Matrin-3	7,97
TRINITY_DN27654_c1_g2_i9	Mitochondrial carnitine/acylcarnitine carrier protein	7,97
TRINITY_DN26312_c1_g4_i6	Serine/threonine-protein phosphatase PP1-gamma catalytic suk	7,97
TRINITY_DN29302_c1_g2_i8	FGGY carbohydrate kinase domain-containing protein	7,97
TRINITY_DN25030_c1_g1_i11	Glutamate receptor 3	7,97
TRINITY_DN28803_c0_g1_i16	Solute carrier family 2, facilitated glucose transporter member !	7,97
TRINITY_DN30591_c0_g1_i12	Zinc finger protein 609	7,97
TRINITY_DN32223_c0_g3_i8	Solute carrier family 22 member 2	7,97
TRINITY_DN30110_c0_g1_i4	Ras-related protein Rab-33B	7,97
TRINITY_DN32574_c2_g1_i1	Prostaglandin reductase 2	7,97
TRINITY_DN28936_c0_g1_i7	Phosphatidate cytidylyltransferase 2	7,97
TRINITY_DN29830_c0_g1_i7	Carboxypeptidase B2	7,97
TRINITY_DN31417_c0_g2_i23	Aryl hydrocarbon receptor nuclear translocator	7,97
TRINITY_DN30017_c0_g1_i6	NFU1 iron-sulfur cluster scaffold homolog, mitochondrial	7,97 7,97
TRINITY_DN30921_c1_g2_i7		7,97 7,97
_	Unconventional myosin-X Saring/thyroning protein kingse Nek I	
TRINITY_DN29357_c2_g1_i5	Serine/threonine-protein kinase Nek1	7,97
TRINITY_DN31394_c2_g1_i12	Aftiphilin Mitsology dried discussions are a granical	7,97
TRINITY_DN32109_c0_g1_i26	Mitochondrial dicarboxylate carrier	7,97
TRINITY_DN32079_c1_g2_i3	Cation-independent mannose-6-phosphate receptor	7,97
TRINITY_DN22732_c0_g1_i8	NPC1-like intracellular cholesterol transporter 1	7,97

TRINITY_DN30305_c0_g1_i10	Cdc42 effector protein 4	7,97
TRINITY_DN27137_c0_g1_i9	DnaJ homolog subfamily B member 1	7,97
TRINITY_DN24859_c0_g1_i9	Ras-related protein Rab-5A	7,97
TRINITY_DN25602_c2_g2_i5	Fatty acid-binding protein, heart	7,97
TRINITY_DN32382_c0_g1_i11	Synergin gamma	7,97
TRINITY_DN32746_c2_g1_i1	Nuclear receptor corepressor 1	7,97
TRINITY_DN27692_c1_g1_i19	NAD-dependent protein deacylase sirtuin-5, mitochondrial	7,97
TRINITY_DN28012_c3_g1_i23	DnaJ homolog subfamily C member 16	7,97
TRINITY_DN30393_c0_g1_i3	RNA helicase aquarius	7,97
	<u>-</u>	
TRINITY_DN25944_c1_g1_i4	Peroxisome proliferator-activated receptor gamma coactivator-	7,97
TRINITY_DN26983_c0_g1_i13	Serine/arginine-rich splicing factor 11	7,97
TRINITY_DN30843_c2_g2_i3	Histone-lysine N-methyltransferase NSD3	7,97
TRINITY_DN28611_c2_g2_i14	DnaJ homolog subfamily C member 5	7,97
TRINITY_DN24101_c3_g2_i6	TSC22 domain family protein 3	7,97
TRINITY_DN30829_c1_g2_i1	Surfeit locus protein 4	7,97
TRINITY_DN17072_c0_g1_i1	Zinc finger homeobox protein 3	7,97
TRINITY_DN30342_c0_g1_i9	Helicase SKI2W	7,97
TRINITY_DN25400_c0_g1_i4	Rapamycin-insensitive companion of mTOR	7,97
TRINITY_DN30497_c1_g2_i7	Protein FAM89A	7,97
TRINITY_DN27318_c1_g1_i8	Leucine-rich repeat-containing protein 57	7,97
TRINITY_DN28458_c1_g1_i4	Macrophage receptor MARCO	7,97
TRINITY_DN31366_c0_g1_i7	Serine/threonine-protein phosphatase 4 catalytic subunit B	7,97
TRINITY_DN31052_c0_g1_i7	Retrovirus-related Pol polyprotein from transposon 297	7,97
TRINITY_DN29363_c0_g1_i2		7,97
	DnaJ homolog subfamily A member 3, mitochondrial	
TRINITY_DN31958_c0_g1_i10	Ubiquitin-associated protein 2-like	7,97
TRINITY_DN26226_c1_g2_i3	CD82 antigen	7,97
TRINITY_DN27637_c0_g2_i4	Serine-protein kinase ATM	7,97
TRINITY_DN29809_c0_g1_i2	Pogo transposable element with ZNF domain	7,97
TRINITY_DN7437_c0_g1_i1	Collagen alpha-1(XI) chain	8,01
TRINITY_DN32789_c0_g2_i17	2-oxoglutarate dehydrogenase, mitochondrial	8,01
TRINITY_DN21964_c0_g1_i1	RING finger protein 44	8,01
TRINITY_DN30677_c1_g3_i3	Long-chain fatty acid transport protein 1	8,01
TRINITY_DN28528_c0_g1_i8	Ubiquitin carboxyl-terminal hydrolase 45	8,01
TRINITY_DN22948_c0_g1_i7	Ras-related and estrogen-regulated growth inhibitor-like protei	8,01
TRINITY_DN25345_c0_g1_i7	TBC1 domain family member 16	8,01
TRINITY_DN32731_c0_g2_i1	Heterogeneous nuclear ribonucleoprotein C	8,01
TRINITY_DN24766_c0_g1_i3	Ellis-van Creveld syndrome protein homolog	8,01
TRINITY_DN29179_c2_g1_i4	TGF-beta receptor type-2	8,01
TRINITY_DN25367_c0_g1_i6	UDP-glucuronosyltransferase 1-1	8,01
_	Alpha-methylacyl-CoA racemase	
TRINITY_DN26832_c6_g1_i2	1 2 2	8,01
TRINITY_DN29499_c0_g1_i12	Dynein regulatory complex subunit 7	8,01
TRINITY_DN31378_c0_g1_i13	Cyclic AMP-dependent transcription factor ATF-6 beta	8,01
TRINITY_DN23869_c0_g1_i5	Rab-like protein 3	8,01
TRINITY_DN22825_c0_g1_i2	Adenosine kinase	8,01
TRINITY_DN30247_c3_g1_i6	Tyrosine 3-monooxygenase	8,01
TRINITY_DN28863_c0_g2_i1	Xylulose kinase	8,01
TRINITY_DN30641_c0_g1_i6	Uncharacterized protein KIAA1522 homolog	8,01
TRINITY_DN33020_c6_g1_i3	Acyl-CoA synthetase family member 2, mitochondrial	8,01
TRINITY_DN29945_c3_g3_i2	Fucolectin	8,01
TRINITY_DN32396_c0_g1_i6	Torsin-1B	8,01
TRINITY_DN25358_c0_g1_i3	E3 ubiquitin-protein ligase RNF115	8,01
TRINITY_DN32927_c1_g1_i7	Protein kinase C and casein kinase substrate in neurons protein	8,01
TRINITY_DN32128_c1_g1_i4	Peptidyl-glycine alpha-amidating monooxygenase A	8,01
TRINITY_DN31209_c0_g1_i7	Nicastrin	8,01
TRINITY_DN23933_c2_g2_i9	Surface protein	8,01
TRINITY_DN31249_c1_g2_i12	Transmembrane protease serine 6	8,01
_	_	
TRINITY_DN25655_c1_g1_i3	DnaJ homolog subfamily B member 9	8,01
TRINITY_DN28195_c0_g2_i6	Alpha-actinin-1	8,01
TRINITY_DN31205_c0_g1_i7	ATP-binding cassette sub-family A member 1	8,01

TRINITY_DN30490_c1_g3_i4	ATP-dependent Clp protease ATP-binding subunit clpX-like, mi	8,01
TRINITY_DN24751_c2_g1_i4	Formin-binding protein 1	8,01
TRINITY_DN27367_c1_g1_i1	Proprotein convertase subtilisin/kexin type 5	8,01
TRINITY_DN31081_c1_g1_i2	Phytanoyl-CoA dioxygenase domain-containing protein 1	8,01
TRINITY_DN31434_c3_g3_i1	Transcription termination factor 1a, mitochondrial	8,01
TRINITY_DN33033_c1_g1_i5	DmX-like protein 1	8,01
TRINITY_DN29118_c0_g1_i5	Centrosomal protein of 63 kDa	8,01
TRINITY_DN28585_c0_g2_i3	Enhancer of mRNA-decapping protein 4	8,01
TRINITY_DN31514_c1_g2_i7	G patch domain-containing protein 8	8,02
TRINITY_DN30175_c0_g1_i4	Enoyl-[acyl-carrier-protein] reductase, mitochondrial	8,06
TRINITY_DN29676_c1_g3_i9	ATP-dependent 6-phosphofructokinase, platelet type	8,06
TRINITY_DN31561_c2_g1_i5	Alanine aminotransferase 2-like	8,06
TRINITY_DN29765_c0_g1_i7	· ·	8,06
_	Splicing factor, arginine/serine-rich 15 (Fragment)	
TRINITY_DN25035_c3_g1_i7	Prolyl 4-hydroxylase subunit alpha-1	8,06
TRINITY_DN27267_c0_g2_i5	Mannan-binding lectin serine protease 2	8,06
TRINITY_DN31517_c0_g1_i3	Oxysterol-binding protein-related protein 9	8,06
TRINITY_DN26915_c0_g1_i1	RING finger protein 11	8,06
TRINITY_DN28933_c0_g1_i7	Sodium-independent sulfate anion transporter	8,06
TRINITY_DN26323_c2_g1_i1	Ubiquitin domain-containing protein UBFD1	8,06
TRINITY_DN25423_c0_g1_i6	Protein FAM114A2	8,06
TRINITY_DN23383_c0_g2_i11	Heterogeneous nuclear ribonucleoprotein A/B	8,06
TRINITY_DN32687_c1_g2_i3	Phosphatidylinositol 4-kinase alpha	8,06
TRINITY_DN26082_c0_g1_i1	60S ribosomal export protein NMD3	8,06
TRINITY_DN28048_c0_g2_i3	Probable E3 ubiquitin-protein ligase MID2	8,06
TRINITY_DN31702_c1_g1_i16	Uveal autoantigen with coiled-coil domains and ankyrin repeat:	8,06
TRINITY_DN27834_c1_g1_i5	Myosin-10	8,06
TRINITY_DN26336_c7_g2_i2	Chromodomain-helicase-DNA-binding protein 6	8,06
TRINITY_DN29242_c0_g1_i10	LON peptidase N-terminal domain and RING finger protein 1	8,06
TRINITY_DN28738_c0_g2_i6	Branched-chain-amino-acid aminotransferase, mitochondrial	8,06
TRINITY_DN32380_c0_g1_i9	Putative ATP-dependent RNA helicase an3	8,06
		8,06
TRINITY_DN33225_c5_g1_i9	Protein unc-93 homolog B1	
TRINITY_DN22470_c0_g1_i4	U6 snRNA-associated Sm-like protein LSm1	8,06
TRINITY_DN24508_c4_g2_i4	Acyl-protein thioesterase 1	8,06
TRINITY_DN30336_c1_g1_i6	Transcriptional regulator ATRX	8,06
TRINITY_DN27263_c0_g1_i1	Serine/threonine-protein kinase MARK2	8,06
TRINITY_DN30418_c0_g1_i1	Palmitoyltransferase ZDHHC9	8,06
TRINITY_DN28935_c2_g2_i11	Genetic suppressor element 1	8,06
TRINITY_DN29062_c0_g3_i4	Importin-8	8,06
TRINITY_DN23382_c0_g1_i6	Transmembrane protein 209	8,06
TRINITY_DN26637_c2_g1_i14	Polycomb group RING finger protein 5-A	8,06
TRINITY_DN21421_c0_g1_i3	Beta-1,3-glucosyltransferase	8,06
TRINITY_DN24533_c2_g4_i3	Vesicle-associated membrane protein-associated protein A	8,06
TRINITY_DN26809_c0_g1_i3	Protein kinase C iota type	8,06
TRINITY_DN33110_c2_g2_i14	Myomegalin	8,06
TRINITY_DN28516_c0_g1_i2	Autophagy protein 5	8,06
TRINITY_DN26917_c0_g1_i11	Transposon Ty3-I Gag-Pol polyprotein	8,06
TRINITY_DN28214_c0_g1_i12	RNA polymerase II subunit A C-terminal domain phosphatase	8,06
TRINITY_DN22878_c0_g1_i2		8,06
_	PDZ and LIM domain protein 4	
TRINITY_DN26794_c1_g2_i3	Epidermal growth factor-like protein 7	8,06
TRINITY_DN30132_c0_g1_i12	L-fucose kinase	8,06
TRINITY_DN20829_c0_g1_i3	Activated CDC42 kinase 1	8,06
TRINITY_DN31829_c1_g1_i7	Caskin-2	8,06
TRINITY_DN29117_c1_g2_i4	Cyclin-K	8,06
TRINITY_DN30110_c0_g1_i8	Ras-related protein Rab-33B	8,06
TRINITY_DN32050_c1_g2_i6	Intersectin-1	8,06
TRINITY_DN29896_c1_g1_i4	BMP-2-inducible protein kinase	8,06
TRINITY_DN30313_c1_g4_i2	Bromodomain adjacent to zinc finger domain protein 1A	8,06
TRINITY_DN25356_c0_g1_i2	Neurocan core protein	8,10
TRINITY_DN30299_c0_g1_i9	Putative monooxygenase p33MONOX	8,10
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TRINITY DN20171 -2 -1 :2	112	0.10
TRINITY_DN29171_c2_g1_i2	U2 small nuclear ribonucleoprotein A'	8,10
TRINITY_DN32677_c0_g1_i4	DENN domain-containing protein 5B	8,10
TRINITY_DN28371_c1_g2_i3	Sodium- and chloride-dependent creatine transporter 1	8,10
TRINITY_DN27657_c1_g1_i8	Collectin-11	8,10
TRINITY_DN24008_c0_g1_i3	Dexamethasone-induced protein homolog	8,10
TRINITY_DN24970_c0_g1_i2	Cytochrome P450 2C20	8,10
TRINITY_DN30007_c1_g1_i3	Endoplasmic reticulum mannosyl-oligosaccharide 1,2-alpha-mc	8,10
TRINITY_DN30256_c0_g1_i3	40S ribosomal protein S3-A	8,10
TRINITY_DN28993_c1_g2_i3	Complement component C7	8,10
TRINITY_DN24846_c0_g1_i5	Methyl-CpG-binding domain protein 2	8,10
TRINITY_DN33190_c2_g1_i10	RNA-directed DNA polymerase from mobile element jockey	8,10
TRINITY_DN29141_c0_g1_i1	Pleckstrin homology domain-containing family M member 2	8,10
TRINITY_DN32001_c2_g1_i8	Apolipoprotein B-100	8,10
TRINITY_DN31616_c0_g2_i2	Zinc finger protein 567	8,10
TRINITY_DN25554_c0_g1_i16	Serine/threonine-protein phosphatase PP1-gamma catalytic sul	8,10
TRINITY_DN28811_c1_g3_i1	Ribosomal protein S6 kinase alpha-4	8,10
TRINITY_DN30781_c0_g1_i1	Phosphatidylinositol 4,5-bisphosphate 3-kinase catalytic subuni	8,10
TRINITY_DN30336_c1_g1_i3	Transcriptional regulator ATRX	8,10
TRINITY_DN22509_c0_g1_i2	Protein BTG3	8,10
TRINITY_DN30569_c0_g1_i16	Leptin receptor gene-related protein	8,10
TRINITY_DN25790_c1_g2_i5	Tumor necrosis factor ligand superfamily member 10	8,10
TRINITY_DN32544_c0_g2_i6	Alpha-adducin	8,10
TRINITY_DN32288_c0_g2_i4	HEAT repeat-containing protein 1	8,10
TRINITY_DN30759_c0_g2_i13	Malate synthase 1, glyoxysomal	8,10
TRINITY_DN29862_c2_g3_i1	Transmembrane protein 117	8,10
TRINITY_DN27639_c1_g3_i1	Nibrin	8,10
TRINITY_DN29685_c1_g1_i9	Ubiquitin carboxyl-terminal hydrolase 47	8,10
TRINITY_DN28229_c2_g1_i4	Centromere protein J	8,10
TRINITY_DN25788_c1_g3_i1	Manganese-dependent ADP-ribose/CDP-alcohol diphosphatase	8,10
TRINITY_DN29191_c2_g1_i18	BRISC and BRCA1-A complex member 2	8,10
TRINITY_DN32882_c1_g1_i5	Inositol 1,4,5-trisphosphate receptor type 1	8,10
TRINITY_DN27339_c1_g1_i8	Protein cornichon homolog 1	8,10
TRINITY_DN33282_c5_g1_i1	Regulator of microtubule dynamics protein 3	8,10
TRINITY_DN31276_c0_g1_i14	Exosome RNA helicase MTR4	8,10
TRINITY_DN22731_c0_g1_i2	Zinc finger protein 436	8,14
TRINITY_DN31258_c2_g1_i1	TOM1-like protein 2	8,14
TRINITY_DN28371_c1_g2_i6	Sodium- and chloride-dependent creatine transporter I	8,14
TRINITY_DN28371_c1_g2_i14	Sodium- and chloride-dependent creatine transporter I Sodium- and chloride-dependent creatine transporter I	8,14
TRINITY_DN25847_c0_g2_i2	Protein 4.1	8,14
TRINITY_DN32844_c3_g1_i10	5'-nucleotidase domain-containing protein 2	8,14
TRINITY_DN32314_c5_g1_i5	Protein AMBP (Fragment)	8,14
TRINITY_DN28172_c2_g4_i1	Phosphatidylinositol 3,4,5-trisphosphate 5-phosphatase 2A	
TRINITY_DN28631_c1_g1_i9		8,14
TRINITY_DN17863_c0_g1_i2	SET domain-containing protein 5 Protein ALP1-like	8,14 8,14
TRINITY_DN31505_c0_g1_i14	Insulin receptor	8,14
TRINITY_DN29807_c0_g1_i4	Zinc finger BED domain-containing protein 1	8,14
TRINITY_DN32851_c1_g3_i2	Signal transducer and activator of transcription 1	8,14
TRINITY_DN28646_c0_g1_i2	Phosphatidylinositol-binding clathrin assembly protein	8,14
TRINITY_DN27557_c2_g1_i4	Serine/threonine-protein kinase TAO1	8,14
TRINITY_DN22086_c0_g1_i6	Phosphatidylglycerophosphatase and protein-tyrosine phosphat	8,14
TRINITY_DN32611_c1_g2_i5	Death-inducer obliterator 1	8,14
TRINITY_DN29732_c0_g1_i11	Phosphatidate phosphatase LPIN2	8,14
TRINITY_DN33055_c1_g1_i5	2-(3-amino-3-carboxypropyl)histidine synthase subunit 2	8,14
TRINITY_DN28269_c0_g1_i4	Arfaptin-2	8,14
TRINITY_DN32298_c3_g1_i8	Unconventional myosin-IXb	8,14
TRINITY_DN31771_c0_g1_i4	RAB6A-GEF complex partner protein 1	8,14
TRINITY_DN22200_c0_g1_i4	Protein FAM204A	8,14
TRINITY_DN31738_c2_g2_i11	Transposon Tf2-11 polyprotein	8,14
TRINITY_DN27578_c0_g1_i7	Myotubularin-related protein 7	8,14

TDINITY DN20200 at at :4	Multinesicular hada subunit 124	8,14
TRINITY_DN28280_c1_g1_i4	Multivesicular body subunit 12A	8,14
TRINITY_DN28177_c2_g2_i15	Tumor necrosis factor receptor superfamily member 5	
TRINITY_DN25418_c2_g2_i6	Transposon Tf2-9 polyprotein Petrovinus polytod Polyprotein from tuguen again 412	8,14
TRINITY_DN26917_c0_g1_i4	Retrovirus-related Pol polyprotein from transposon 412	8,14
TRINITY_DN25731_c4_g1_i6	Phosphatidylinositol 3,4,5-trisphosphate 3-phosphatase and du	8,14
TRINITY_DN31103_c0_g3_i10	Ras-related GTP-binding protein C	8,14
TRINITY_DN29919_c0_g3_i2	Death-associated protein 1	8,14
TRINITY_DN32046_c0_g1_i14	Choline kinase alpha	8,14
TRINITY_DN32433_c1_g1_i4	Coiled-coil-helix-coiled-coil-helix domain-containing protein 7	8,14
TRINITY_DN25998_c1_g5_i4	F-BAR domain only protein 1	8,14
TRINITY_DN31030_c0_g1_i3	Putative Polycomb group protein ASXL2	8,14
TRINITY_DN26290_c1_g2_i1	Heterogeneous nuclear ribonucleoprotein R	8,14
TRINITY_DN30655_c2_g1_i11	Pseudouridylate synthase 7 homolog	8,14
TRINITY_DN30992_c0_g1_i3	Poly [ADP-ribose] polymerase 14	8,14
TRINITY_DN27039_c0_g1_i4	Protein FAM234A	8,14
TRINITY_DN24546_c1_g2_i17	Mitochondrial fission factor homolog B	8,14
TRINITY_DN30314_c0_g1_i9	Mini-chromosome maintenance complex-binding protein	8,14
TRINITY_DN31812_c0_g1_i6	Chromatin-remodeling ATPase INO80	8,14
TRINITY_DN32326_c1_g1_i5	Putative E3 ubiquitin-protein ligase UNKL	8,18
TRINITY_DN26452_c0_g2_i12	Histone-lysine N-methyltransferase EHMT2	8,18
TRINITY_DN28232_c0_g2_i9	Nicotinamide N-methyltransferase	8,18
TRINITY_DN31482_c0_g1_i8	[Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme	8,18
TRINITY_DN32616_c3_g1_i7	Alpha-2-antiplasmin	8,18
TRINITY_DN25059_c0_g1_i1	Zinc finger X-chromosomal protein	8,18
TRINITY_DN30429_c0_g1_i5	Coiled-coil domain-containing protein 77	8,18
TRINITY_DN27788_c0_g1_i1	RUN and SH3 domain-containing protein 1	8,18
TRINITY_DN29767_c0_g1_i5	Exportin-2	8,18
TRINITY_DN26605_c0_g1_i12	Transcription factor 7-like 2	8,18
TRINITY_DN30027_c1_g2_i2	FGGY carbohydrate kinase domain-containing protein	8,18
TRINITY_DN31787_c4_g2_i10	SUN domain-containing protein 1	8,18
TRINITY_DN25778_c0_g3_i5	Zinc transporter Slc39a7 (Fragment)	8,18
TRINITY_DN28916_c1_g1_i5	E3 SUMO-protein ligase RanBP2 (Fragment)	8,18
TRINITY_DN27774_c7_g1_i4	Androgen receptor	8,18
TRINITY_DN32311_c1_g1_i27	Histone deacetylase 4	8,18
TRINITY_DN29174_c0_g1_i2	Transmembrane protein 68	8,18
TRINITY_DN28347_c0_g1_i8	Septin-2	8,18
TRINITY_DN28660_c1_g1_i2	Transmembrane protein 79	8,18
TRINITY_DN30975_c2_g1_i5	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subun	8,18
TRINITY_DN28111_c2_g5_i1	Phosphoenolpyruvate carboxykinase, cytosolic [GTP]	8,18
TRINITY DN29038 c1 g1 i7	Transmembrane gamma-carboxyglutamic acid protein 1	8,18
TRINITY_DN24858_c0_g1_i3	Ninjurin-1	8,18
TRINITY_DN24576_c1_g2_i13	Ras-related protein Rab-28	8,18
TRINITY_DN24868_c4_g2_i4	Mastermind-like protein 3	8,18
TRINITY_DN25035_c3_g1_i17	Prolyl 4-hydroxylase subunit alpha-1	8,18
TRINITY_DN28950_c0_g2_i6	Cyclin-C	8,18
TRINITY_DN29688_c0_g1_i7	DIS3-like exonuclease 2	8,18
TRINITY_DN29727_c2_g1_i6	ETS-related transcription factor Elf-2	8,18
TRINITY_DN31075_c1_g1_i2	Ras GTPase-activating protein-binding protein 1	8,18
TRINITY_DN32881_c1_g1_i2	Receptor-type tyrosine-protein phosphatase C	8,18
TRINITY_DN32415_c1_g1_i5	Very long-chain specific acyl-CoA dehydrogenase, mitochondria	8,18
TRINITY_DN32024_c1_g1_i8	Chromodomain-helicase-DNA-binding protein 1	8,18
TRINITY_DN33162_c1_g1_i10	Transmembrane protein KIAA1109 homolog	8,18
TRINITY_DN29216_c4_g1_i5	Major facilitator superfamily domain-containing protein 9	8,18
TRINITY_DN30599_c1_g2_i6	Metaxin-1	8,18
TRINITY_DN25951_c0_g1_i9	UPF0538 protein C2orf76 homolog	8,18
TRINITY_DN24138_c0_g1_i3	BEN domain-containing protein 3	8,18
TRINITY_DN30311_c1_g2_i14	CMP-N-acetylneuraminate-beta-1,4-galactoside alpha-2,3-sialy	8,18
TRINITY_DN32025_c2_g1_i9	Immunoglobulin lambda-like polypeptide 1	8,18
TRINITY_DN26472_c0_g1_i2	2-phosphoxylose phosphatase 1	8,22
	I F F F	-,

TRINITY DN24421 -2 -2 :2	III.;; i.; III.	0.22
TRINITY_DN24421_c2_g2_i3	Ubiquitin-like protein FUBI	8,22
TRINITY_DN29764_c1_g3_i1	ATP-binding cassette sub-family G member 2	8,22
TRINITY_DN22066_c0_g1_i1	Uncharacterized protein C7orf26 homolog	8,22
TRINITY_DN30655_c2_g1_i27	Pseudouridylate synthase 7 homolog	8,22
TRINITY_DN25306_c2_g1_i5	Interferon-stimulated 20 kDa exonuclease-like 2	8,22
TRINITY_DN27984_c2_g1_i2	Tyrosine-protein kinase transforming protein Fps	8,22
TRINITY_DN26397_c0_g1_i1	Vacuolar protein sorting-associated protein 29	8,22
TRINITY_DN30276_c0_g1_i5	Tyrosine-protein phosphatase non-receptor type 1	8,22
TRINITY_DN25110_c0_g1_i8	E3 ubiquitin-protein ligase BRE1B	8,22
TRINITY_DN28179_c2_g1_i3	F-box only protein 4	8,22
TRINITY_DN28229_c2_g1_i6	Centromere protein J	8,22
TRINITY_DN25606_c0_g1_i3	Class I histocompatibility antigen, F10 alpha chain	8,22
TRINITY_DN24224_c0_g1_i9	E3 ubiquitin-protein ligase TRIM33	8,22
TRINITY_DN28705_c0_g1_i5	Hepatoma-derived growth factor-related protein 2	8,22
TRINITY_DN26264_c2_g1_i4	Zinc finger protein 300	8,22
TRINITY_DN28333_c1_g1_i24	Patatin-like phospholipase domain-containing protein 2	8,22
TRINITY_DN28723_c0_g2_i3	Serine/threonine-protein kinase tousled-like 2	8,22
TRINITY_DN29164_c0_g2_i1	Phospholipid phosphatase 4	8,22
TRINITY_DN26729_c3_g2_i9	Protein NLRC3	8,22
TRINITY_DN23175_c0_g1_i5	Adenylate kinase isoenzyme 6	8,22
TRINITY_DN28430_c1_g1_i15	Mitochondrial uncoupling protein 2	8,22
TRINITY_DN33238_c5_g1_i16	Telomerase protein component 1	8,22
TRINITY_DN28263_c0_g1_i9	Inositol hexakisphosphate and diphosphoinositol-pentakisphosp	8,22
TRINITY_DN23404_c0_g1_i4		8,22
	Persulfide dioxygenase ETHE1, mitochondrial	
TRINITY_DN29621_c0_g1_i5	Ras-related protein Rab-8B	8,22
TRINITY_DN31378_c0_g1_i27	Cyclic AMP-dependent transcription factor ATF-6 beta	8,22
TRINITY_DN27581_c0_g1_i6	Adiponectin receptor protein 1	8,22
TRINITY_DN28800_c1_g1_i6	Heat shock protein HSP 90-alpha 1	8,22
TRINITY_DN32546_c0_g1_i11	E3 ubiquitin-protein ligase TTC3	8,22
TRINITY_DN28368_c1_g1_i19	Protein FAM13A	8,22
TRINITY_DN31417_c0_g2_i22	Aryl hydrocarbon receptor nuclear translocator	8,22
TRINITY_DN32475_c5_g1_i7	UTPglucose-1-phosphate uridylyltransferase	8,22
TRINITY_DN29183_c0_g1_i2	ADP-ribosylation factor-binding protein GGA3	8,22
TRINITY_DN30411_c1_g1_i7	Transposon Ty3-I Gag-Pol polyprotein	8,22
TRINITY_DN30585_c0_g1_i14	Hemopexin	8,22
TRINITY_DN28371_c1_g2_i17	Sodium- and chloride-dependent creatine transporter 1	8,22
TRINITY_DN32693_c7_g1_i11	Interleukin-1 receptor type 2	8,22
TRINITY_DN25960_c3_g3_i3	B-cell receptor CD22	8,22
TRINITY_DN24510_c0_g1_i1	Transmembrane emp24 domain-containing protein 5	8,22
TRINITY_DN28166_c1_g1_i2	CAAX prenyl protease 2	8,22
TRINITY_DN27877_c0_g1_i1	Histone H4 transcription factor	8,22
TRINITY_DN25458_c0_g1_i4	RNA-binding protein 34	8,25
TRINITY_DN24920_c4_g1_i7	Immune-associated nucleotide-binding protein 4	8,25
TRINITY_DN23926_c1_g1_i6	Transposable element Tcb1 transposase	8,25
TRINITY_DN31903_c0_g1_i20	Fanconi anemia group A protein homolog	8,25
TRINITY_DN26407_c0_g2_i3	Voltage-dependent anion-selective channel protein 2	8,25
TRINITY_DN28898_c0_g3_i2	CXXC-type zinc finger protein 5	8,25
TRINITY_DN31579_c0_g2_i5	APOBEC1 complementation factor	8,25
TRINITY_DN23398_c1_g1_i4	Redox-regulatory protein FAM213A	8,25
TRINITY_DN29606_c0_g1_i2	Collagen alpha-1(V) chain	8,25
TRINITY_DN28401_c0_g1_i3		
	Unconventional myosin-If V type proton ATPase catalytic subunit A	8,25 8,25
TRINITY_DN25915_c0_g1_i1	V-type proton ATPase catalytic subunit A Cyclin dependent kings 2 associated protein I	8,25
TRINITY_DN21569_c0_g1_i1	Cyclin-dependent kinase 2-associated protein 1	8,25
TRINITY_DN27588_c0_g1_i9	Nucleolar and coiled-body phosphoprotein 1	8,25
TRINITY_DN23526_c0_g1_i8	Vacuole membrane protein 1	8,25
TRINITY_DN26614_c0_g1_i14	Rho-related BTB domain-containing protein 3	8,25
TRINITY_DN29430_c6_g1_i4	Y+L amino acid transporter 2	8,25
TRINITY_DN33264_c7_g1_i8	Complement component receptor 1-like protein	8,25
TRINITY_DN26453_c0_g1_i8	Multidrug resistance-associated protein 5	8,25

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TRINITY_DN26913_c1_g1_i1	39S ribosomal protein L28, mitochondrial	8,25
TRINITY_DN22958_c0_g1_i6	Abl interactor 1	8,25
TRINITY_DN28536_c0_g1_i9	SRSF protein kinase 1	8,25
TRINITY_DN31651_c0_g1_i7	Zinc finger and BTB domain-containing protein 10	8,25
TRINITY_DN31953_c3_g2_i11	PAN2-PAN3 deadenylation complex subunit pan3	8,25
TRINITY_DN28616_c0_g2_i7	Probable RNA-directed DNA polymerase from transposon BS	8,25
TRINITY_DN31873_c0_g1_i8	DAP3-binding cell death enhancer 1	8,25
TRINITY_DN27676_c0_g2_i5	GRAM domain-containing protein 2A	8,25
TRINITY_DN27026_c0_g1_i9	Transmembrane protein 268	8,25
TRINITY_DN33020_c6_g1_i2	Acyl-CoA synthetase family member 2, mitochondrial	8,25
TRINITY_DN25994_c3_g1_i3	Protein FAM126B	8,25
TRINITY_DN28755_c2_g1_i5	NADP-dependent malic enzyme	8,25
TRINITY_DN29392_c0_g1_i2	RNA-binding protein 39	8,25
TRINITY_DN26377_c0_g1_i3	Secretory carrier-associated membrane protein 3	8,25
TRINITY_DN25933_c0_g1_i3	Alpha-1,3/1,6-mannosyltransferase ALG2	8,25
TRINITY_DN29231_c0_g1_i1	Transmembrane protein adipocyte-associated 1 homolog	8,25
TRINITY_DN25244_c0_g1_i2	Protein canopy-1	8,25
TRINITY_DN25435_c0_g1_i13	ATP-dependent RNA helicase DHX36	8,25
TRINITY_DN30991_c1_g2_i3	Sarcoplasmic/endoplasmic reticulum calcium ATPase 2	8,25
TRINITY_DN24659_c0_g1_i2	Histone-binding protein RBBP7	8,25
TRINITY_DN30451_c0_g1_i6	E3 ubiquitin-protein ligase CHFR	8,25
TRINITY_DN30949_c0_g1_i4	BUB3-interacting and GLEBS motif-containing protein ZNF20%	8,25
TRINITY_DN30572_c0_g1_i13	Lysosomal alpha-glucosidase	8,25
TRINITY_DN29178_c2_g1_i3	Serine hydroxymethyltransferase, mitochondrial	8,29
TRINITY_DN28903_c0_g1_i16	Eukaryotic translation initiation factor 3 subunit K	8,29
TRINITY_DN27303_c0_g3_i5	Nucleolin	8,29
TRINITY_DN28631_c1_g1_i13	SET domain-containing protein 5	8,29
TRINITY_DN28580_c4_g1_i1	Interferon regulatory factor 2	8,29
TRINITY_DN31007_c3_g1_i2	DnaJ homolog subfamily C member 5	8,29
TRINITY_DN28883_c3_g1_i10	WASH complex subunit 4	8,29
TRINITY_DN32423_c1_g2_i3	Endosomal/lysosomal potassium channel TMEM175	8,29
TRINITY_DN28504_c0_g1_i1	Metaxin-2	8,29
TRINITY_DN30380_c4_g1_i4	10 kDa heat shock protein, mitochondrial	8,29
	Protein Mdm4	
TRINITY_DN32924_c1_g1_i15		8,29
TRINITY_DN27940_c0_g2_i5	Complement C3 (Fragment)	8,29
TRINITY_DN29628_c0_g1_i10	Cytochrome c-type heme lyase	8,29
TRINITY_DN31052_c0_g1_i6	Transposon Ty3-G Gag-Pol polyprotein	8,29
TRINITY_DN30555_c2_g4_i2	Carnitine O-palmitoyltransferase 1, liver isoform	8,29
TRINITY_DN24274_c0_g1_i1	Mannosyl-oligosaccharide glucosidase	8,29
TRINITY_DN32340_c0_g1_i16	Peptidyl-prolyl cis-trans isomerase D	8,29
TRINITY_DN30262_c1_g1_i11	Protein kinase C-binding protein 1	8,29
TRINITY_DN23083_c0_g1_i7	Mitochondrial ribonuclease P catalytic subunit	8,29
TRINITY_DN31603_c1_g1_i20	G-protein-signaling modulator 2	8,29
TRINITY_DN29389_c1_g1_i13	Tetratricopeptide repeat protein 13	8,29
TRINITY_DN32582_c0_g1_i1	RING finger protein 10	8,29
TRINITY_DN26146_c0_g1_i7	L-serine dehydratase/L-threonine deaminase	8,29
TRINITY_DN30093_c0_g1_i4	CUE domain-containing protein 1	8,29
TRINITY_DN30321_c0_g1_i11	Nuclear transcription factor Y subunit gamma	8,29
TRINITY_DN22901_c0_g2_i5	snRNA-activating protein complex subunit 4	8,29
TRINITY_DN27823_c1_g1_i13	Protein NOXP20	8,29
TRINITY_DN32338_c1_g1_i1	MORC family CW-type zinc finger protein 3	8,29
TRINITY_DN32302_c1_g1_i16	ATP-dependent RNA helicase DDX42	8,29
TRINITY_DN28784_c0_g1_i4	Arf-GAP domain and FG repeat-containing protein 1	8,29
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TRINITY_DN28540_c1_g1_i13	HCLS1-associated protein X-1	8,29
TRINITY_DN31220_c1_g1_i2	Tax1-binding protein 1 homolog B	8,29
TRINITY_DN26848_c0_g1_i12	Peroxisome proliferator-activated receptor delta	8,29
TRINITY_DN30419_c0_g1_i11	Urotensin-2 receptor	8,29
TRINITY_DN32311_c1_g1_i2	Histone deacetylase 4	8,29
TRINITY_DN30470_c0_g1_i5	Sodium/potassium-transporting ATPase subunit alpha-1	8,29

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TRINITY_DN28233_c1_g1_i14	Glycogen synthase kinase-3 beta	8,29
TRINITY_DN28443_c0_g1_i10	Zinc finger protein 92 homolog	8,29
TRINITY_DN28801_c0_g1_i5	FTS and Hook-interacting protein homolog	8,29
TRINITY_DN31126_c0_g1_i4	Heterogeneous nuclear ribonucleoprotein U-like protein 2	8,29
TRINITY_DN30418_c0_g1_i5	Palmitoyltransferase ZDHHC9	8,29
TRINITY_DN27034_c1_g2_i10	Sodium-dependent multivitamin transporter	8,29
TRINITY_DN29351_c1_g3_i2	Low-density lipoprotein receptor-related protein 5	8,32
TRINITY_DN32271_c1_g1_i14	Coxsackievirus and adenovirus receptor homolog	8,32
TRINITY_DN33150_c2_g2_i27	Oxidation resistance protein 1	8,32
TRINITY_DN33187_c0_g1_i6	E3 ubiquitin-protein ligase HERC2	8,32
TRINITY_DN29949_c1_g3_i7	Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial	8,32
TRINITY_DN24429_c1_g1_i9	H-2 class II histocompatibility antigen, I-E beta chain	8,32
TRINITY_DN28165_c2_g1_i7		8,32
_	Ribosomal protein S6 kinase alpha-3	
TRINITY_DN30949_c0_g1_i1	BUB3-interacting and GLEBS motif-containing protein ZNF20%	8,32
TRINITY_DN23444_c0_g1_i8	Cerebellin-3	8,32
TRINITY_DN27858_c2_g3_i3	Transposable element Tcb1 transposase	8,32
TRINITY_DN26425_c3_g1_i3	Translocation protein SEC63 homolog	8,32
TRINITY_DN30802_c0_g1_i3	TRPM8 channel-associated factor homolog	8,32
TRINITY_DN25474_c0_g2_i4	Sentrin-specific protease 1	8,32
TRINITY_DN31276_c0_g1_i4	Exosome RNA helicase MTR4	8,32
TRINITY_DN27940_c1_g2_i6	D-glucuronyl C5-epimerase B	8,32
TRINITY_DN28782_c0_g1_i10	H-2 class II histocompatibility antigen, E-U alpha chain	8,32
TRINITY_DN25200_c2_g1_i31	GTPase IMAP family member 7	8,32
TRINITY_DN24527_c0_g1_i10	Multiple epidermal growth factor-like domains protein 6	8,32
TRINITY_DN23415_c0_g1_i4	Nuclear pore complex protein Nup85	8,36
TRINITY_DN31058_c0_g2_i11	Moesin	8,36
TRINITY_DN27403_c0_g1_i10	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial	8,36
TRINITY_DN30684_c3_g1_i3	Enhancer of polycomb homolog 1	8,36
TRINITY_DN32660_c0_g1_i10	Protein scribble homolog	8,36
TRINITY_DN31647_c0_g1_i5	Methionine synthase	8,36
TRINITY_DN27015_c0_g2_i7	Myocyte-specific enhancer factor 2D homolog	8,36
TRINITY_DN26828_c0_g2_i2	Upstream stimulatory factor 2	8,36
TRINITY_DN27360_c0_g1_i2		8,36
	Desumoylating isopeptidase 1	
TRINITY_DN30262_c1_g1_i8	Protein kinase C-binding protein 1	8,36
TRINITY_DN25778_c0_g3_i8	Zinc transporter Slc39a7 (Fragment)	8,36
TRINITY_DN28798_c0_g1_i4	Transmembrane protein 180	8,36
TRINITY_DN25997_c0_g1_i3	MAGUK p55 subfamily member 6	8,36
TRINITY_DN27560_c0_g1_i5	EARP-interacting protein homolog	8,36
TRINITY_DN33095_c2_g1_i15	Mitogen-activated protein kinase kinase kinase 3	8,36
TRINITY_DN27917_c0_g1_i4	Ubiquitin-fold modifier 1	8,36
TRINITY_DN31393_c1_g1_i13	Activin receptor type-2A	8,36
TRINITY_DN23420_c0_g1_i1	Proliferation marker protein Ki-67	8,36
TRINITY_DN30164_c0_g1_i1	Periodic tryptophan protein 2 homolog	8,36
TRINITY_DN27686_c3_g1_i5	MORC family CW-type zinc finger protein 2A	8,36
TRINITY_DN25171_c0_g1_i9	Complement C3 (Fragment)	8,36
TRINITY_DN29301_c0_g1_i2	Structural maintenance of chromosomes protein 1A	8,36
TRINITY_DN28346_c0_g3_i2	Dolichyl-diphosphooligosaccharideprotein glycosyltransferas	8,36
TRINITY_DN31134_c0_g1_i4	Drebrin	8,36
TRINITY_DN26841_c0_g1_i10	SPRY domain-containing SOCS box protein 3	8,36
TRINITY_DN26020_c1_g2_i1	Phosphoenolpyruvate carboxykinase, cytosolic [GTP]	8,36
TRINITY_DN31560_c1_g2_i10	Pre-mRNA cleavage complex 2 protein Pcf11	8,36
TRINITY_DN22948_c0_g1_i10	Ras-related and estrogen-regulated growth inhibitor-like protei.	8,36
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TRINITY_DN27995_c0_g1_i7	Nuclear respiratory factor 1 Cyanina myslootida hinding protein subunit alpha 13	8,36
TRINITY_DN26342_c1_g2_i4	Guanine nucleotide-binding protein subunit alpha-13	8,36
TRINITY_DN31622_c1_g1_i6	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 3	8,36
TRINITY_DN27998_c0_g1_i13	IGF-like family receptor 1	8,36
TRINITY_DN31945_c1_g1_i5	Protein diaphanous homolog 2	8,36
TRINITY_DN32564_c3_g1_i3	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	8,36
TRINITY_DN32502_c2_g1_i12	Ethanolamine-phosphate phospho-lyase	8,36

TRINITY DN22426 .1 .1 '0	D	0.26
TRINITY_DN32426_c1_g1_i9	Protein transport protein Sec24C	8,36
TRINITY_DN25030_c1_g1_i3	Iodotyrosine deiodinase 1	8,36
TRINITY_DN24617_c2_g1_i1	Testican-3	8,36
TRINITY_DN27840_c0_g1_i11	Cell division cycle protein 27 homolog	8,36
TRINITY_DN32980_c3_g5_i15	Arf-GAP with coiled-coil, ANK repeat and PH domain-containing	8,36
TRINITY_DN26998_c2_g5_i5	Transmembrane protein 14C	8,36
TRINITY_DN22421_c0_g1_i5	RNA-binding protein 4.1	8,36
TRINITY_DN28187_c0_g2_i1	PHD finger protein 13	8,36
TRINITY_DN27189_c1_g1_i12	Lysine-specific demethylase 5A	8,36
TRINITY_DN30242_c0_g1_i3	Rho guanine nucleotide exchange factor 1	8,39
TRINITY_DN27406_c0_g1_i8	pre-mRNA 3' end processing protein WDR33	8,39
TRINITY_DN33021_c1_g1_i6	Cysteine-rich with EGF-like domain protein 1	8,39
TRINITY_DN29805_c1_g2_i5	[Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme	8,39
TRINITY_DN25976_c0_g1_i13	Nephrocystin-3	8,39
_	Platelet endothelial cell adhesion molecule	8,39
TRINITY_DN26289_c0_g1_i10		
TRINITY_DN22652_c0_g1_i7	Protein LYRIC	8,39
TRINITY_DN27947_c0_g1_i5	Endoplasmic reticulum-Golgi intermediate compartment protein	8,39
TRINITY_DN31380_c0_g1_i1	Zinc finger protein ZXDC	8,39
TRINITY_DN33110_c2_g2_i13	Myomegalin	8,39
TRINITY_DN25594_c3_g2_i7	Mitogen-activated protein kinase kinase kinase 2	8,39
TRINITY_DN27774_c7_g1_i7	Androgen receptor (Fragment)	8,39
TRINITY_DN29377_c1_g1_i9	KN motif and ankyrin repeat domain-containing protein 1	8,39
TRINITY_DN30132_c0_g1_i1	L-fucose kinase	8,39
TRINITY_DN31821_c2_g1_i9	Choline kinase alpha	8,39
TRINITY_DN29940_c2_g1_i10	Specifically androgen-regulated gene protein	8,39
TRINITY_DN31328_c1_g1_i15	PHD finger protein 21A	8,39
TRINITY_DN25171_c0_g1_i7	Complement C3 (Fragment)	8,39
TRINITY_DN32303_c0_g1_i9	Endoplasmic reticulum mannosyl-oligosaccharide 1,2-alpha-ma	8,39
TRINITY_DN29219_c0_g1_i4	Zinc finger and BTB domain-containing protein 21	8,39
TRINITY_DN30980_c0_g2_i6	Lysine-specific demethylase 2A	8,39
TRINITY_DN31065_c3_g1_i7	Cytoskeleton-associated protein 4	8,39
TRINITY_DN26934_c2_g1_i13	Insulin-like growth factor II	8,39
TRINITY_DN32264_c1_g1_i10	Diacylglycerol O-acyltransferase 2	8,39
_		
TRINITY_DN31499_c0_g2_i7	Attractin	8,39
TRINITY_DN32978_c0_g1_i11	ValinetRNA ligase	8,39
TRINITY_DN27288_c0_g1_i7	U3 small nucleolar RNA-associated protein 4 homolog	8,39
TRINITY_DN29147_c1_g2_i6	Inhibin beta B chain	8,43
TRINITY_DN33063_c4_g3_i4	Bone morphogenetic protein 1	8,43
TRINITY_DN30453_c0_g1_i5	Chromobox protein homolog 1	8,43
TRINITY_DN27783_c0_g1_i3	Anthrax toxin receptor 2	8,43
TRINITY_DN25805_c0_g1_i4	Seipin	8,43
TRINITY_DN29303_c0_g1_i6	E3 ubiquitin-protein ligase MYCBP2	8,43
TRINITY_DN24557_c0_g1_i2	Chromobox protein homolog 8	8,43
TRINITY_DN24625_c1_g3_i1	Probable low-specificity L-threonine aldolase 2	8,43
TRINITY_DN29460_c0_g1_i4	Long-chain fatty acid transport protein 6	8,43
TRINITY_DN27015_c0_g3_i1	Membrane-associated phosphatidylinositol transfer protein 2	8,43
TRINITY_DN27135_c0_g1_i4	Carbohydrate deacetylase	8,43
TRINITY_DN29093_c0_g1_i3	Biotinprotein ligase	8,43
TRINITY_DN28170_c0_g1_i9	Sperm-associated antigen 1A	8,43
TRINITY_DN29677_c1_g1_i4	Nectin-4	8,43
TRINITY_DN29965_c1_g1_i5	T-cell activation inhibitor, mitochondrial	8,43
TRINITY_DN27850_c0_g1_i4	Heparan sulfate glucosamine 3-O-sulfotransferase 3B1	8,43
TRINITY_DN32308_c1_g2_i9	60 kDa lysophospholipase	8,43
TRINITY_DN23905_c0_g1_i3		
_	Zinc finger FYVE domain-containing protein 1	8,43
TRINITY_DN26354_c2_g3_i4	Epidermal retinol dehydrogenase 2	8,43
TRINITY_DN29249_c0_g1_i12	CD276 antigen	8,43
TRINITY_DN28960_c0_g1_i8	Interferon-induced GTP-binding protein Mx	8,43
TRINITY_DN28614_c1_g1_i2	Inorganic pyrophosphatase	8,43
TRINITY_DN32298_c3_g1_i6	Unconventional myosin-IXb	8,43

FD 11 1 FT	** 1	0.42
TRINITY_DN22968_c0_g1_i5	Uncharacterized protein C1orf21 homolog	8,43
TRINITY_DN29916_c0_g2_i7	Actin filament-associated protein 1-like 1	8,43
TRINITY_DN24604_c1_g1_i4	Nuclear factor 1 B-type	8,43
TRINITY_DN28127_c1_g4_i12	Bifunctional heparan sulfate N-deacetylase/N-sulfotransferase 2	8,43
TRINITY_DN28070_c1_g1_i10	Dystonin	8,43
TRINITY_DN29069_c4_g1_i8	Retrotransposon-derived protein PEG10	8,43
TRINITY_DN28903_c0_g1_i12	Eukaryotic translation initiation factor 3 subunit K	8,43
TRINITY_DN32646_c1_g1_i8	CAD protein	8,43
TRINITY_DN31936_c0_g2_i6	1-acyl-sn-glycerol-3-phosphate acyltransferase gamma	8,43
TRINITY_DN28879_c0_g1_i6	Protein unc-45 homolog A	8,43
TRINITY_DN33144_c6_g1_i9	Zinc finger MYM-type protein 2	8,43
_	E3 ubiquitin-protein ligase HERC2	8,43
TRINITY_DN33187_c0_g1_i9		
TRINITY_DN30084_c0_g1_i10	Zinc fingers and homeoboxes protein 3	8,43
TRINITY_DN30356_c0_g1_i4	Protein YIF1A	8,43
TRINITY_DN27440_c4_g4_i9	F-box only protein 21	8,43
TRINITY_DN32965_c1_g1_i5	Cytochrome P450 2F2	8,45
TRINITY_DN31440_c0_g1_i9	Rab3 GTPase-activating protein non-catalytic subunit	8,46
TRINITY_DN28573_c2_g1_i9	Serine incorporator 5	8,46
TRINITY_DN26020_c1_g1_i11	Sepiapterin reductase	8,46
TRINITY_DN25268_c0_g1_i4	WD repeat-containing protein 20	8,46
TRINITY_DN28708_c0_g1_i1	Sialin	8,46
TRINITY_DN30380_c4_g1_i7	10 kDa heat shock protein, mitochondrial	8,46
TRINITY_DN31340_c2_g2_i10	Sterol O-acyltransferase 1	8,46
TRINITY_DN28942_c2_g1_i5	Signal peptide peptidase-like 3	8,46
TRINITY_DN30248_c0_g1_i3	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyli	8,46
TRINITY_DN28618_c1_g1_i12	E3 ubiquitin-protein ligase TRAF7	8,46
TRINITY_DN31980_c1_g1_i9	Cytochrome P450 2J2	8,46
TRINITY_DN30958_c0_g2_i2	Nuclear receptor ROR-beta	8,46
TRINITY_DN24996_c1_g1_i15	Retinoic acid receptor RXR-beta-A	8,46
_		8,46
TRINITY_DN31923_c0_g1_i7	AT-rich interactive domain-containing protein 1B	
TRINITY_DN29966_c4_g1_i3	Mitochondrial glutamate carrier 1	8,46
TRINITY_DN24794_c1_g1_i2	Transmembrane and TPR repeat-containing protein 3	8,46
TRINITY_DN31677_c2_g1_i1	Solute carrier family 25 member 47-B	8,46
TRINITY_DN27998_c0_g2_i8	Equilibrative nucleoside transporter 1	8,46
TRINITY_DN27020_c0_g1_i10	Heterogeneous nuclear ribonucleoprotein M	8,46
TRINITY_DN25126_c0_g1_i4	Exportin-4	8,46
TRINITY_DN32616_c3_g1_i10	Alpha-2-antiplasmin	8,46
TRINITY_DN30009_c2_g4_i6	Sideroflexin-1	8,46
TRINITY_DN31757_c2_g1_i2	DEP domain-containing mTOR-interacting protein	8,46
TRINITY_DN26752_c2_g3_i4	Vacuolar protein sorting-associated protein 4A	8,46
TRINITY_DN24209_c0_g1_i4	26S proteasome regulatory subunit 4	8,46
TRINITY_DN30616_c1_g1_i13	Ras-related protein Rab-43	8,46
TRINITY_DN33078_c0_g1_i16	Serine/threonine-protein kinase MRCK beta	8,48
TRINITY_DN31294_c0_g1_i3	Cyclin-L1	8,49
TRINITY_DN32612_c1_g2_i11	Kynureninase	8,49
TRINITY_DN27224_c2_g1_i5	Perilipin-2	8,49
_	•	8,49
TRINITY_DN25868_c0_g1_i16	ZZ-type zinc finger-containing protein 3	
TRINITY_DN31874_c0_g1_i2	Peroxiredoxin-5, mitochondrial	8,49
TRINITY_DN25562_c0_g1_i4	Kelch-like protein 15	8,49
TRINITY_DN29289_c0_g1_i15	Ubiquitin carboxyl-terminal hydrolase MINDY-1	8,49
TRINITY_DN33014_c1_g1_i4	Omega-amidase NIT2	8,49
TRINITY_DN32671_c1_g2_i1	Ubiquitin-conjugating enzyme E2 E1	8,49
TRINITY_DN26830_c2_g2_i10	Tetratricopeptide repeat protein 17	8,49
TRINITY_DN31256_c1_g2_i8	E3 ubiquitin-protein ligase HECTD1	8,49
TRINITY_DN30790_c1_g1_i1	Tight junction-associated protein 1	8,49
TRINITY_DN30433_c0_g1_i5	Endoplasmic reticulum resident protein 44	8,49
TRINITY_DN28504_c0_g1_i6	Metaxin-2	8,49
TRINITY_DN30928_c0_g1_i2	Glycogen phosphorylase, liver form	8,49
TRINITY_DN27903_c0_g1_i8	Phosphatidylinositol 4-kinase type 2-beta	8,49
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TRINITY_DN26752_c2_g3_i2	Vacuolar protein sorting-associated protein 4A	8,49
TRINITY_DN30162_c1_g1_i14	Transport and Golgi organization protein 1 homolog	8,49
TRINITY_DN28004_c0_g1_i5	A-kinase anchor protein 11	8,52
TRINITY_DN28412_c2_g1_i7	Complement C1q tumor necrosis factor-related protein 1	8,52
TRINITY_DN32631_c0_g1_i7	Lysosome membrane protein 2	8,52
TRINITY_DN30055_c0_g1_i12	Pleckstrin homology domain-containing family A member 5	8,52
TRINITY_DN32097_c1_g1_i4	Adapter protein CIKS	8,52
TRINITY_DN24327_c0_g1_i12	Transcription factor PU.1	8,52
TRINITY_DN29711_c3_g2_i4	Protein lifeguard 1	8,52
TRINITY_DN29286_c0_g2_i7	Pinin	8,52
TRINITY_DN25461_c3_g2_i17	Polyadenylate-binding protein-interacting protein 2B	8,52
TRINITY_DN23718_c0_g1_i3	T-complex protein 1 subunit beta	8,52
_		8,52
TRINITY_DN27980_c0_g1_i4	DNA polymerase beta	
TRINITY_DN32855_c1_g1_i18	Ras-related protein Rab-23	8,52
TRINITY_DN29235_c0_g1_i11	Coiled-coil domain-containing protein 97	8,52
TRINITY_DN28074_c1_g2_i4	Mitochondrial Rho GTPase 1-A	8,52
TRINITY_DN29297_c1_g1_i2	Homeodomain-interacting protein kinase 3	8,52
TRINITY_DN30719_c0_g1_i5	Histone acetyltransferase KAT6B	8,52
TRINITY_DN32817_c2_g1_i9	Unconventional myosin-Vb	8,52
TRINITY_DN30567_c0_g1_i2	Plexin-C1	8,52
TRINITY_DN32004_c3_g2_i8	Ribosomal protein S6 kinase alpha-6	8,55
TRINITY_DN30401_c0_g1_i7	Polyphosphoinositide phosphatase	8,55
TRINITY_DN24290_c0_g1_i4	GDP-fucose protein O-fucosyltransferase 1	8,55
TRINITY_DN25744_c0_g2_i2	Nuclear receptor subfamily 2 group F member 1-A	8,55
TRINITY_DN30362_c0_g1_i13	SWI/SNF-related matrix-associated actin-dependent regulator c	8,55
TRINITY_DN28450_c1_g1_i10	Choline transporter-like protein 1	8,55
TRINITY_DN24590_c0_g1_i5	CD9 antigen	8,55
TRINITY_DN27064_c0_g1_i5	Ras-related GTP-binding protein A	8,55
TRINITY_DN24163_c2_g1_i23	von Willebrand factor A domain-containing protein 7	8,55
TRINITY_DN30365_c1_g1_i10	DNA transposase THAP9	8,55
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TRINITY_DN22809_c0_g1_i2	Pleiotropic regulator 1	8,55
TRINITY_DN32189_c0_g1_i14	Anion exchange protein 2	8,55
TRINITY_DN25032_c1_g1_i12	Activating molecule in BECN1-regulated autophagy protein 1	8,55
TRINITY_DN28046_c0_g1_i17	Retrovirus-related Pol polyprotein from transposon 17.6	8,55
TRINITY_DN26570_c0_g2_i11	Plastin-3	8,55
TRINITY_DN33056_c3_g2_i2	Serine/threonine-protein kinase ULK1	8,55
TRINITY_DN26659_c0_g2_i7	SMC5-SMC6 complex localization factor protein 2	8,55
TRINITY_DN27904_c1_g1_i6	Carboxy-terminal domain RNA polymerase II polypeptide A sm	8,55
TRINITY_DN32463_c0_g1_i7	Replication stress response regulator SDE2	8,55
TRINITY_DN32180_c0_g2_i3	Complement C3 (Fragment)	8,55
TRINITY_DN31590_c0_g1_i8	Ectonucleoside triphosphate diphosphohydrolase 4	8,55
TRINITY_DN32792_c1_g1_i16	Alkaline ceramidase 3	8,55
TRINITY_DN30633_c0_g1_i8	Ubinuclein-2	8,55
TRINITY_DN29809_c0_g1_i8	Pogo transposable element with ZNF domain	8,55
TRINITY_DN32817_c2_g1_i8	Unconventional myosin-Vb	8,55
TRINITY_DN32398_c0_g1_i19	Dyslexia-associated protein KIAA0319-like protein	8,55
TRINITY_DN31670_c0_g3_i9	Ligand-dependent nuclear receptor corepressor-like protein	8,55
TRINITY_DN26196_c0_g1_i14	Bromodomain-containing protein 3	8,55
TRINITY_DN28515_c2_g1_i6	Intersectin-1	8,58
TRINITY_DN30080_c2_g2_i9	Protein SPT2 homolog	8,58
TRINITY_DN29098_c1_g1_i2	WD repeat-containing protein 37	8,58
TRINITY_DN28687_c1_g4_i1	Bile acid receptor	8,58
TRINITY_DN28371_c1_g2_i8	Sodium- and chloride-dependent creatine transporter 1	8,58
TRINITY_DN23663_c1_g1_i8	Transcription factor Dp-1	8,58
TRINITY_DN32729_c0_g1_i15	Putative ATP-dependent RNA helicase DHX57	8,58
TRINITY_DN32840_c3_g1_i7	Hypoxia up-regulated protein 1	8,58
TRINITY_DN31771_c0_g1_i1	RAB6A-GEF complex partner protein 1	8,58
TRINITY_DN29159_c2_g1_i4	Purine-rich element-binding protein gamma	8,58
TRINITY_DN30156_c1_g1_i8	Phosphoglycerate mutase 1	8,58

TRINITY_DN25364_c4_g1_i2	Eukaryotic translation initiation factor 4E type 3	8,58
TRINITY_DN33119_c2_g1_i27	Syndecan-2	8,58
TRINITY_DN26768_c4_g1_i3	Eukaryotic translation initiation factor 4E-binding protein 3-lik	8,58
TRINITY_DN25777_c0_g1_i4	Eukaryotic translation initiation factor 1b	8,58
TRINITY_DN27581_c0_g1_i1	Adiponectin receptor protein 1	8,58
TRINITY_DN24913_c0_g1_i3	F-box only protein 8	8,58
TRINITY_DN28960_c0_g1_i18	Interferon-induced GTP-binding protein Mx	8,58
TRINITY_DN29718_c0_g1_i4	Zinc finger protein 280D	8,58
TRINITY_DN32277_c0_g1_i10	C2 domain-containing protein 2	8,58
TRINITY_DN24839_c4_g2_i1	TIP41-like protein	8,58
TRINITY_DN31272_c0_g1_i19	Ubiquitin-protein ligase E3B	8,58
TRINITY_DN32920_c1_g1_i2	Nuclear receptor coactivator 6	8,58
TRINITY_DN27916_c1_g1_i7	Guanine nucleotide-binding protein $G(t)$ subunit alpha-1	8,58
TRINITY_DN28070_c1_g1_i34 TRINITY_DN28085_c0_g2_i7	Dystonin Solute carrier family 25 member 44	8,58 8,58
TRINITY_DN25394_c0_g1_i2	Major histocompatibility complex class I-related gene protein	8,58
TRINITY_DN28861_c1_g3_i2	Lysosome-associated membrane glycoprotein 1	8,58
TRINITY_DN30615_c0_g1_i2	OTU domain-containing protein 4	8,61
TRINITY_DN32403_c1_g1_i2	Dedicator of cytokinesis protein 8	8,61
TRINITY_DN25441_c3_g5_i2	NAD(P)(+)arginine ADP-ribosyltransferase 2	8,61
TRINITY_DN25855_c0_g1_i1	Activin receptor type-1B	8,61
TRINITY_DN29933_c0_g1_i2	La-related protein 4B	8,61
TRINITY_DN25024_c0_g3_i2	BCL-6 corepressor	8,61
TRINITY_DN28834_c1_g1_i11	Phosphorylated CTD-interacting factor 1	8,61
TRINITY_DN28516_c0_g1_i3	Autophagy protein 5	8,61
TRINITY_DN25753_c0_g1_i1	T-box transcription factor TBX2b	8,61
TRINITY_DN30812_c0_g1_i6	Cullin-9	8,61
TRINITY_DN32424_c1_g1_i5	Pericentriolar material 1 protein	8,61
TRINITY_DN31374_c0_g2_i2	2-oxoglutarate dehydrogenase, mitochondrial	8,61
TRINITY_DN31110_c1_g1_i21	Histone-lysine N-methyltransferase, H3 lysine-79 specific	8,61
TRINITY_DN29516_c2_g1_i11	Complement factor H-related protein 1	8,61
TRINITY_DN30158_c1_g1_i4	Son of sevenless homolog 1	8,61
TRINITY_DN30565_c0_g1_i8	Histone-lysine N-methyltransferase setd3	8,61
TRINITY_DN30675_c0_g1_i2	Breakpoint cluster region protein	8,61
TRINITY_DN30557_c5_g1_i6	Solute carrier family 35 member E2	8,61
TRINITY_DN32426_c1_g1_i2	Protein transport protein Sec24C	8,61
TRINITY_DN27148_c1_g3_i1	Coiled-coil domain-containing protein 50	8,64
TRINITY_DN32906_c2_g1_i11	Lysosome-associated membrane glycoprotein 1	8,64
TRINITY_DN29906_c0_g2_i1	Uncharacterized protein C11orf24 homolog	8,64
TRINITY_DN26655_c0_g2_i8	Growth hormone-inducible transmembrane protein	8,64
TRINITY_DN28408_c1_g2_i5	H(+)/Cl(-) exchange transporter 5	8,64
TRINITY_DN31589_c1_g1_i14	Coagulation factor V	8,64
TRINITY_DN31172_c0_g1_i24	OTU domain-containing protein 3	8,64
TRINITY_DN32020_c4_g1_i7	Junction-mediating and -regulatory protein	8,64
TRINITY_DN30921_c1_g2_i17	Unconventional myosin-X	8,64
TRINITY_DN28996_c0_g1_i11	Egl nine homolog 1	8,64
TRINITY_DN27087_c0_g1_i3	Sodium-coupled neutral amino acid transporter 3	8,64
TRINITY_DN22081_c0_g1_i2	Fibroblast growth factor receptor-like 1	8,64
TRINITY_DN27361_c0_g1_i4	Cyclin-Y	8,64
TRINITY_DN32367_c2_g1_i2	ERBB receptor feedback inhibitor 1	8,64
TRINITY_DN27921_c2_g2_i7	B-cell receptor-associated protein 29	8,64
TRINITY_DN29960_c0_g1_i6 TRINITY_DN30927_c1_g3_i4	Ufm1-specific protease 2 Protein LYRIC	8,64 8,64
TRINITY_DN26987_c0_g1_i9	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	8,64
TRINITY_DN26611_c2_g2_i3	Ras and Rab interactor 3	8,67
TRINITY_DN30433_c0_g3_i2	F-box/LRR-repeat protein 17	8,67
TRINITY_DN31110_c1_g1_i3	Histone-lysine N-methyltransferase, H3 lysine-79 specific	8,67
TRINITY_DN31322_c0_g2_i3	Ran-binding protein 10	8,67
TRINITY_DN27362_c0_g2_i2	Synaptojanin-2-binding protein	8,67
	The state of the s	-,

TRINITY_DN3098_c2_g1_9 Pathion/transferase ZDHHC18 8,67 TRINITY_DN37893_c0_g1_i11 Mamonse-binding protein 8,67 TRINITY_DN33697_c1_g1_i2 Protein timspoor protein Sec24D 8,67 TRINITY_DN33697_c1_g1_i2 Protein timspoor protein Sec24D 8,67 TRINITY_DN32978_c1_g1_i2 Protein dissulfide-isomerase A6 8,67 TRINITY_DN32978_c1_g1_i2 Fritzle-8 8,67 TRINITY_DN32620_c0_g1_i1 Zymin-specific demethyluses pht2 8,67 TRINITY_DN32620_c0_g1_i1 TRINITY_DN32620_c0_g1_i1 Trinity_DN3283_c1_g1_i1 Trinity_DN3283	TD1) WTV D1(2007 2 2 10	D 1 1 1 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	0.67
TRINITY_DN32890_c1_g2_i1	_		
TRINITY DN32983_cl_g2_12			
TRINITY_DN31990_cl_gl_i1		~ *	
TRINITY DN2700 c.O. g. i.i. Queusoine salvage protein S.67			
TRINITY DN3260_c0_g1.i1		*	
TRINITY DN32620_c0_g2_i13	TRINITY_DN29783_c1_g2_i2	Protein disulfide-isomerase A6	8,67
TRINITY_DN31281_c2_gl_i2	TRINITY_DN22700_c0_g1_i1	Queuosine salvage protein	8,67
RRINITY_DN30283_c_g_l_i1	TRINITY_DN32620_c0_g2_i13	Lysine-specific demethylase phf2	8,67
TRINITY_DN30482_cl_gl_i1	TRINITY_DN31971_c2_g1_i2	Frizzled-8	8,67
TRINITY_DN30482_cl_gl_i1	TRINITY DN29271 c0 g1 i1	AT-rich interactive domain-containing protein 5B	8,69
TRINITY_DN3040_C0_g1_i1 Sarcoplasmic/endoplasmic reticulum calcium ATPase 1 8,69 TRINITY_DN2886_C0_g2_i1 Miogen-activated protein kinase kinase kinase 13 8,69 TRINITY_DN2885_C0_g1_i1 Transposm Ty3-G Gag_Pol polyprotein 8,69 TRINITY_DN3175_C0_g1_i1 Long-chain fatty acid transport protein 8,69 TRINITY_DN3165_C0_g2_i4 Eukaryotic translation initiation factor 4E transporter 8,69 TRINITY_DN3167_C0_g1_i16 Commide synthase 8,69 TRINITY_DN3167_C0_g1_i16 Commide synthase 8,69 TRINITY_DN3167_C0_g1_i16 Commide synthase 8,69 TRINITY_DN3167_C0_g1_i16 Commide synthase 8,69 TRINITY_DN3167_C0_g1_i6 Cacin 8,69 TRINITY_DN3167_C0_g1_i6 Cacin 8,69 TRINITY_DN3167_C0_g1_i6 Cacin 8,69 TRINITY_DN3168_C0_g2_i3 Ras GTPase-activating protein 8,69 TRINITY_DN32601_C0_g1_i7 Eukaryotic translation-associated protein 8,69 Axin interactor, dorsalization-associated protein 8,69 TRINITY_DN32601_C0_g1_i7 Eukapylasmic reticulum mannosyl-oligosaccharide 1,2-alpha-ma 8,72 TRINITY_DN32601_C0_g1_i7 Eukapylasmic reticulum mannosyl-oligosaccharide 1,2-alpha-ma 8,72 TRINITY_DN3107_C0_g1_i1 Eukapylasmic reticulum mannosyl-oligosaccharide 1,2-alpha-ma 8,72 Echinoderm microtubule-associated protein-like 3 8,72 Echinoderm microtubule-associated protein-like 3 8,72 Echinoderm microtubule-associated protein-like 3 8,72 Echinoderm microtubule-associated protein 8,			
RRINITY_DN28896_C0_g2_i1	_	•	
TRINITY_DN28256_C0_g1_i1			
TRINITY_DN31752_c0_g1_i1	_		
TRINITY_DN31452_c0_gl_i1	_	ž .	
TRINITY_DN3145_0_g_1_i15	_		
TRINITY_DN31367_c0_gl_i16	_	•	
TRINITY_DN3137_c0_gl_i6			
TRINITY_DN36512_c0_g1_i6	_		
TRINITY_DN33137_c5_g1_i8 TRINITY_DN31313_c1_g1_g1_i4 Retinol dehydrogenase 8 8.69 TRINITY_DN31483_02_g1_i7 TRINITY_DN32691_c2_g1_i7 Palmitoyltransferase ZDHHC5 RENITY_DN29699_c0_g1_i1 TRINITY_DN29699_c0_g1_i1 TRINITY_DN32303_c0_g1_i7 TRINITY_DN32303_c0_g1_i7 TRINITY_DN32959_c4_g1_i10 TRINITY_DN31177_c0_g1_i17 SEC14-like protein 2 TRINITY_DN31177_c0_g1_i17 SEC14-like protein 2 TRINITY_DN31177_c0_g1_i17 TRINITY_DN3194_c2_g1_i9 Arithmenine-protein kinase N2 TRINITY_DN3194_c2_g1_i3 Uncharacterized protein KIAA0930 homolog R72 TRINITY_DN32425_c0_g1_i3 TRINITY_DN32425_c0_g1_i3 TRINITY_DN32425_c0_g1_i3 TRINITY_DN32425_c0_g1_i4 TRINITY_DN32301_cd_g1_i4 TRINITY_DN32301_cd_g1_i4 TRINITY_DN32301_cd_g1_i4 TRINITY_DN32301_cd_g1_i4 TRINITY_DN23051_c0_g1_i4 TRINITY_DN23051_c0_g1_i4 TRINITY_DN25063_c0_g1_i2 TRINITY_DN2698_c0_g1_i2 TRINITY_DN2698_c0_g1_i2 TRINITY_DN28066_c0_g1_i4 TRINITY_DN28066_c0_g1_i4 TRINITY_DN28060_c1_g1_i1 TRINITY_DN28060_c1_g1_i1 TRINITY_DN28060_c1_g1_i1 TRINITY_DN28060_c1_g1_i1 TRINITY_DN28060_c0_g1_i4 TRINITY_DN28060_c0_g1_i4 TRINITY_DN28060_c0_g1_i4 TRINITY_DN28060_c0_g1_i4 TRINITY_DN28060_c0_g1_i5 TRINITY_DN28060_c0_g1_i4 TRINITY_DN28060_c0_g1_i5 TRINITY_DN28060_c0_g1_i6 Activating transcription factor 7-interacting protein 1 R75 TRINITY_DN29091_c0_g1_i6 Activating transcription factor 11 R75 TRINITY_DN209091_c1_g1_i6 Activating transcription factor 11 R75 TRINITY_DN209131_5_c1_g1_i4 Activating transcrip	_	•	
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TRINITY_DN25972_c0_g1_i1	TRINITY_DN29579_c4_g1_i10	Serine/threonine-protein kinase N2	8,72
TRINITY_DN31394_c2_g1_i9	TRINITY_DN31177_c0_g1_i17	SEC14-like protein 2	8,72
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TRINITY_DN25363_c0_g1_i4	_		
TRINITY_DN23051_c0_g1_i4			
TRINITY_DN26960_c1_g2_i10 Proliferation-associated protein 2G4 8,72 TRINITY_DN29698_c0_g1_i2 H-2 class II histocompatibility antigen, I-A beta chain 8,72 TRINITY_DN26737_c5_g1_i4 Protein transport protein Sec31A 8,75 TRINITY_DN28066_c0_g1_i4 Mothers against decapentaplegic homolog 9 8,75 TRINITY_DN25193_c0_g2_i8 Small glutamine-rich tetratricopeptide repeat-containing protein 8,75 TRINITY_DN28070_c1_g1_i12 Dystonin 8,75 TRINITY_DN25027_c0_g1_i5 R3H domain-containing protein 2 8,75 TRINITY_DN28080_c0_g1_i4 Type I phosphatidylinositol 4,5-bisphosphate 4-phosphatase-B 8,75 TRINITY_DN26252_c1_g1_i8 Focal adhesion kinase 1 8,75 TRINITY_DN26983_c0_g1_i12 Serine/arginine-rich splicing factor 11 8,75 TRINITY_DN27249_c1_g2_i4 Gag-Pol polyprotein 8,75 TRINITY_DN25974_c0_g1_i6 Activating transcription factor 7-interacting protein 1 8,75 TRINITY_DN30423_c0_g1_i2 Cytochrome P450 1A3 8,75 TRINITY_DN32011_c0_g1_i3 Receptor-type tyrosine-protein phosphatase alpha 8,75 TRINITY_DN26991_c1_g1_i6 Low-density lipoprotein receptor-related protein 1 8,75 TRINITY_DN29283_c0_g1_i1 Nuclear receptor-interacting protein 1 8,75 TRINITY_DN31315_c1_g1_i4 Transmembrane and coiled-coil domains protein 1 8,75		ž v v	
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TRINITY_DN25193_c0_g2_i8			
TRINITY_DN28070_c1_g1_i12 Dystonin			
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TRINITY_DN25974_c0_g1_i6 Activating transcription factor 7-interacting protein 1 TRINITY_DN30423_c0_g1_i2 Cytochrome P450 1A3 TRINITY_DN32011_c0_g1_i3 Receptor-type tyrosine-protein phosphatase alpha R75 TRINITY_DN26991_c1_g1_i6 Low-density lipoprotein receptor-related protein 1 TRINITY_DN29283_c0_g1_i1 Nuclear receptor-interacting protein 1 TRINITY_DN31315_c1_g1_i4 Transmembrane and coiled-coil domains protein 1 8,75	TRINITY_DN26983_c0_g1_i12	Serine/arginine-rich splicing factor 11	8,75
TRINITY_DN30423_c0_g1_i2 Cytochrome P450 1A3 8,75 TRINITY_DN32011_c0_g1_i3 Receptor-type tyrosine-protein phosphatase alpha 8,75 TRINITY_DN26991_c1_g1_i6 Low-density lipoprotein receptor-related protein 1 8,75 TRINITY_DN29283_c0_g1_i1 Nuclear receptor-interacting protein 1 8,75 TRINITY_DN31315_c1_g1_i4 Transmembrane and coiled-coil domains protein 1 8,75	TRINITY_DN27249_c1_g2_i4	Gag-Pol polyprotein	8,75
TRINITY_DN32011_c0_g1_i3Receptor-type tyrosine-protein phosphatase alpha8,75TRINITY_DN26991_c1_g1_i6Low-density lipoprotein receptor-related protein 18,75TRINITY_DN29283_c0_g1_i1Nuclear receptor-interacting protein 18,75TRINITY_DN31315_c1_g1_i4Transmembrane and coiled-coil domains protein 18,75	TRINITY_DN25974_c0_g1_i6	Activating transcription factor 7-interacting protein 1	8,75
TRINITY_DN32011_c0_g1_i3Receptor-type tyrosine-protein phosphatase alpha8,75TRINITY_DN26991_c1_g1_i6Low-density lipoprotein receptor-related protein 18,75TRINITY_DN29283_c0_g1_i1Nuclear receptor-interacting protein 18,75TRINITY_DN31315_c1_g1_i4Transmembrane and coiled-coil domains protein 18,75	TRINITY_DN30423_c0_g1_i2		8,75
TRINITY_DN26991_c1_g1_i6Low-density lipoprotein receptor-related protein 18,75TRINITY_DN29283_c0_g1_i1Nuclear receptor-interacting protein 18,75TRINITY_DN31315_c1_g1_i4Transmembrane and coiled-coil domains protein 18,75	_	•	
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TRINITY_DN31315_c1_g1_i4 Transmembrane and coiled-coil domains protein 1 8,75	_		
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TD 11 17 D 1100 CCT 0 4 144		0.5
TRINITY_DN32665_c0_g1_i11	Immunoglobulin superfamily member 8	8,76
TRINITY_DN29721_c1_g1_i8	Transmembrane protein 184C	8,77
TRINITY_DN32852_c1_g1_i16	Selenoprotein Pb	8,77
TRINITY_DN31886_c0_g1_i3	La-related protein 4	8,77
TRINITY_DN27182_c0_g1_i13	Small G protein signaling modulator 3	8,77
TRINITY_DN29794_c1_g1_i6	GATOR complex protein DEPDC5	8,77
TRINITY_DN23260_c0_g1_i1	Leptin receptor	8,77
TRINITY_DN30890_c1_g1_i3	Endoplasmic reticulum membrane sensor NFE2L1	8,77
TRINITY_DN30983_c0_g1_i1	Interleukin-17 receptor C	8,77
TRINITY_DN33123_c2_g1_i1	Nucleoside diphosphate kinase 7	8,77
TRINITY_DN26063_c1_g1_i14	SH2B adapter protein 3	8,77
TRINITY_DN27148_c1_g3_i6	Coiled-coil domain-containing protein 50	8,77
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TRINITY_DN30073_c1_g1_i5	Adenylate cyclase type 9	8,77
TRINITY_DN28648_c1_g1_i3	Insulin-like growth factor 2 mRNA-binding protein 2	8,77
TRINITY_DN30082_c0_g1_i3	Nuclear migration protein nudC	8,77
TRINITY_DN28927_c1_g1_i4	Proline-, glutamic acid- and leucine-rich protein 1	8,77
TRINITY_DN28602_c0_g1_i7	Cell surface glycoprotein MUC18	8,77
TRINITY_DN32508_c0_g3_i7	Rap guanine nucleotide exchange factor 2	8,77
TRINITY_DN28343_c1_g1_i7	Synaptotagmin-like protein 1	8,80
TRINITY_DN27403_c0_g1_i3	Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial	8,80
TRINITY_DN27432_c0_g1_i1	Nucleolar MIF4G domain-containing protein 1	8,80
TRINITY_DN27667_c1_g1_i15	Prolyl 3-hydroxylase OGFOD1	8,80
TRINITY_DN30610_c0_g1_i9	Mitochondrial fission regulator 1	8,80
TRINITY_DN23787_c2_g2_i2	RING finger protein 215	8,80
TRINITY_DN26361_c0_g1_i2	Protein ABHD11	8,80
TRINITY_DN29123_c2_g1_i18	LON peptidase N-terminal domain and RING finger protein 2	8,80
TRINITY_DN25531_c4_g1_i3	Transcription factor 12	8,80
TRINITY_DN28878_c3_g2_i2	Aladin	8,80
TRINITY_DN33321_c5_g1_i14	Dystonin	8,80
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TRINITY_DN31029_c3_g3_i4	Betainehomocysteine S-methyltransferase 1	8,80
TRINITY_DN24772_c4_g3_i11	GPI mannosyltransferase 1	8,80
TRINITY_DN26415_c0_g1_i2	Tetratricopeptide repeat protein 36	8,80
TRINITY_DN28022_c2_g3_i3	SAP30-binding protein	8,80
TRINITY_DN28165_c2_g3_i5	Ribosomal protein S6 kinase alpha-1	8,80
TRINITY_DN30958_c0_g1_i11	Cytochrome c-type heme lyase	8,80
TRINITY_DN28177_c2_g2_i11	Tumor necrosis factor receptor superfamily member 5	8,80
TRINITY_DN25505_c1_g1_i4	Complement factor B	8,80
TRINITY_DN33174_c9_g2_i3	1,25-dihydroxyvitamin $D(3)$ 24 -hydroxylase, mitochondrial	8,82
TRINITY_DN30272_c0_g2_i3	Retinoblastoma-binding protein 5	8,82
TRINITY_DN30503_c1_g1_i12	Protein sel-1 homolog 1	8,82
TRINITY_DN26182_c1_g1_i4	Transmembrane protein 19	8,82
TRINITY_DN23772_c0_g1_i4	Carbohydrate sulfotransferase 14	8,82
TRINITY_DN24641_c0_g1_i4	Serpin H1	8,82
TRINITY_DN25606_c0_g1_i1	Class I histocompatibility antigen, F10 alpha chain	8,82
TRINITY_DN31042_c2_g1_i2	D-aspartate oxidase	8,82
TRINITY_DN29451_c0_g1_i4	Zinc transporter 9	8,82
TRINITY_DN27995_c0_g1_i11	Nuclear respiratory factor 1	8,82
TRINITY_DN27973_c0_g1_i9	PHD finger protein 3	8,82
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TRINITY_DN31748_c0_g1_i3	Acetyl-CoA acetyltransferase, mitochondrial	8,82
TRINITY_DN26379_c1_g3_i2	Stonustoxin subunit beta	8,82
TRINITY_DN24895_c4_g1_i6	Probable RNA-binding protein EIF1AD	8,82
TRINITY_DN28106_c1_g1_i1	B-cell lymphoma 6 protein homolog	8,82
TRINITY_DN29018_c0_g2_i11	Cytosolic purine 5'-nucleotidase	8,82
TRINITY_DN30968_c2_g1_i6	Ubiquitin carboxyl-terminal hydrolase 47	8,85
TRINITY_DN24863_c2_g1_i3	BCL2/adenovirus E1B 19 kDa protein-interacting protein 3	8,85
TRINITY_DN26920_c0_g1_i16	Katanin p80 WD40 repeat-containing subunit B1	8,85
TRINITY_DN31647_c0_g1_i15	Methionine synthase	8,85
TRINITY_DN31685_c1_g1_i6	Protein FAM126B	8,85
TRINITY_DN24766_c0_g1_i6	Limbin	8,85

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TRINITY_DN28683_c0_g1_i3	PWWP domain-containing protein 2A	8,85
TRINITY_DN32598_c1_g1_i1	FACT complex subunit SPT16	8,85
TRINITY_DN28498_c1_g1_i8	Kynureninase	8,85
TRINITY_DN28892_c0_g3_i2	Guanine nucleotide-binding protein $G(t)$ subunit alpha-1	8,85
TRINITY_DN29178_c2_g1_i12	Serine hydroxymethyltransferase, mitochondrial	8,85
TRINITY_DN26872_c0_g1_i2	Double-stranded RNA-binding protein Staufen homolog 1	8,85
TRINITY_DN30467_c0_g1_i5	Cysteine-rich with EGF-like domain protein 2	8,85
TRINITY_DN32706_c0_g1_i4	1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase epsi	8,85
TRINITY_DN27802_c0_g1_i4	Suppression of tumorigenicity 5 protein	8,85
TRINITY_DN31413_c2_g1_i2	mRNA export factor	8,85
TRINITY_DN28179_c1_g1_i18	Poly(A) polymerase gamma	8,85
TRINITY_DN28994_c0_g1_i7	Ras-related protein Rab-22A	8,87
TRINITY_DN27938_c0_g1_i7		8,87
	Autophagy-related protein 13	
TRINITY_DN29832_c3_g1_i2	Rho guanine nucleotide exchange factor 6	8,87
TRINITY_DN28315_c0_g1_i9	Protein MB21D2	8,87
TRINITY_DN24161_c3_g2_i5	Kelch-like protein 28	8,87
TRINITY_DN32507_c1_g1_i15	TOM1-like protein 2	8,87
TRINITY_DN28864_c2_g1_i9	CCR4-NOT transcription complex subunit 1	8,87
TRINITY_DN32372_c2_g1_i1	Protein Shroom1	8,87
TRINITY_DN31510_c0_g1_i4	FYVE, RhoGEF and PH domain-containing protein 5	8,87
TRINITY_DN28024_c2_g1_i4	Guanine nucleotide-binding protein subunit alpha-11	8,87
TRINITY_DN30762_c8_g1_i1	U2 snRNP-associated SURP motif-containing protein	8,87
TRINITY_DN24384_c0_g1_i1	Serine/threonine-protein kinase 17A	8,87
TRINITY_DN31065_c3_g1_i10	Cytoskeleton-associated protein 4	8,87
TRINITY_DN30271_c0_g3_i1	14-3-3 protein beta/alpha-A	8,87
TRINITY_DN22913_c0_g1_i3	Armadillo repeat-containing protein 8	8,87
TRINITY_DN31193_c0_g1_i7	Guanine nucleotide-binding protein G(i) subunit alpha-2	8,90
TRINITY_DN28371_c1_g4_i21	Reticulocalbin-1	8,90
TRINITY_DN31980_c1_g1_i27	Cytochrome P450 2J2	8,90
TRINITY_DN28016_c0_g1_i3	Thyroid receptor-interacting protein 6	8,90
TRINITY_DN20829_c0_g1_i2	Activated CDC42 kinase 1	8,90
TRINITY_DN30414_c0_g2_i9	Serine/threonine-protein kinase D2	8,90
TRINITY_DN32808_c2_g1_i9	Cell death activator CIDE-A	8,90
TRINITY_DN32072_c0_g2_i4	Cyclic AMP receptor-like protein A	8,90
TRINITY_DN27203_c2_g1_i7	3'(2'),5'-bisphosphate nucleotidase 1	8,90
TRINITY_DN28185_c0_g1_i1	DnaJ homolog subfamily C member 1	8,90
TRINITY_DN23495_c0_g1_i10	Fatty acid-binding protein, heart	8,90
TRINITY_DN31490_c2_g1_i20	Transposon Tf2-9 polyprotein	8,90
TRINITY_DN25057_c4_g1_i4	Leucine-rich repeat-containing protein 58	8,90
TRINITY_DN31590_c0_g1_i3	Ectonucleoside triphosphate diphosphohydrolase 4	8,92
TRINITY_DN27545_c0_g1_i9	Rho guanine nucleotide exchange factor 7	8,92
TRINITY_DN30946_c0_g2_i2	Perforin-1	8,92
TRINITY_DN30131_c0_g2_i6	Signal transducing adapter molecule 1	8,92
TRINITY_DN31296_c1_g1_i4	Tectonin beta-propeller repeat-containing protein 2	8,92
TRINITY_DN30069_c0_g3_i2	Glycogen [starch] synthase, muscle	8,92
TRINITY_DN29890_c2_g1_i1	Endoplasmic reticulum aminopeptidase 1	8,92
TRINITY_DN32817_c2_g1_i1	Unconventional myosin-Vb	8,92
TRINITY_DN24405_c0_g1_i1	Gelsolin	8,92
TRINITY_DN24530_c0_g1_i6	Ras association domain-containing protein 1	8,92
TRINITY_DN28482_c1_g1_i8	Regulation of nuclear pre-mRNA domain-containing protein 2	8,92
TRINITY_DN26347_c0_g1_i5	SH3 domain-binding glutamic acid-rich-like protein 3	8,92
TRINITY_DN32676_c1_g2_i20	Phospholipid hydroperoxide glutathione peroxidase	8,92
<u> </u>		8,92 8,92
TRINITY_DN29537_c0_g3_i4	Complement factor H	
TRINITY_DN25316_c1_g2_i2	Lysophosphatidic acid receptor 1	8,92
TRINITY_DN32783_c3_g2_i5	Integrin beta-1	8,92
TRINITY_DN32974_c1_g1_i2	Kinesin-like protein KIF13B	8,94
TRINITY_DN23867_c1_g4_i2	Adenosylhomocysteinase 3	8,94
TRINITY_DN28879_c0_g1_i4	Protein unc-45 homolog A	8,94
TRINITY_DN27321_c1_g2_i1	Acidic leucine-rich nuclear phosphoprotein 32 family member E	8,94

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TRINITY_DN31843_c0_g1_i4	Multiple PDZ domain protein	8,94
TRINITY_DN25634_c2_g1_i1	NEDD8-conjugating enzyme UBE2F	8,94
TRINITY_DN32395_c1_g2_i7	Biogenesis of lysosome-related organelles complex 1 subunit 6	8,94
TRINITY_DN31880_c2_g1_i11	Growth factor receptor-bound protein 10	8,94
TRINITY_DN27008_c1_g1_i12	Serine/arginine-rich splicing factor 12	8,94
TRINITY_DN27843_c0_g1_i2	Protein PRR14L	8,97
TRINITY_DN26696_c1_g2_i1	SLIT-ROBO Rho GTPase-activating protein 3	8,97
TRINITY_DN25276_c0_g1_i3	Pleckstrin homology domain-containing family A member 2	8,97
TRINITY_DN26377_c0_g1_i2	Secretory carrier-associated membrane protein 3	8,97
TRINITY_DN14246_c0_g1_i1	Collagen alpha-1(XI) chain	8,97
TRINITY_DN30696_c1_g1_i19	Zinc finger CCCH domain-containing protein 4	8,97
TRINITY_DN28369_c1_g1_i9		8,97
_	Hippocampus abundant transcript 1 protein	
TRINITY_DN28091_c1_g3_i2	Alpha-1,2-mannosyltransferase ALG9	8,97
TRINITY_DN33119_c2_g1_i20	Syndecan-2	8,97
TRINITY_DN29234_c2_g1_i3	Probable ATP-dependent RNA helicase DDX27	8,97
TRINITY_DN27039_c0_g1_i1	Protein FAM234A	8,97
TRINITY_DN30796_c0_g1_i5	CDGSH iron-sulfur domain-containing protein 1	8,97
TRINITY_DN25897_c0_g1_i1	P2X purinoceptor 4	8,97
TRINITY_DN29129_c5_g1_i27	Homogentisate 1,2-dioxygenase	8,99
TRINITY_DN32784_c0_g2_i13	E3 ubiquitin-protein ligase RNF185	8,99
TRINITY_DN31881_c2_g1_i5	Calmodulin-regulated spectrin-associated protein 3	8,99
TRINITY_DN28549_c1_g1_i18	Fibronectin type III and SPRY domain-containing protein 2	8,99
TRINITY_DN33305_c3_g1_i8	Complement C3 (Fragment)	8,99
TRINITY_DN29226_c0_g1_i5	Membrane-bound transcription factor site-2 protease	8,99
TRINITY_DN29313_c4_g3_i22	Zinc finger protein ubi-d4	8,99
TRINITY_DN30748_c0_g1_i7	Zinc transporter ZIP9	8,99
TRINITY_DN30719_c0_g1_i10	Histone acetyltransferase KAT6B	8,99
TRINITY_DN27007_c5_g1_i5	Limb region 1 homolog-like protein	8,99
TRINITY_DN27018_c0_g1_i9		8,99
_	Uridine-cytidine kinase-like 1	
TRINITY_DN21126_c0_g1_i2	ATP synthase subunit f, mitochondrial	8,99
TRINITY_DN29120_c0_g2_i1	FACT complex subunit SSRP1	8,99
TRINITY_DN28797_c0_g1_i12	Serine/threonine-protein kinase ppk4	8,99
TRINITY_DN32203_c3_g1_i3	Serine/threonine-protein kinase WNK1	9,01
TRINITY_DN31166_c0_g1_i13	Protein polybromo-1	9,01
TRINITY_DN31358_c0_g1_i5	Sprouty-related, EVH1 domain-containing protein 2	9,01
TRINITY_DN32945_c2_g2_i16	Tricarboxylate transport protein, mitochondrial	9,01
TRINITY_DN32477_c0_g1_i11	Myosin light chain kinase, smooth muscle (Fragment)	9,01
TRINITY_DN29383_c2_g1_i14	Carnitine O-acetyltransferase	9,01
TRINITY_DN33033_c1_g1_i9	DmX-like protein 1	9,01
TRINITY_DN30723_c0_g1_i3	Mitogen-activated protein kinase kinase kinase 3	9,01
TRINITY_DN31006_c2_g1_i10	Flotillin-2a	9,01
TRINITY_DN25897_c0_g1_i22	Copine-4	9,01
TRINITY_DN24678_c3_g3_i4	Protein ANTAGONIST OF LIKE HETEROCHROMATIN PROT	9,01
TRINITY_DN25778_c0_g3_i4	Zinc transporter Slc39a7 (Fragment)	9,01
TRINITY_DN25699_c0_g1_i9	Nucleolar protein 58	9,01
TRINITY_DN32712_c2_g1_i2	•	9,01
_	MRG/MORF4L-binding protein	
TRINITY_DN25731_c4_g1_i12	Phosphatidylinositol 3,4,5-trisphosphate 3-phosphatase and du	9,01
TRINITY_DN23495_c0_g1_i11	Fatty acid-binding protein, heart	9,01
TRINITY_DN33321_c5_g1_i16	Dystonin	9,01
TRINITY_DN26517_c1_g2_i7	Interferon alpha/beta receptor 2	9,01
TRINITY_DN32687_c1_g1_i6	Phosphatidylinositol 4-kinase alpha	9,03
TRINITY_DN31178_c2_g1_i11	Uridine-cytidine kinase-like 1	9,03
TRINITY_DN25420_c1_g1_i4	Protein lifeguard 2	9,03
TRINITY_DN29828_c0_g1_i1	Dual specificity testis-specific protein kinase 2	9,03
TRINITY_DN31165_c3_g1_i9	Thyroid hormone receptor-associated protein 3	9,03
TRINITY_DN31137_c1_g2_i1	Calpain-1 catalytic subunit	9,03
TRINITY_DN28571_c2_g2_i16	Transducin-like enhancer protein 3-B	9,03
TRINITY_DN30247_c3_g1_i26	Tyrosine 3-monooxygenase	9,03
TRINITY_DN24917_c0_g1_i6	Solute carrier family 25 member 46	9,03

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TRINITY_DN31836_c1_g1_i6	SEC14-like protein 1	9,03
TRINITY_DN24380_c2_g1_i2	Coiled-coil domain-containing protein 3	9,03
TRINITY_DN32777_c0_g1_i9	Histone-lysine N-methyltransferase 2A	9,03
TRINITY_DN29247_c0_g1_i6	Vacuolar protein sorting-associated protein 13B	9,03
TRINITY_DN29767_c0_g1_i2	Exportin-2	9,03
TRINITY_DN26488_c0_g1_i5	Rho guanine nucleotide exchange factor 18	9,05
TRINITY_DN30210_c0_g1_i3	RING finger and SPRY domain-containing protein 1	9,05
TRINITY_DN26454_c0_g1_i5	DNA-binding protein RFX7	9,05
TRINITY_DN32106_c0_g2_i3	Epidermal growth factor receptor kinase substrate 8	9,05
TRINITY_DN30687_c0_g1_i17	Cyclin-G-associated kinase	9,05
TRINITY_DN32877_c0_g1_i2	Disabled homolog 1	9,05
TRINITY_DN29495_c2_g1_i3	Ubiquitin carboxyl-terminal hydrolase CYLD	9,05
_		
TRINITY_DN31893_c1_g2_i6	E3 ubiquitin-protein ligase MIB2	9,05
TRINITY_DN28357_c2_g3_i8	Zinc finger protein 711	9,05
TRINITY_DN29653_c1_g1_i6	Cold shock domain-containing protein E1	9,05
TRINITY_DN25271_c2_g1_i6	Choline-phosphate cytidylyltransferase B	9,07
TRINITY_DN23077_c0_g2_i2	Transposon Tf2-9 polyprotein	9,07
TRINITY_DN31838_c0_g1_i11	Focal adhesion kinase 1	9,07
TRINITY_DN33003_c1_g1_i16	Choline kinase alpha	9,07
TRINITY_DN33240_c9_g1_i11	Complement C3 (Fragment)	9,07
TRINITY_DN30242_c0_g1_i16	Rho guanine nucleotide exchange factor 1	9,07
TRINITY_DN29643_c1_g2_i1	Arylsulfatase D	9,07
TRINITY_DN32396_c0_g1_i10	Torsin-1A	9,07
TRINITY_DN28195_c0_g1_i2	Alpha-actinin-4	9,07
TRINITY_DN25140_c0_g1_i1	Cytoplasmic protein NCK1	9,07
TRINITY_DN31623_c0_g1_i8	Retrovirus-related Pol polyprotein from type-1 retrotransposab	9,07
		9,07
TRINITY_DN31703_c0_g1_i11	3 beta-hydroxysteroid dehydrogenase type 7	
TRINITY_DN29113_c0_g1_i9	Mitogen-activated protein kinase kinase kinase 10	9,09
TRINITY_DN30473_c3_g1_i8	Ecotropic viral integration site 5 protein homolog	9,09
TRINITY_DN25554_c0_g1_i17	Serine/threonine-protein phosphatase PP1-beta	9,09
TRINITY_DN28924_c0_g1_i1	RNA-binding protein 20	9,09
TRINITY_DN27852_c0_g1_i6	Hamartin	9,09
TRINITY_DN29123_c2_g1_i2	LON peptidase N-terminal domain and RING finger protein 2	9,09
TRINITY_DN32611_c1_g2_i8	Death-inducer obliterator 1	9,09
TRINITY_DN32302_c1_g1_i7	ATP-dependent RNA helicase DDX42	9,09
TRINITY_DN33238_c5_g1_i17	Telomerase protein component 1	9,09
TRINITY_DN23941_c0_g1_i1	Alpha-N-acetylglucosaminidase	9,09
TRINITY_DN29468_c1_g2_i5	U3 small nucleolar ribonucleoprotein protein MPP10	9,09
TRINITY_DN31262_c2_g1_i7	Solute carrier family 25 member 36-A	9,11
TRINITY_DN29921_c0_g1_i12	Partitioning defective 3 homolog	9,11
TRINITY_DN25515_c1_g2_i4	Monocarboxylate transporter 2	9,11
TRINITY_DN26877_c1_g5_i4	Elongation factor 1-delta	9,11
TRINITY_DN24822_c2_g1_i8		
	Vitelline membrane outer layer protein 1 homolog	9,11
TRINITY_DN29157_c0_g2_i4	RNA-binding motif, single-stranded-interacting protein 2	9,11
TRINITY_DN29135_c1_g1_i6	WD repeat-containing protein 26	9,11
TRINITY_DN25715_c4_g1_i9	Importin subunit alpha-3	9,11
TRINITY_DN31748_c0_g1_i17	Acetyl-CoA acetyltransferase, mitochondrial	9,11
TRINITY_DN32708_c2_g1_i1	Period circadian protein homolog 2	9,11
TRINITY_DN30348_c2_g1_i1	Keratinocyte-associated transmembrane protein 2	9,13
TRINITY_DN26252_c1_g1_i2	Focal adhesion kinase 1	9,13
TRINITY_DN32729_c0_g1_i9	Putative ATP-dependent RNA helicase DHX57	9,13
TRINITY_DN26102_c1_g1_i5	Zinc finger protein 24	9,13
TRINITY_DN30365_c1_g1_i19	DNA transposase THAP9	9,13
TRINITY_DN29870_c0_g1_i2	Legumain	9,13
TRINITY_DN28018_c4_g1_i7	L-xylulose reductase	9,13
TRINITY_DN27557_c2_g4_i4	Serine/threonine-protein kinase TAO1	9,13
TRINITY_DN32245_c0_g1_i13	Thioredoxin-interacting protein	9,13
TRINITY_DN26845_c2_g1_i10	Coagulation factor IX	9,13
TRINITY_DN32530_c1_g1_i6	DDB1- and CUL4-associated factor 6	9,15
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TDINITY DN22777 a0 a1 :1	History Insing N mothyltagasforage 24	9,15
TRINITY_DN32777_c0_g1_i1	Histone-lysine N-methyltransferase 2A	9,15
TRINITY_DN25915_c0_g1_i7 TRINITY_DN29938_c1_g1_i10	V-type proton ATPase catalytic subunit A	9,15
	Interleukin enhancer-binding factor 3 homolog	
TRINITY_DN25126_c0_g1_i1	Exportin-4	9,15
TRINITY_DN32311_c1_g1_i34	Histone deacetylase 4	9,15
TRINITY_DN23439_c0_g1_i1	Lysine-specific demethylase 6A	9,15
TRINITY_DN28779_c2_g1_i4	OCIA domain-containing protein 1	9,15
TRINITY_DN26016_c2_g1_i11	Carbohydrate sulfotransferase 10	9,15
TRINITY_DN29268_c5_g2_i7	CDK5 and ABL1 enzyme substrate 1	9,17
TRINITY_DN30712_c0_g1_i5	Sodium- and chloride-dependent GABA transporter 1	9,17
TRINITY_DN30876_c5_g1_i1	cGMP-inhibited 3',5'-cyclic phosphodiesterase B	9,17
TRINITY_DN30655_c2_g1_i12	Pseudouridylate synthase 7 homolog	9,17
TRINITY_DN25641_c3_g2_i4	Pyrroline-5-carboxylate reductase 2	9,17
TRINITY_DN30273_c0_g1_i4	Rab11 family-interacting protein 4B	9,17
TRINITY_DN23790_c0_g1_i2	Methylmalonic aciduria type A protein, mitochondrial	9,17
TRINITY_DN30883_c0_g2_i5	Polyadenylate-binding protein 1	9,19
TRINITY_DN29377_c2_g1_i9	Apolipoprotein B-100	9,19
TRINITY_DN30482_c0_g1_i4	Phosphatidylinositol 4-kinase beta	9,19
TRINITY_DN26570_c0_g2_i7	Plastin-3	9,19
TRINITY_DN32583_c1_g1_i10	Exostosin-1b	9,19
TRINITY_DN28430_c1_g1_i10	Mitochondrial uncoupling protein 2	9,19
TRINITY_DN33132_c1_g1_i6	Peptidyl-prolyl cis-trans isomerase FKBP5	9,21
TRINITY_DN30713_c3_g1_i6	Serine/threonine-protein kinase WNK1	9,21
TRINITY_DN24636_c4_g2_i1	Pleckstrin homology-like domain family B member 1 (Fragment	9,21
TRINITY_DN31589_c1_g1_i3	Coagulation factor V	9,21
TRINITY_DN24930_c2_g5_i17	Leukocyte cell-derived chemotaxin-2	9,21
TRINITY_DN27946_c0_g2_i3	Endoribonuclease LACTB2	9,21
TRINITY_DN28293_c0_g1_i4	Copine-8	9,21
TRINITY_DN28141_c1_g2_i8	PH domain leucine-rich repeat-containing protein phosphatase	9,21
TRINITY_DN31810_c0_g1_i1	Protein tweety homolog 3	9,21
TRINITY_DN29994_c1_g1_i3	LINE-1 retrotransposable element ORF2 protein	9,21
TRINITY_DN31175_c0_g1_i9	D-2-hydroxyglutarate dehydrogenase, mitochondrial	9,23
TRINITY_DN33320_c4_g1_i13	Protein PRRC2B	9,23
TRINITY_DN26239_c0_g1_i6	Adipocyte enhancer-binding protein 1	9,23
TRINITY_DN32243_c2_g1_i6	Centrosomal protein of 152 kDa	9,23
TRINITY_DN26302_c0_g1_i3	UDP-N-acetylglucosamine transporter	9,23
TRINITY_DN29458_c0_g1_i14	Phosphatidate phosphatase LPIN1	9,23
TRINITY_DN30321_c0_g1_i13	Nuclear transcription factor Y subunit gamma	9,23
TRINITY_DN24962_c0_g2_i5	Cytochrome c oxidase subunit 7A-related protein, mitochondria	9,23
TRINITY_DN32496_c0_g1_i9	Serine/threonine-protein phosphatase 1 regulatory subunit 10	9,25
TRINITY_DN23526_c0_g1_i4	Vacuole membrane protein 1	9,25
TRINITY_DN29178_c2_g2_i3	Ataxin-7-like protein 1	9,25
TRINITY_DN31407_c1_g1_i5	Peroxisomal sarcosine oxidase	9,25
TRINITY_DN26885_c0_g1_i4	Epidermal growth factor receptor kinase substrate 8-like protein	9,23
TRINITY_DN32912_c2_g2_i2		9,27
	Peroxisome proliferator-activated receptor alpha	
TRINITY_DN26323_c2_g1_i4	Ubiquitin domain-containing protein UBFD1	9,27
TRINITY_DN32477_c0_g1_i14	Myosin light chain kinase, smooth muscle (Fragment)	9,27
TRINITY_DN26565_c0_g2_i4	Lissencephaly-1 homolog A	9,27
TRINITY_DN27188_c0_g1_i2	Cohesin subunit SA-2	9,27
TRINITY_DN28519_c0_g1_i6	Serine/threonine-protein phosphatase 2A 56 kDa regulatory sul	9,29
TRINITY_DN27265_c1_g1_i6	Atlastin-2	9,29
TRINITY_DN28540_c1_g1_i18	HCLS1-associated protein X-1	9,29
TRINITY_DN26961_c0_g1_i2	RNA cytidine acetyltransferase	9,29
TRINITY_DN32556_c0_g1_i8	Lysine-specific demethylase 6A	9,29
TRINITY_DN27629_c0_g1_i2	Peroxidasin	9,29
TRINITY_DN28143_c0_g1_i10	Arginase-2, mitochondrial	9,29
TRINITY_DN24316_c1_g1_i2	Ubiquitin carboxyl-terminal hydrolase 4	9,29
TRINITY_DN31272_c0_g1_i4	Ubiquitin-protein ligase E3B	9,29
TRINITY_DN26528_c1_g3_i1	Mitogen-activated protein kinase kinase kinase 3	9,29

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TRINITY_DN32863_c0_g1_i15	Nischarin Christian Communication of the Communicat	9,29
TRINITY_DN26288_c0_g1_i9	Chromatin target of PRMT1 protein	9,29
TRINITY_DN29118_c0_g1_i8	Centrosomal protein of 63 kDa	9,30
TRINITY_DN23386_c0_g1_i2	Ribokinase	9,30
TRINITY_DN32628_c0_g2_i1	NCK-interacting protein with SH3 domain	9,30
TRINITY_DN29234_c2_g1_i2	Probable ATP-dependent RNA helicase DDX27	9,30
TRINITY_DN28637_c0_g1_i8	Solute carrier family 43 member 3	9,32
TRINITY_DN31903_c0_g1_i4	Serine/threonine-protein kinase ULK3	9,32
TRINITY_DN28497_c1_g1_i14	Globoside alpha-1,3-N-acetylgalactosaminyltransferase 1	9,32
TRINITY_DN25778_c0_g4_i1	Inositol-pentakisphosphate 2-kinase	9,34
TRINITY_DN29626_c0_g2_i4	Sorbin and SH3 domain-containing protein 1	9,34
TRINITY_DN24906_c0_g1_i1	ATP synthase $F(0)$ complex subunit C3, mitochondrial	9,34
TRINITY_DN32473_c0_g1_i6	Protein IMPACT-B	9,34
TRINITY_DN31513_c0_g1_i9	Peroxisomal targeting signal 1 receptor	9,36
TRINITY_DN28466_c1_g1_i10	Protein TOPAZ1	9,36
TRINITY_DN28880_c0_g1_i17	Insulin-induced gene 2 protein	9,36
TRINITY_DN27637_c0_g3_i4	Serine-protein kinase ATM	9,36
TRINITY_DN31936_c0_g2_i4	1-acyl-sn-glycerol-3-phosphate acyltransferase gamma	9,36
TRINITY_DN29176_c7_g1_i8	Ribosome-releasing factor 2, mitochondrial	9,36
TRINITY_DN27843_c0_g1_i8	Protein PRR14L	9,36
TRINITY_DN28797_c0_g1_i6	Serine/threonine-protein kinase ppk4	9,36
TRINITY_DN25482_c2_g1_i3	Bromodomain-containing protein 2	9,36
TRINITY_DN33303_c6_g1_i4	Cytochrome P450 2K6	9,36
TRINITY_DN31066_c0_g1_i2	Spermatogenesis-associated protein 13	9,36
TRINITY_DN27928_c0_g1_i4	Polynucleotide 5'-hydroxyl-kinase NOL9	9,37
TRINITY_DN31103_c0_g1_i7	Collagen type IV alpha-3-binding protein	9,37
TRINITY_DN27011_c0_g1_i6	UTPglucose-1-phosphate uridylyltransferase	9,37
TRINITY_DN29650_c0_g1_i11	Tyrosine-protein kinase JAK2	9,37
TRINITY_DN26658_c4_g1_i2	1-aminocyclopropane-1-carboxylate synthase-like protein 1	9,37
TRINITY_DN32079_c1_g2_i2	Cation-independent mannose-6-phosphate receptor	9,37
TRINITY_DN28419_c1_g2_i4	AN1-type zinc finger protein 5	9,37
TRINITY_DN23962_c0_g1_i12	ATP-dependent RNA helicase DHX33	9,37
TRINITY_DN27470_c0_g1_i2	Splicing factor, arginine/serine-rich 15 (Fragment)	9,37
TRINITY_DN33123_c2_g1_i20	Nucleoside diphosphate kinase 7	9,37
TRINITY_DN27936_c0_g1_i4	Protein phosphatase 1 regulatory subunit 1B	9,37
TRINITY_DN29663_c4_g1_i4	Aldehyde oxidase 1	9,39
TRINITY_DN27840_c0_g1_i8	Cell division cycle protein 27 homolog	9,39
TRINITY_DN29680_c0_g1_i3	Ornithine decarboxylase antizyme 2	9,39
TRINITY_DN29503_c2_g1_i4	General transcription and DNA repair factor IIH helicase subw	9,39
TRINITY_DN25791_c4_g1_i10	Mannose-P-dolichol utilization defect 1 protein	9,39
TRINITY_DN29977_c0_g1_i4	Bifunctional coenzyme A synthase	9,41
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TRINITY_DN32781_c0_g1_i7	SUN domain-containing	9,41
TRINITY_DN32763_c1_g1_i18	SOSS complex subunit B2	9,42
TRINITY_DN31683_c0_g1_i4	Acidic mammalian chitinase	9,42
TRINITY_DN24100_c2_g5_i8	Vesicle-associated membrane protein 2	9,42
TRINITY_DN32450_c0_g1_i8	Lysine-specific demethylase 2B	9,42
TRINITY_DN30909_c0_g2_i7	DNA-binding protein RFX5	9,44
TRINITY_DN25715_c4_g1_i7	Importin subunit alpha-3	9,44
TRINITY_DN22735_c0_g2_i2	Nucleolar complex protein 2 homolog	9,44
TRINITY_DN29258_c0_g2_i1	Rab GTPase-binding effector protein 1	9,44
TRINITY_DN28651_c0_g2_i13	NLR family member X1	9,44
TRINITY_DN28546_c0_g1_i10	Rho GTPase-activating protein 12	9,44
TRINITY_DN24536_c1_g1_i5	Eukaryotic translation initiation factor 4 gamma 1	9,46
TRINITY_DN32859_c1_g1_i7	Tight junction protein ZO-1	9,46
TRINITY_DN23235_c1_g4_i12	H-2 class I histocompatibility antigen, D-K alpha chain	9,46
TRINITY_DN26023_c0_g2_i3	Methylglutaconyl-CoA hydratase, mitochondrial	9,46
TRINITY_DN32903_c1_g2_i7	UPF0183 protein C16orf70	9,47
TRINITY_DN31886_c0_g1_i2	La-related protein 4	9,47
TRINITY_DN23920_c0_g1_i2	Retrovirus-related Pol polyprotein from transposon TNT 1-94	9,47

TRINITY_DN31471_c1_g2_i13	Protein KIAA0100	9,47
TRINITY_DN26970_c1_g1_i7	Phosphorylase b kinase gamma catalytic chain, skeletal muscle	9,47
TRINITY_DN30792_c1_g1_i1	Transcription factor SOX-4	9,47
TRINITY_DN26208_c0_g1_i2	Serine/threonine-protein kinase PAK 2	9,49
TRINITY_DN27544_c0_g1_i22	Dynamin-2	9,49
TRINITY_DN28923_c0_g1_i2	TATA-binding protein-associated factor 172	9,50
TRINITY_DN32395_c1_g2_i1	Biogenesis of lysosome-related organelles complex 1 subunit 6	9,50
TRINITY_DN32817_c2_g1_i11	Unconventional myosin-Vb	9,50
TRINITY_DN32746_c2_g1_i31	Nuclear receptor corepressor 1	9,50
TRINITY_DN29793_c0_g1_i1	Ectonucleotide pyrophosphatase/phosphodiesterase family mem	9,50
TRINITY_DN28080_c0_g1_i10		9,50
	Type I phosphatidylinositol 4,5-bisphosphate 4-phosphatase-B	
TRINITY_DN31451_c1_g2_i1	F-box/WD repeat-containing protein 11	9,50
TRINITY_DN25606_c0_g3_i2	Major histocompatibility complex class I-related gene protein	9,50
TRINITY_DN32781_c0_g1_i6	SUN domain-containing	9,50
TRINITY_DN25473_c3_g1_i4	S-adenosylmethionine synthase isoform type-1	9,50
TRINITY_DN32865_c1_g1_i31	HLA class I histocompatibility antigen, alpha chain F	9,50
TRINITY_DN32738_c1_g1_i2	Ornithine decarboxylase	9,52
TRINITY_DN26425_c3_g1_i4	Translocation protein SEC63 homolog	9,53
TRINITY_DN31708_c0_g1_i15	Transmembrane protein 181	9,53
TRINITY_DN28498_c1_g1_i7	Kynureninase	9,55
TRINITY_DN30748_c0_g1_i8	Zinc transporter ZIP9	9,55
TRINITY_DN30984_c0_g1_i7	Ephexin-1	9,56
TRINITY_DN28925_c2_g2_i14	[Pyruvate dehydrogenase (acetyl-transferring)] kinase isozyme	9,56
TRINITY_DN33224_c10_g1_i14	Equilibrative nucleoside transporter 1	9,56
TRINITY_DN28332_c1_g3_i1	Gem-associated protein 6	9,58
TRINITY_DN33238_c5_g2_i2	TOM1-like protein 2	9,58
TRINITY_DN29636_c0_g2_i9	Plakophilin-3	9,58
TRINITY_DN22389_c0_g1_i2		9,58
	ER membrane protein complex subunit 1	
TRINITY_DN26699_c1_g2_i11	Exocyst complex component 6B	9,58
TRINITY_DN30184_c0_g1_i2	Bone morphogenetic protein receptor type-2	9,59
TRINITY_DN28239_c0_g1_i5	RNA exonuclease 1 homolog	9,59
TRINITY_DN26853_c1_g1_i4	DNA-directed RNA polymerase III subunit RPC8	9,61
TRINITY_DN33041_c1_g1_i9	Proteasome activator complex subunit 4B	9,61
TRINITY_DN31027_c2_g2_i12	Erlin-2	9,61
TRINITY_DN32253_c3_g2_i7	D-glucuronyl C5-epimerase	9,61
TRINITY_DN30348_c2_g1_i6	Keratinocyte-associated transmembrane protein 2	9,61
TRINITY_DN31165_c3_g1_i3	Thyroid hormone receptor-associated protein 3	9,61
TRINITY_DN30206_c1_g1_i4	Catenin beta-1	9,61
TRINITY_DN27855_c1_g1_i9	Terminal uridylyltransferase 4	9,61
TRINITY_DN30708_c1_g2_i15	MOB kinase activator 2	9,61
TRINITY_DN28526_c1_g1_i7	Rab5 GDP/GTP exchange factor	9,61
TRINITY_DN32242_c0_g2_i3	Baculoviral IAP repeat-containing protein 2	9,62
TRINITY_DN32978_c0_g1_i15	ValinetRNA ligase	9,62
TRINITY_DN30067_c1_g1_i7	Ubiquitin carboxyl-terminal hydrolase 42	9,62
TRINITY_DN29732_c1_g1_i4	Phosphatidate phosphatase LPIN2	9,62
TRINITY_DN27726_c2_g1_i9	Coatomer subunit zeta-1	9,62
TRINITY_DN26415_c0_g1_i4	Tetratricopeptide repeat protein 36	9,64
TRINITY_DN25956_c0_g2_i2	Actin-related protein 2-B	9,64
TRINITY_DN26327_c0_g1_i3	Rho-related GTP-binding protein RhoQ	9,64
TRINITY_DN25171_c0_g1_i10	Complement C3 (Fragment)	9,65
TRINITY_DN30227_c0_g1_i5	ATP-binding cassette sub-family C member 9	9,65
TRINITY_DN33001_c0_g1_i3	G protein-coupled receptor kinase 5	9,65
TRINITY_DN25960_c3_g1_i2	Retrovirus-related Pol polyprotein from transposon 297	9,66
TRINITY_DN28435_c1_g1_i6	Probable ubiquitin carboxyl-terminal hydrolase FAF-X	9,66
TRINITY_DN30797_c1_g1_i1	Coenzyme Q-binding protein COQ10 homolog, mitochondrial	9,66
TRINITY_DN31879_c0_g2_i13	Heterogeneous nuclear ribonucleoprotein A1	9,66
TRINITY_DN33118_c2_g1_i1	Protein PRRC2A	9,68
TRINITY_DN27960_c0_g1_i1	Protoporphyrinogen oxidase	9,68
TRINITY_DN31838_c0_g1_i20	Focal adhesion kinase 1	9,68
11d1 11 1 _D1 10 10 50 _C0 _g1 _120	1 COM MINUSCON INNUSC 1	7,00

TD D WTW D W 2000 0 1 11		0.60
TRINITY_DN23800_c0_g1_i1	Collagen alpha-3(VI) chain	9,68
TRINITY_DN24647_c0_g1_i1	E3 ubiquitin-protein ligase znrf3	9,69
TRINITY_DN31786_c0_g1_i3	Crooked neck-like protein 1	9,69
TRINITY_DN25606_c0_g6_i2	Major histocompatibility complex class I-related gene protein	9,71
TRINITY_DN30162_c1_g1_i3	Transport and Golgi organization protein 1 homolog	9,71
TRINITY_DN25796_c0_g1_i20	FERM, ARHGEF and pleckstrin domain-containing protein 2	9,71
TRINITY_DN32224_c5_g1_i8	Mitochondrial glycine transporter A	9,71
TRINITY_DN32975_c0_g2_i1	E1A-binding protein p400	9,71
TRINITY_DN30752_c1_g1_i1	Signal recognition particle subunit SRP68	9,71
TRINITY_DN32033_c0_g1_i6	MLX-interacting protein	9,72
TRINITY_DN32880_c1_g2_i9	Golgin subfamily A member 2	9,72
TRINITY_DN31975_c0_g1_i2	Dynamin-like 120 kDa protein, mitochondrial	9,72
TRINITY_DN29223_c0_g1_i2	Keratin, type I cytoskeletal 18	9,72
TRINITY_DN31368_c1_g3_i5	MAP/microtubule affinity-regulating kinase 3	9,73
TRINITY_DN28410_c2_g1_i15	Immunoglobulin mu heavy chain	9,73
TRINITY_DN30544_c0_g1_i2	Junctional adhesion molecule A	9,76
TRINITY_DN29537_c0_g3_i13	Complement factor H	9,76
TRINITY_DN26920_c0_g1_i15	Katanin p80 WD40 repeat-containing subunit B1	9,76
TRINITY_DN32936_c1_g2_i3	Uncharacterized protein KIAA0232	9,77
TRINITY_DN25558_c0_g3_i3	Transcription factor AP-1	9,77
TRINITY_DN32660_c0_g1_i27	Protein scribble homolog	9,78
TRINITY_DN30999_c0_g1_i3	Cullin-5	9,78
TRINITY_DN29302_c1_g2_i13	FGGY carbohydrate kinase domain-containing protein	9,78
TRINITY_DN25164_c2_g2_i14	Methylthioribose-1-phosphate isomerase	9,78
TRINITY_DN32746_c2_g1_i15	Nuclear receptor corepressor 1	9,80
TRINITY_DN24377_c1_g2_i12	Pancreatic progenitor cell differentiation and proliferation factor	9,80
TRINITY_DN29607_c0_g1_i3		9,80
_	Magnesium transporter protein 1	
TRINITY_DN29147_c1_g2_i3	Inhibin beta B chain	9,81
TRINITY_DN29146_c0_g1_i16	Apoptosis-inducing factor 1, mitochondrial	9,82
TRINITY_DN32687_c1_g1_i8	Phosphatidylinositol 4-kinase alpha	9,83
TRINITY_DN32582_c0_g1_i5	RING finger protein 10	9,83
TRINITY_DN28808_c0_g1_i7	Phosphatidate phosphatase LPIN2	9,83
TRINITY_DN26825_c0_g1_i3	Uncharacterized protein FAM241A	9,83
TRINITY_DN25694_c1_g1_i1	Ubiquitin-conjugating enzyme E2 N	9,83
TRINITY_DN28341_c0_g1_i11	SEC23-interacting protein	9,85
TRINITY_DN32452_c0_g1_i13	Ubiquitin carboxyl-terminal hydrolase 32	9,85
TRINITY_DN27799_c0_g1_i9	CUGBP Elav-like family member 1	9,86
TRINITY_DN29898_c0_g2_i6	Pleckstrin homology domain-containing family G member 5	9,86
TRINITY_DN27106_c1_g2_i18	Flavin-containing monooxygenase FMO GS-OX-like 5	9,86
TRINITY_DN25778_c0_g3_i1	Zinc transporter Slc39a7 (Fragment)	9,86
TRINITY_DN28730_c0_g1_i4	CCAAT/enhancer-binding protein gamma	9,86
TRINITY_DN26220_c0_g1_i2	Transcription factor SPT20 homolog	9,86
TRINITY_DN26853_c1_g1_i1	DNA-directed RNA polymerase III subunit RPC8	9,86
TRINITY_DN30314_c0_g2_i3	Heat shock factor protein 1	9,87
TRINITY_DN31523_c0_g1_i12	Phosphorylase b kinase regulatory subunit alpha, skeletal musc	9,87
TRINITY_DN32835_c0_g2_i12	Fibronectin type-III domain-containing protein 3a	9,87
TRINITY_DN30477_c0_g1_i2	A-kinase anchor protein 12	9,87
TRINITY_DN26830_c2_g2_i4	Tetratricopeptide repeat protein 17	9,88
TRINITY_DN31561_c2_g1_i18	Alanine aminotransferase 2-like	9,88
TRINITY_DN28512_c0_g1_i1	HLA class I histocompatibility antigen, alpha chain F	9,89
TRINITY_DN28782_c0_g1_i3	H-2 class II histocompatibility antigen, E-D alpha chain	9,89
TRINITY_DN29032_c1_g2_i2	Microtubule-associated proteins 1A/1B light chain 3A	9,89
TRINITY_DN24236_c2_g1_i12	Cadherin EGF LAG seven-pass G-type receptor 3	9,89
TRINITY_DN25162_c0_g1_i8	Geranylgeranyl transferase type-1 subunit beta	9,91
TRINITY_DN32015_c1_g1_i9	4-hydroxyphenylpyruvate dioxygenase	9,91
TRINITY_DN26891_c0_g1_i8	Lipoprotein lipase	9,92
TRINITY_DN29589_c1_g1_i9	Patatin-like phospholipase domain-containing protein 2	9,94
TRINITY_DN31535_c1_g1_i2	Transposon TX1 uncharacterized 149 kDa protein	9,94
TRINITY_DN32882_c1_g1_i4	Inositol 1,4,5-trisphosphate receptor type 1	9,95
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TDINITY DN21500 at at :15	Consulation factor V	9,95
TRINITY_DN31589_c1_g1_i15	Coagulation factor V	
TRINITY_DN33029_c2_g1_i3	Dihydropyrimidine dehydrogenase [NADP(+)]	9,96
TRINITY_DN25244_c0_g1_i3	Protein canopy-1	9,96
TRINITY_DN26523_c0_g1_i3	Insulin receptor substrate 1-B	9,96
TRINITY_DN32685_c2_g1_i2	Erbin	9,98
TRINITY_DN31994_c0_g1_i7	Rho-associated protein kinase 1	9,98
TRINITY_DN30547_c0_g1_i1	Nuclear protein 1	9,99
TRINITY_DN31547_c5_g3_i4	EH domain-containing protein 3	9,99
TRINITY_DN29339_c0_g1_i1	CD81 protein	9,99
TRINITY_DN28508_c0_g1_i2	Phosphoserine aminotransferase	10,00
TRINITY_DN32330_c0_g1_i14	TBC1 domain family member 2B	10,00
TRINITY_DN27012_c0_g3_i1	Roquin-1	10,01
TRINITY_DN32224_c5_g1_i11	Mitochondrial glycine transporter A	10,01
TRINITY_DN31057_c0_g1_i4	Endoribonuclease Dicer	10,02
TRINITY_DN29440_c0_g1_i4	Glutamine and serine-rich protein 1	10,02
TRINITY_DN25885_c3_g1_i6	Guanine nucleotide-binding protein $G(s)$ subunit alpha isoform:	10,03
TRINITY_DN30452_c0_g2_i8	DnaJ homolog subfamily A member 1	10,03
TRINITY_DN24984_c2_g3_i1	Serrate RNA effector molecule homolog	10,03
TRINITY_DN33106_c4_g2_i1	Leukemia inhibitory factor receptor	10,04
TRINITY_DN32978_c0_g1_i1	ValinetRNA ligase	10,04
TRINITY_DN33282_c5_g1_i9	Regulator of microtubule dynamics protein 3	10,04
TRINITY_DN29935_c0_g1_i6	Mannose-P-dolichol utilization defect 1 protein	10,06
TRINITY_DN32245_c0_g1_i1	Thioredoxin-interacting protein	10,06
TRINITY_DN30462_c0_g1_i2	Excitatory amino acid transporter 1	10,07
TRINITY_DN25394_c0_g1_i4	BOLA class I histocompatibility antigen, alpha chain BL3-7	10,08
TRINITY_DN32660_c0_g1_i21	Protein scribble homolog	10,08
TRINITY_DN32731_c0_g2_i6	Heterogeneous nuclear ribonucleoprotein C	10,08
TRINITY_DN30270_c3_g1_i5	Lysocardiolipin acyltransferase 1	10,08
TRINITY_DN27565_c0_g1_i2	COUP transcription factor 2	10,08
TRINITY_DN28233_c1_g1_i6	Glycogen synthase kinase-3 beta	10,08
TRINITY_DN31535_c2_g1_i8	Signal transducer and activator of transcription 5B	10,08
TRINITY_DN33238_c5_g1_i15	Telomerase protein component 1	10,08
	Pantothenate kinase 1	10,09
TRINITY_DN32558_c0_g1_i6	Protein Shroom1	
TRINITY_DN32372_c2_g1_i9	Protein KIAA0100	10,10
TRINITY_DN31471_c1_g2_i12		10,10
TRINITY_DN28964_c1_g1_i8	Exosome complex exonuclease RRP44	10,10
TRINITY_DN30452_c0_g2_i16	DnaJ homolog subfamily A member 1	10,10
TRINITY_DN31934_c0_g2_i10	Glycerophosphocholine phosphodiesterase GPCPD1	10,10
TRINITY_DN25644_c0_g1_i3	Prohibitin-2	10,11
TRINITY_DN29922_c2_g1_i6	6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase	10,11
TRINITY_DN31043_c0_g1_i3	Arsenite methyltransferase	10,12
TRINITY_DN32882_c1_g1_i10	Inositol 1,4,5-trisphosphate receptor type 1	10,12
TRINITY_DN28551_c1_g1_i5	Nuclear receptor coactivator 6	10,13
TRINITY_DN27398_c0_g1_i5	Pre-mRNA-splicing factor SLU7	10,13
TRINITY_DN31748_c0_g1_i9	Acetyl-CoA acetyltransferase, mitochondrial	10,14
TRINITY_DN25606_c0_g5_i2	Major histocompatibility complex class I-related gene protein	10,14
TRINITY_DN23763_c5_g3_i4	Exopolyphosphatase PRUNE1	10,14
TRINITY_DN28375_c1_g1_i3	Trafficking kinesin-binding protein 1	10,16
TRINITY_DN32749_c0_g1_i12	Protein TASOR	10,18
TRINITY_DN32452_c0_g1_i2	Ubiquitin carboxyl-terminal hydrolase 32	10,18
TRINITY_DN32660_c0_g1_i5	Protein scribble homolog	10,18
TRINITY_DN25520_c0_g1_i2	E3 ubiquitin-protein ligase SMURF2	10,19
TRINITY_DN31561_c2_g1_i3	Alanine aminotransferase 2-like	10,19
TRINITY_DN31395_c0_g1_i11	ATP-binding cassette sub-family A member 5	10,22
TRINITY_DN29412_c0_g1_i3	KAT8 regulatory NSL complex subunit 3	10,22
TRINITY_DN32200_c2_g1_i3	Forkhead box protein O4	10,23
TRINITY_DN32912_c2_g2_i1	Peroxisome proliferator-activated receptor alpha	10,24
TRINITY_DN32459_c2_g1_i9	Zinc finger FYVE domain-containing protein 1	10,24
TRINITY_DN26235_c0_g1_i40	5'-AMP-activated protein kinase subunit beta-2	10,26
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TRINITY DN22000 -0 -1 :10	History Institut November of CETD2	10.26
TRINITY_DN32080_c0_g1_i10	Histone-lysine N-methyltransferase SETD2	10,26
TRINITY_DN30280_c1_g1_i2	Bifunctional heparan sulfate N-deacetylase/N-sulfotransferase 2	10,27
TRINITY_DN28647_c4_g2_i4	Sestrin-1	10,27
TRINITY_DN28229_c2_g1_i11	Centromere protein J	10,27
TRINITY_DN26379_c1_g1_i6	Protein transport protein Sec61 subunit gamma	10,29
TRINITY_DN32978_c0_g1_i5	ValinetRNA ligase	10,29
TRINITY_DN32835_c0_g2_i2	Fibronectin type-III domain-containing protein 3a	10,29
TRINITY_DN31366_c0_g1_i5	Serine/threonine-protein phosphatase 4 catalytic subunit B	10,29
TRINITY_DN31786_c0_g2_i19	Ligand of Numb protein X 2	10,29
TRINITY_DN29751_c0_g1_i3	Autophagy-related protein 16-1	10,29
TRINITY_DN28754_c0_g2_i12	Aspartyl aminopeptidase	10,32
TRINITY_DN25391_c0_g1_i8	Glutamine synthetase	10,32
TRINITY_DN33279_c1_g1_i4	RNA-binding protein 4	10,32
TRINITY_DN29517_c0_g2_i24	Rho guanine nucleotide exchange factor 38	10,32
TRINITY_DN26106_c1_g2_i6	Transposon Tf2-6 polyprotein	10,33
TRINITY_DN33080_c3_g1_i1	Vitamin K-dependent gamma-carboxylase	10,34
TRINITY_DN26034_c1_g1_i6	Ubiquitin carboxyl-terminal hydrolase 7	10,34
TRINITY_DN32763_c1_g1_i20	SOSS complex subunit B2	10,35
TRINITY_DN27403_c0_g1_i13	Long-chain-fatty-acidCoA ligase ACSBG1	10,35
TRINITY_DN25039_c0_g1_i3	Elongation of very long chain fatty acids protein 4	10,36
TRINITY_DN32944_c0_g1_i3	Vacuolar protein sorting-associated protein 13C	10,36
_		10,36
TRINITY_DN30503_c1_g1_i3	Protein sel-1 homolog 1	
TRINITY_DN32502_c2_g2_i9	UDP-glucuronosyltransferase 2A1	10,37
TRINITY_DN28797_c0_g1_i15	Serine/threonine-protein kinase ppk4	10,38
TRINITY_DN31620_c0_g3_i1	Deoxynucleoside triphosphate triphosphohydrolase SAMHD1	10,39
TRINITY_DN28027_c1_g2_i6	Tectonin beta-propeller repeat-containing protein 1	10,39
TRINITY_DN24675_c2_g1_i9	Retinol-binding protein 2	10,41
TRINITY_DN33104_c6_g1_i10	Histidine ammonia-lyase	10,41
TRINITY_DN28185_c0_g1_i9	DnaJ homolog subfamily C member 1	10,44
TRINITY_DN32457_c1_g1_i8	Alpha-mannosidase 2	10,45
TRINITY_DN29252_c1_g1_i7	Coatomer subunit delta	10,46
TRINITY_DN30128_c1_g1_i1	Alpha-aminoadipic semialdehyde dehydrogenase	10,46
TRINITY_DN31058_c0_g1_i6	Ezrin	10,46
TRINITY_DN30450_c2_g1_i4	Neurogenic locus notch homolog protein 2	10,46
TRINITY_DN32757_c0_g2_i1	Baculoviral IAP repeat-containing protein 6	10,47
TRINITY_DN32228_c0_g1_i11	Translation initiation factor eIF-2B subunit epsilon	10,47
TRINITY_DN29010_c0_g1_i3	Eukaryotic translation initiation factor 5B	10,48
TRINITY_DN32406_c2_g1_i4	Transcription elongation factor SPT6	10,48
TRINITY_DN33194_c1_g1_i10	Very-long-chain (3R)-3-hydroxyacyl-CoA dehydratase 2	10,49
TRINITY_DN33088_c3_g1_i1	Inter-alpha-trypsin inhibitor heavy chain H3	10,49
TRINITY_DN32245_c0_g1_i3	Thioredoxin-interacting protein	10,49
TRINITY_DN29793_c1_g1_i3	Choline transporter-like protein 2	10,50
TRINITY_DN32906_c2_g1_i6	Apoptotic chromatin condensation inducer in the nucleus	10,50
TRINITY_DN32191_c3_g1_i1	Sulfite oxidase, mitochondrial	10,51
TRINITY_DN27341_c0_g1_i4	F-box only protein 33	10,51
TRINITY_DN32757_c0_g2_i3	Baculoviral IAP repeat-containing protein 6	10,52
TRINITY_DN29688_c1_g2_i9	Methylmalonyl-CoA mutase, mitochondrial	10,53
TRINITY_DN30276_c0_g1_i3	Tyrosine-protein phosphatase non-receptor type 1	10,53
TRINITY_DN31197_c0_g1_i1	Coatomer subunit beta'	10,54
TRINITY_DN29583_c0_g1_i9	Serine/threonine-protein kinase SIK3 homolog	10,55
TRINITY_DN32679_c1_g1_i1	Inactive histone-lysine N-methyltransferase 2E	10,56
TRINITY_DN32970_c1_g1_i4	Regulator of nonsense transcripts 2	10,58
TRINITY_DN32970_c1_g1_i5	Sodium-coupled neutral amino acid transporter 4	10,58
TRINITY_DN31065_c3_g1_i8	Cytoskeleton-associated protein 4	10,61
TRINITY_DN31027_c2_g2_i7	Erlin-2 Protein phosphatasa II	10,61
TRINITY_DN25744_c0_g1_i3	Protein phosphatase 1L	10,61
TRINITY_DN27201_c1_g3_i2	Cyclic AMP-dependent transcription factor ATF-5	10,62
TRINITY_DN26205_c2_g2_i3	Palmitoyltransferase ZDHHC18	10,63
TRINITY_DN28355_c3_g2_i2	Dual specificity tyrosine-phosphorylation-regulated kinase 1A	10,63

TRINITY DN3320 c. g. g. li 6	TED IN HERV DAY 2001 2 1 16	TH FO. 11	10.61
TRINITY_DN2986_c0_g2_i3			
TRINITY DN2993.6.0_gl_35			
TRINITY_DN2982_c3_g1_3		•	
RRINITY DN32536_cl_gl_identity Membrane-associated guanylate kinase, WW and PDZ domain- 10,66		<u> •</u>	
TRINITY DN28952_63_g1_39	_		
TRINITY DN3165_c3_g1_i9			
RRINITY_DN28375_el_gl_i2	_	Arylsulfatase B	10,67
TRINITY_DN2623R_c0_gl_i3 Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial 10,69 TRINITY_DN29229_c0_gl_i36 Histidine ammonia-lyase 10,70 10,70 10,71 10,71 10,72 10,71 10,72 10,72 10,72 10,73 10,72 10,73 10,72 10,73 10,72 10,73 10,73 10,72 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,73 10,74 10,74 10,75	TRINITY_DN31065_c3_g1_i9	Cytoskeleton-associated protein 4	10,68
RRINITY_DN2929_c0_g1_i6	TRINITY_DN28375_c1_g1_i2	Trafficking kinesin-binding protein 1	10,68
TRINITY_DN2929_c0_g2_i36	TRINITY_DN26378_c0_g1_i1	Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial	10,69
TRINITY_DN31889_cl_gl_i3	TRINITY_DN29379_c0_g1_i7	Splicing factor 3B subunit 1	10,70
TRINITY_DN32338_cl_gl_i7	TRINITY_DN29229_c0_g2_i36	Histidine ammonia-lyase	10,70
TRINITY_DN32338_cl_gl_i7	TRINITY_DN31589_c1_g1_i4	Coagulation factor V	10,71
TRINITY_DN32149_c0_g1_i5	TRINITY DN29079 c1 g3 i3	SWI/SNF complex subunit SMARCC1	10,72
TRINITY_DN32449_c0_g1_i5			
TRINITY_DN32457_cl_gl_il			
TRINITY_DN25913_c6_gl_i6	_		
TRINITY_DN35913_c6_g1_i6		*	
TRINITY_DN31459_cl_gl_i8			
TRINITY_DN32189_c0_g1_i3			
TRINITY_DN32189_c0_g1_i3			
TRINITY_DN2493_c2_gd_i2		· · ·	
TRINITY_DN24931_c2_g4_i12		* *	
TRINITY_DN26798_c0_g2_i10			
TRINITY_DN32587_cl_g2_i11			
TRINITY_DN25244_e0_gl_i1	_		
TRINITY_DN25394_c0_gl_i30			
TRINITY_DN30262_cl_gl_i9 MAU2 chromatid cohesion factor homolog 10,95 TRINITY_DN30262_cl_gl_i7 Protein kinase C-binding protein 1 10,97 TRINITY_DN3104_c6_gl_i7 Histidine ammonia-lyase 10,98 TRINITY_DN30404_c1_g2_i3 ATP-dependent 6-phosphofructokinase, liver type 11,00 TRINITY_DN29810_c0_gl_i8 Glutamine-tRNA ligase 11,02 TRINITY_DN31543_cl_gl_i1 Solute carrier family 2, facilitated glucose transporter member: 11,04 TRINITY_DN31655_c3_gl_i8 TBCI domain family member 8B 11,09 TRINITY_DN3065_c3_gl_i3 TBCI domain family member 8B 11,09 TRINITY_DN30762_c8_gl_i4 U2 snRNP-associated Drotein 4 11,09 TRINITY_DN32403_cl_gl_i10 Dedicator of cytokinesis protein 8 11,11 TRINITY_DN32403_cl_gl_i1 CTD small phosphatase-like protein 11,12 TRINITY_DN3290_cl_gl_i1 CTD small phosphatase-like protein 11,12 TRINITY_DN3290_cl_gl_i2 Heterogeneous nuclear ribonucleoprotein M 11,17 TRINITY_DN33318_c5_g2_i11 Critochrome P450 IAI 11,22 TRINITY_DN332205_cl_gl_i3 Cytochrome P450 IAI 11,22		* *	
TRINITY_DN33104_c6_g1_i7 Protein kinase C-binding protein 1 10,97 TRINITY_DN33104_c6_g1_i7 Histidine ammonia-lyase 10,98 TRINITY_DN36404_c1_g2_i3 ATP-dependent 6-phosphofructokinase, liver type 11,00 TRINITY_DN29810_c0_g1_i8 Keratinocyte-associated protein 2 11,00 TRINITY_DN31543_c1_g1_i1 Solute carrier family 2, facilitated glucose transporter member . 11,04 TRINITY_DN31065_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN31065_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN32062_c8_g1_i4 U2 snRNP-associated SURP motif-containing protein 11,10 TRINITY_DN32403_c1_g1_i10 Dedicator of cytokinesis protein 8 11,11 TRINITY_DN32647_c4_g2_j9 Sestrin-1 11,12 TRINITY_DN31290_c1_g1_i1 CTO small phosphatase-like protein 11,12 TRINITY_DN30365_c1_g1_i26 DNA transposase THAP9 11,13 TRINITY_DN32225_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN32225_c0_g1_i1 Cytochrome P450 IA1 11,22 TRINITY_DN3021_c1_g4_i2 Fibrillin-1 11,23 TRINITY_DN31920_c1_g1_i16 <td>_</td> <td></td> <td></td>	_		
TRINITY_DN30404_c6_g1_i7 Histidine ammonia-lyase 11,00 TRINITY_DN30404_c1_g2_i3 ATP-dependent 6-phosphofructokinase, liver type 11,00 TRINITY_DN26924_c3_g1_i2 Keratinocyte-associated protein 2 11,00 TRINITY_DN29810_c0_g1_i8 Glutamine-tRNA ligase 11,02 TRINITY_DN31543_c1_g1_i1 Solute carrier family 2, facilitated glucose transporter member: 11,04 TRINITY_DN29635_c0_g3_i8 TBC1 domain family member 8B 11,05 TRINITY_DN31065_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN30762_c8_g1_i4 U2 snRNP-associated SURP motif-containing protein 11,10 TRINITY_DN32403_c1_g1_i10 Dedicator of cytokinesis protein 8 11,11 TRINITY_DN32647_c4_g2_i9 Sestrin-1 11,12 TRINITY_DN31290_c1_g1_i1 CTD small phosphatase-like protein 11,12 TRINITY_DN30365_c1_g1_i26 DNA transposase THAP9 11,13 TRINITY_DN32707_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN323318_c5_g2_i15 Chloride transport protein 6 11,21 TRINITY_DN30423_c0_g1_i3 Cytochrome P450 lA1 11,22 TRINITY_DN30507_c1_g1_i5 TOM1-like protein 2 11,23 TRINITY_DN30201_c1_g4_i2 Fibrillin-1 TRINITY_DN30210_c1_g1_i1 Exitatory amino acid transporter 1 11,30 TRINITY_DN302120_c1_g1_i1 Exitatory amino acid transcription 5B 11,36 TRINITY_DN31275_c1_g1_i1 Cluconventional myosin-lXb 11,36 TRINITY_DN31540_c0_g1_i7 Glucosidase 2 subunit beta 11,36 TRINITY_DN3190_c1_g1_i122 Glucosidase 2 subunit beta 11,43 TRINITY_DN3190_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN3190_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN3190_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN3190_c1_g1_i3 CJuconventional myosin-lXb 11,49			
TRINITY_DN30404_cl_g2_i3	_		
TRINITY_DN26924_c3_g1_i2	_	· · · · · · · · · · · · · · · · · · ·	
TRINITY_DN31543_c1_g1_i1 Solute carrier family 2, facilitated glucose transporter member : 11,04 TRINITY_DN3165_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN31065_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN31065_c3_g1_i1 U2 snRNP-associated SURP motif-containing protein 11,10 TRINITY_DN32403_c1_g1_i10 Dedicator of cytokinesis protein 8 11,11 TRINITY_DN31290_c1_g1_i1 CTD small phosphatase-like protein 11,12 TRINITY_DN31290_c1_g1_i2 DNA transposase THAP9 11,13 TRINITY_DN30365_c1_g1_i26 DNA transposase THAP9 11,13 TRINITY_DN32797_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN32295_c0_g1_i1 Large neutral amino acids transporter small subunit 3 11,19 TRINITY_DN3225_c0_g1_i3 Cytochrome P450 IAI 11,22 TRINITY_DN30423_c0_g1_i3 Cytochrome P450 IAI 11,22 TRINITY_DN30423_c0_g1_i3 Epithelial splicing regulatory protein 2 11,23 TRINITY_DN30402_c0_g1_i10 Excitatory amino acid transporter I 11,24 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,44 TRINITY_DN31920_c1_g1_i12 E3 ubiquitin/ISG15 ligase TRIM25 11,44			
TRINITY_DN31543_c1_g1_i1 Solute carrier family 2, facilitated glucose transporter member : 11,04 TRINITY_DN29635_c0_g3_i8 TBC1 domain family member 8B 11,05 TRINITY_DN31065_c3_g1_i2 Cytoskeleton-associated protein 4 11,09 TRINITY_DN30762_c8_g1_i4 U2 snRNP-associated SURP motif-containing protein 11,10 TRINITY_DN32403_c1_g1_i10 Dedicator of cytokinesis protein 8 11,11 TRINITY_DN32647_c4_g2_i9 Sestrin-1 TRINITY_DN31290_c1_g1_i1 CTD small phosphatase-like protein 11,12 TRINITY_DN30365_c1_g1_i26 DNA transposase THAP9 11,13 TRINITY_DN32797_c0_g1_i1 Heterogeneous nuclear ribonucleoprotein M 11,17 TRINITY_DN32797_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN32295_c0_g1_i1 Large neutral amino acids transporter small subunit 3 11,19 TRINITY_DN3318_c5_g2_i15 Chloride transport protein 6 11,21 TRINITY_DN30423_c0_g1_i3 Cytochrome P450 IA1 TRINITY_DN30423_c0_g1_i3 TOM1-like protein 2 TRINITY_DN302507_c1_g1_i5 TOM1-like protein 2 TRINITY_DN3026277_c3_g1_i1 Epithelial splicing regulatory protein 2 11,23 TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i6 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31275_c1_g1_i12 Unconventional myosin-IXb TRINITY_DN31275_c1_g1_i12 Golgin subfamily A member 4 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31275_c1_g1_i12 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31275_c1_g1_i12 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31275_c1_g1_i12 E3 ubiquitin/ISG15 ligase TRIM25 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25	_	· · · · · · · · · · · · · · · · · · ·	
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TRINITY_DN31065_c3_g1_i2			
TRINITY_DN30762_c8_g1_i4		* *	
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TRINITY_DN27020_c0_g1_i2 Heterogeneous nuclear ribonucleoprotein M 11,17 TRINITY_DN32797_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN32225_c0_g1_i1 Large neutral amino acids transporter small subunit 3 11,19 TRINITY_DN33318_c5_g2_i15 Chloride transport protein 6 11,21 TRINITY_DN30423_c0_g1_i3 Cytochrome P450 IAI 11,22 TRINITY_DN32507_c1_g1_i5 TOM1-like protein 2 11,23 TRINITY_DN32507_c3_g1_i1 Epithelial splicing regulatory protein 2 11,24 TRINITY_DN33021_c1_g4_i2 Fibrillin-1 11,27 TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN31920_c1_g1_i8 Alpha-actinin-4 11,49			
TRINITY_DN32797_c0_g1_i1 Fatty acid synthase 11,18 TRINITY_DN32225_c0_g1_i1 Large neutral amino acids transporter small subunit 3 11,19 TRINITY_DN33318_c5_g2_i15 Chloride transport protein 6 11,21 TRINITY_DN30423_c0_g1_i3 Cytochrome P450 1A1 11,22 TRINITY_DN32507_c1_g1_i5 TOM1-like protein 2 11,23 TRINITY_DN32507_c3_g1_i1 Epithelial splicing regulatory protein 2 11,24 TRINITY_DN3021_c1_g4_i2 Fibrillin-1 11,27 TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN3195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN30365_c1_g1_i26	DNA transposase THAP9	11,13
TRINITY_DN32225_c0_g1_i1	TRINITY_DN27020_c0_g1_i2	Heterogeneous nuclear ribonucleoprotein M	11,17
TRINITY_DN33318_c5_g2_i15	TRINITY_DN32797_c0_g1_i1	Fatty acid synthase	11,18
TRINITY_DN30423_c0_g1_i3	TRINITY_DN32225_c0_g1_i1	Large neutral amino acids transporter small subunit 3	11,19
TRINITY_DN32507_c1_g1_i5	TRINITY_DN33318_c5_g2_i15	Chloride transport protein 6	11,21
TRINITY_DN26277_c3_g1_i1 Epithelial splicing regulatory protein 2 11,24 TRINITY_DN33021_c1_g4_i2 Fibrillin-1 11,27 TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN30423_c0_g1_i3	Cytochrome P450 1A1	11,22
TRINITY_DN33021_c1_g4_i2 Fibrillin-1 11,27 TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN32507_c1_g1_i5	TOM1-like protein 2	11,23
TRINITY_DN30462_c0_g1_i10 Excitatory amino acid transporter 1 11,30 TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN26277_c3_g1_i1	Epithelial splicing regulatory protein 2	11,24
TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN33021_c1_g4_i2	Fibrillin-1	11,27
TRINITY_DN31920_c1_g1_i16 E3 ubiquitin/ISG15 ligase TRIM25 11,34 TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	TRINITY_DN30462_c0_g1_i10	Excitatory amino acid transporter 1	11,30
TRINITY_DN31535_c2_g1_i5 Signal transducer and activator of transcription 5B 11,36 TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49			
TRINITY_DN32298_c3_g1_i12 Unconventional myosin-IXb 11,36 TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	_		
TRINITY_DN31275_c1_g1_i12 Glucosidase 2 subunit beta 11,36 TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49			
TRINITY_DN31549_c0_g1_i7 Golgin subfamily A member 4 11,43 TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49			
TRINITY_DN31920_c1_g1_i22 E3 ubiquitin/ISG15 ligase TRIM25 11,44 TRINITY_DN28195_c0_g1_i8 Alpha-actinin-4 11,49	_		
TRINITY_DN28195_c0_g1_i8			
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TRINITY_DN31579_c0_g2_i1	APOBEC1 complementation factor	11,53
TRINITY_DN29580_c0_g1_i4	Ubiquitin carboxyl-terminal hydrolase 5	11,53
TRINITY_DN25107_c0_g1_i1	T-complex protein 1 subunit delta	11,55
TRINITY_DN29333_c2_g1_i3	Calcium-binding protein 39-like	11,59
TRINITY_DN32475_c5_g2_i9	UTPglucose-1-phosphate uridylyltransferase	11,64
TRINITY_DN29800_c1_g4_i1	Aspartate aminotransferase, cytoplasmic	11,66
TRINITY_DN25515_c1_g1_i10	Monocarboxylate transporter 1	11,66
TRINITY_DN25992_c0_g1_i5	Protein SCAF11	11,76
TRINITY_DN33071_c2_g1_i6	UDP-N-acetylglucosaminepeptide N-acetylglucosaminyltransj	11,87
TRINITY_DN25394_c0_g1_i34	Class I histocompatibility antigen, F10 alpha chain	11,91
TRINITY_DN25473_c3_g1_i13	S-adenosylmethionine synthase isoform type-1	11,97
TRINITY_DN29810_c0_g1_i3	GlutaminetRNA ligase	11,98
TRINITY_DN30872_c1_g1_i11	Elongation factor 1-alpha	12,05
TRINITY_DN31561_c2_g1_i8	Alanine aminotransferase 2-like	12,06
TRINITY_DN29887_c3_g1_i4	ERO1-like protein beta	12,13
TRINITY_DN27774_c7_g1_i2	Androgen receptor	12,23
TRINITY_DN33320_c4_g1_i4	Protein PRRC2B	12,25
TRINITY_DN32734_c1_g2_i14	Myosin-9	12,30
TRINITY_DN32745_c0_g2_i9	Transferrin receptor protein 1	12,30
TRINITY_DN32676_c1_g2_i3	Phospholipid hydroperoxide glutathione peroxidase	12,32
TRINITY_DN31401_c0_g2_i4	C4b-binding protein alpha chain	12,34
TRINITY_DN32844_c3_g1_i8	5'-nucleotidase domain-containing protein 2	12,37
TRINITY_DN32467_c2_g1_i1	Vitronectin	12,47
TRINITY_DN28588_c6_g4_i3	Sialomucin core protein 24	12,51
TRINITY_DN26041_c2_g1_i6	Cold-inducible RNA-binding protein B	12,51
TRINITY_DN29887_c3_g1_i7	ERO1-like protein beta	12,59
TRINITY_DN31753_c0_g1_i3	Mid1-interacting protein 1A	12,71
TRINITY_DN32734_c1_g2_i8	Myosin-9	12,96
TRINITY_DN32828_c2_g1_i8	Glucagon receptor	13,17
TRINITY_DN27303_c0_g1_i11	60S ribosomal protein L10	13,23
TRINITY_DN33104_c6_g1_i35	Histidine ammonia-lyase	14,78