Microarray Data
Cells + Compound 8.3 / Cells + DMSO @ 3h

# id	ratio	minus (1/ratio)	
ceacam6	12.59	12.587	carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)(CEACAM6) mRNA
redd1	4.786	4.786	regulated in development and DNA damage response 1, mRNA.
il8	2.729	2.7287	interleukin 8 (IL8), mRNA.
cyp4f11	2.287	2.2874	cytochrome P450, family 4, subfamily F, polypeptide 11 (CYP4F11), mRNA.
scnn1a	2.106	2.1058	sodium channel, nonvoltage-gated 1 alpha (SCNN1A), mRNA.
bhlhb2	2.048	2.0478	basic helix-loop-helix domain containing, class B, 2 (BHLHB2), mRNA.
fos	2.036	2.0363	v-fos FBJ murine osteosarcoma viral oncogene homolog (FOS), mRNA.
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znf45	0.5	-2	zinc finger protein 45 (ZNF45), mRNA.
wdr58	0.5	-2	THO complex 6 homolog (Drosophila) (THOC6), mRNA.
dhrs3	0.5	-2	dehydrogenase/reductase (SDR family) member 3 (DHRS3), mRNA.
mgc14141	0.5	-2	transmembrane protein 141 (TMEM141), mRNA.
bbx	0.5	-2	bobby sox homolog (Drosophila) (BBX), mRNA.
zmynd8	0.5	-2.001	zinc finger, MYND-type containing 8 (ZMYND8), transcript variant 3, mRNA.
fer1l4	0.5	-2.001	fer-1-like 4 (C. elegans) (FER1L4) on chromosome 20.
znf618	0.5	-2.002	zinc finger protein 618 (ZNF618), mRNA.
morg1	0.5	-2.002	mitogen-activated protein kinase organizer 1 (MORG1), mRNA.
nup107	0.5	-2.002	nucleoporin 107kDa (NUP107), mRNA.
epb49	0.5	-2.002	erythrocyte membrane protein band 4.9 (dematin) (EPB49), mRNA.
prps1	0.5	-2.002	phosphoribosyl pyrophosphate synthetase 1 (PRPS1), mRNA.
chst13	0.5	-2.002	carbohydrate (chondroitin 4) sulfotransferase 13 (CHST13), mRNA.
rims3	0.499	-2.002	regulating synaptic membrane exocytosis 3 (RIMS3), mRNA.
mrpl52	0.499	-2.002	mitochondrial ribosomal protein L52 (MRPL52)nuclear gene encoding mitochondrial protein, trans. variant 6mRNA
phka2	0.499	-2.003	phosphorylase kinase, alpha 2 (liver) (PHKA2), mRNA.
rnmt	0.499	-2.003	RNA (guanine-7-) methyltransferase (RNMT), mRNA.
kiaa0859	0.499	-2.003	KIAA0859 (KIAA0859), transcript variant 1, mRNA.
rc74	0.499	-2.003	integrator complex subunit 9 (INTS9), mRNA.
slc25a5	0.499	-2.003	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5 (SLC25A5), mRNA.
gprasp2	0.499	-2.003	G protein-coupled receptor associated sorting protein 2 (GPRASP2), transcript variant 2, mRNA.
mxd3	0.499	-2.004	MAX dimerization protein 3 (MXD3), mRNA.
cradd	0.499	-2.004	CASP2 and RIPK1 domain containing adaptor with death domain (CRADD), mRNA.
klhl21	0.499	-2.004	kelch-like 21 (Drosophila) (KLHL21), mRNA.
rnf170	0.499	-2.004	ring finger protein 170 (RNF170), mRNA.
atic	0.499	-2.004	5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase (ATIC), mRNA.
camk1	0.499	-2.005	calcium/calmodulin-dependent protein kinase I (CAMK1), mRNA.
vgf	0.499	-2.005	VGF nerve growth factor inducible (VGF), mRNA.
txnrd1	0.499	-2.005	thioredoxin reductase 1 (TXNRD1), transcript variant 4, mRNA.
mical1	0.499	-2.005	microtubule associated monoxygenase, calponin and LIM domain containing 1 (MICAL1), mRNA.
golga4	0.499	-2.005	golgi autoantigen, golgin subfamily a, 4 (GOLGA4), mRNA.

mgrn1	0.499	-2.005	mahogunin, ring finger 1 (MGRN1), mRNA.
loc644928	0.499	-2.006	PREDICTED: similar to 40S ribosomal protein S26 (LOC644928), mRNA.
commd5	0.499	-2.006	COMM domain containing 5 (COMMD5), mRNA.
scdr10	0.499	-2.006	hydroxysteroid (11-beta) dehydrogenase 1-like (HSD11B1L), transcript variant e, mRNA.
flj14054	0.499	-2.006	hypothetical protein FLJ14054 (FLJ14054), mRNA.
c9orf48	0.498	-2.006	chromosome 9 open reading frame 48 (C9orf48), mRNA.
znrf3	0.498	-2.006	PREDICTED: zinc and ring finger 3, transcript variant 5 (ZNRF3), mRNA.
c15orf24	0.498	-2.007	chromosome 15 open reading frame 24 (C15orf24), mRNA.
adam15	0.498	-2.007	PREDICTED: ADAM metallopeptidase domain 15 (metargidin) (ADAM15), mRNA.
znf142	0.498	-2.007	zinc finger protein 142 (ZNF142), mRNA.
smyd5	0.498	-2.007	SMYD family member 5 (SMYD5), mRNA.
fbxw4	0.498	-2.008	F-box and WD repeat domain containing 4 (FBXW4), mRNA.
golt1a	0.498	-2.008	golgi transport 1 homolog A (S. cerevisiae) (GOLT1A), mRNA.
zcchc3	0.498	-2.008	zinc finger, CCHC domain containing 3 (ZCCHC3), mRNA.
cln3	0.498	-2.008	ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease) (CLN3), mRNA.
tubgcp2	0.498	-2.008	tubulin, gamma complex associated protein 2 (TUBGCP2), mRNA.
golga5	0.498	-2.008	golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA.
st3gal5	0.498	-2.009	ST3 beta-galactoside alpha-2,3-sialyltransferase 5 (ST3GAL5), mRNA.
nacap1	0.498	-2.009	nascent-polypeptide-associated complex alpha polypeptide pseudogene 1 (NACAP1) on chromosome 8.
znf585a	0.498	-2.009	zinc finger protein 585A (ZNF585A), transcript variant 2, mRNA.
thumpd3	0.498	-2.009	THUMP domain containing 3 (THUMPD3), mRNA.
man1b1	0.498	-2.009	mannosidase, alpha, class 1B, member 1 (MAN1B1), mRNA.
dhcr24	0.498	-2.009	24-dehydrocholesterol reductase (DHCR24), mRNA.
slc7a7	0.498	-2.009	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7 (SLC7A7), mRNA.
ndufb7	0.498	-2.01	NADHdehydrogenase(ubiquinone)1bsubcomplex,7,18kDa(NDUFB7),nucl. gene encoding mitochon. protein,mRNA
pacs2	0.498	-2.01	phosphofurin acidic cluster sorting protein 2 (PACS2), mRNA.
ccne1	0.498	-2.01	cyclin E1 (CCNE1), transcript variant 1, mRNA.
sema3b	0.498	-2.01	sema domain, (Ig) domain, short basic domain, secreted, (semaphorin) 3B (SEMA3B), trans.variant 2, mRNA.
cklf	0.498	-2.01	chemokine-like factor (CKLF), transcript variant 6, mRNA.
pkd1	0.497	-2.01	polycystic kidney disease 1 (autosomal dominant) (PKD1), transcript variant 1, mRNA.
lsm3	0.497	-2.01	LSM3 homolog, U6 small nuclear RNA associated (S. cerevisiae) (LSM3), mRNA.
loc642209	0.497	-2.012	PREDICTED: similar to ribosomal protein L13a, transcript variant 1 (LOC642209), mRNA.
nkrf	0.497	-2.012	NF-kappaB repressing factor (NKRF), mRNA.
atp1a1	0.497	-2.012	ATPase, Na+/K+ transporting, alpha 1 polypeptide (ATP1A1), transcript variant 1, mRNA.
nsfl1c	0.497	-2.012	NSFL1 (p97) cofactor (p47) (NSFL1C), transcript variant 3, mRNA.
hs.561357	0.497	-2.012	UI-H-ED0-axn-h-17-0-UI.s1 NCI_CGAP_ED0 cDNA clone UI-H-ED0-axn-h-17-0-UI 3, mRNA sequence
loc116143	0.497	-2.012	hypothetical protein BC014022 (LOC116143), mRNA.
wdfy1	0.497	-2.012	WD repeat and FYVE domain containing 1 (WDFY1), mRNA.
loc388642	0.497	-2.012	PREDIC:similar to Triosephosphate isomerase(TIM)(Triose-phosphate isomerase),trans. variant 3(LOC388642)mRN
loc652826	0.497	-2.012	PREDIC:similar to 26S protease regulatory subunit 6B(MIP224)(MB67-interacting protein)(LOC652826)mRNA.
tmem11	0.497	-2.013	transmembrane protein 11 (TMEM11), mRNA.
luzp1	0.497	-2.013	leucine zipper protein 1 (LUZP1), mRNA.

tbc1d2b	0.497	-2.014	TBC1 domain family, member 2B (TBC1D2B), mRNA.
dgcr14	0.497	-2.014	DiGeorge syndrome critical region gene 14 (DGCR14), mRNA.
rsu1	0.497	-2.014	Ras suppressor protein 1 (RSU1), transcript variant 2, mRNA.
vrk1	0.497	-2.014	vaccinia related kinase 1 (VRK1), mRNA.
mrpl42	0.497	-2.014	mitochondrial ribosomal protein L42 (MRPL42), nuclear gene encoding mitochon. protein, trans.variant 2, mRNA.
c14orf156	0.497	-2.014	chromosome 14 open reading frame 156 (C14orf156), mRNA.
ube3a	0.496	-2.015	ubiquitin protein ligaseE3A(human papilloma virus E6-assoc.protein,Angelman synd.)(UBE3A)trans.variant3mRNA
mgc52110	0.496	-2.015	hypothetical protein MGC52110 (MGC52110), mRNA.
hs.572219	0.496	-2.015	CM3-MT0357-260101-690-b10 MT0357 cDNA, mRNA sequence
loc645489	0.496	-2.015	PREDICTED: hypothetical protein LOC645489 (LOC645489), mRNA.
bub1b	0.496	-2.015	BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast) (BUB1B), mRNA.
slc25a15	0.496	-2.015	solute carrier family 25(mitocho.carrier; ornithine transporter)member 15,nucl.gene encode mitocho.protein,mRN
tsen2	0.496	-2.015	tRNA splicing endonuclease 2 homolog (S. cerevisiae) (TSEN2), mRNA.
anp32b	0.496	-2.015	acidic (leucine-rich) nuclear phosphoprotein 32 family, member B (ANP32B), mRNA.
nbl1	0.496	-2.016	neuroblastoma, suppression of tumorigenicity 1 (NBL1), transcript variant 2, mRNA.
ppp6c	0.496	-2.016	protein phosphatase 6, catalytic subunit (PPP6C), mRNA.
ddx39	0.496	-2.016	DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 (DDX39), transcript variant 2, mRNA.
mrps11	0.496	-2.016	mitochondrial ribosomal protein S11(MRPS11)nuclear gene encoding mitochondrial protein, trans.variant 2,mRNA
ercc1	0.496	-2.017	excision repair cross-complementing rodent repair deficiency, complementation group 1 (includes overlapping antisense sequence) (ERCC1), transcript variant 1, mRNA.
c9orf116	0.496	-2.017	chromosome 9 open reading frame 116 (C9orf116), mRNA.
slc25a1	0.496	-2.017	solute carrier family 25 (mitochondrial carrier; citrate transporter), member 1 (SLC25A1), mRNA.
gpr125	0.496	-2.017	PREDICTED: G protein-coupled receptor 125, transcript variant 5 (GPR125), mRNA.
cdca2	0.496	-2.017	cell division cycle associated 2 (CDCA2), mRNA.
thrap3	0.496	-2.017	thyroid hormone receptor associated protein 3 (THRAP3), mRNA.
ipo4	0.496	-2.017	importin 4 (IPO4), mRNA.
loc653726	0.496	-2.018	PREDICTED: similar to family with sequence similarity 86, member B1 (LOC653726), mRNA.
loc388532	0.496	-2.018	PREDICTED: similar to ribosomal protein L21 isoform 1 (LOC388532), mRNA.
ttf1	0.496	-2.018	transcription termination factor, RNA polymerase I (TTF1), mRNA.
scrib	0.496	-2.018	scribbled homolog (Drosophila) (SCRIB), transcript variant 1, mRNA.
mgc2654	0.496	-2.018	LP8272 (MGC2654), mRNA.
zswim1	0.496	-2.018	zinc finger, SWIM-type containing 1 (ZSWIM1), mRNA.
cpne3	0.496	-2.018	copine III (CPNE3), mRNA.
orc1l	0.495	-2.018	origin recognition complex, subunit 1-like (yeast) (ORC1L), mRNA.
7-Mar	0.495	-2.018	membrane-associated ring finger (C3HC4) 7 (MARCH7), mRNA.
flj12529	0.495	-2.019	pre-mRNA cleavage factor I, 59 kDa subunit (FLJ12529), mRNA.
mrpl21	0.495	-2.019	mitochondrial ribosomal protein L21(MRPL21),nuclear gene encoding mitochondrial protein, trans.variant 3,mRNA
mad2l2	0.495	-2.019	MAD2 mitotic arrest deficient-like 2 (yeast) (MAD2L2), mRNA.
pttg1ip	0.495	-2.019	pituitary tumor-transforming 1 interacting protein (PTTG1IP), mRNA.
c9orf125	0.495	-2.019	chromosome 9 open reading frame 125 (C9orf125), mRNA.
gpr64	0.495	-2.019	G protein-coupled receptor 64 (GPR64), mRNA.
smpdl3a	0.495	-2.019	sphingomyelin phosphodiesterase, acid-like 3A (SMPDL3A), mRNA.

c8orf53	0.495	-2.019	chromosome 8 open reading frame 53 (C8orf53), mRNA.
tmem32	0.495	-2.019	transmembrane protein 32 (TMEM32), mRNA.
pafah1b3	0.495	-2.019	platelet-activating factor acetylhydrolase, isoform Ib, gamma subunit 29kDa (PAFAH1B3), mRNA.
znf146	0.495	-2.02	zinc finger protein 146 (ZNF146), mRNA.
tapbp	0.495	-2.02	TAP binding protein (tapasin) (TAPBP), transcript variant 3, mRNA.
cgi-115	0.495	-2.02	CGI-115 protein (CGI-115), mRNA.
loc647099	0.495	-2.02	PREDICTED: similar to 60S ribosomal protein L23a (LOC647099), mRNA.
ankrd25	0.495	-2.02	ankyrin repeat domain 25 (ANKRD25), mRNA.
dkc1	0.495	-2.02	dyskeratosis congenita 1, dyskerin (DKC1), mRNA.
cope	0.495	-2.02	coatomer protein complex, subunit epsilon (COPE), transcript variant 3, mRNA.
snrpb2	0.495	-2.02	small nuclear ribonucleoprotein polypeptide B'' (SNRPB2), transcript variant 1, mRNA.
chchd4	0.495	-2.02	coiled-coil-helix-coiled-coil-helix domain containing 4 (CHCHD4), mRNA.
hyou1	0.495	-2.021	hypoxia up-regulated 1 (HYOU1), mRNA.
abcb7	0.495	-2.021	ATP-binding cassette, sub-family B(MDR/TAP),member7(ABCB7),nuclr. gene encoding mitochon. protein,mRNA
mrpl49	0.495	-2.021	mitochondrial ribosomal protein L49 (MRPL49), nuclear gene encoding mitochondrial protein, mRNA.
hs.370359	0.495	-2.021	mRNA; cDNA DKFZp686F09166 (from clone DKFZp686F09166)
arnt2	0.495	-2.021	aryl-hydrocarbon receptor nuclear translocator 2 (ARNT2), mRNA.
hs.126768	0.495	-2.022	full-length cDNA clone CS0DI004YB08 of Placenta Cot 25-normalized of (human)
c16orf24	0.495	-2.022	chromosome 16 open reading frame 24 (C16orf24), mRNA.
extl3	0.495	-2.022	exostoses (multiple)-like 3 (EXTL3), mRNA.
c18orf8	0.495	-2.022	chromosome 18 open reading frame 8 (C18orf8), mRNA.
phf5a	0.494	-2.023	PHD finger protein 5A (PHF5A), mRNA.
papss1	0.494	-2.023	3'-phosphoadenosine 5'-phosphosulfate synthase 1 (PAPSS1), mRNA.
ptpn13	0.494	-2.023	protein tyrosine phosphatase,non-receptor type 13 (APO-1/CD95(Fas)-associated phosphatase) (PTPN13), transcript variant 3, mRNA.
pcmt1	0.494	-2.023	protein-L-isoaspartate (D-aspartate) O-methyltransferase (PCMT1), mRNA.
dirc2	0.494	-2.023	disrupted in renal carcinoma 2 (DIRC2), mRNA.
tagln2	0.494	-2.023	transgelin 2 (TAGLN2), mRNA.
myh9	0.494	-2.023	myosin, heavy polypeptide 9, non-muscle (MYH9), mRNA.
hexb	0.494	-2.023	hexosaminidase B (beta polypeptide) (HEXB), mRNA.
ppat	0.494	-2.024	phosphoribosyl pyrophosphate amidotransferase (PPAT), mRNA.
cryz	0.494	-2.024	crystallin, zeta (quinone reductase) (CRYZ), mRNA.
mgc13170	0.494	-2.024	chromosome 19 open reading frame 48 (C19orf48), mRNA.
nov	0.494	-2.024	nephroblastoma overexpressed gene (NOV), mRNA.
exosc5	0.494	-2.024	exosome component 5 (EXOSC5), mRNA.
raly	0.494	-2.024	RNA binding protein, autoantigenic (hnRNP-associated with lethal yellow homolog (mouse)) (RALY), transcript variant 1, mRNA.
rab4a	0.494	-2.024	RAB4A, member RAS oncogene family (RAB4A), mRNA.
pif1	0.494	-2.025	PIF1 5'-to-3' DNA helicase homolog (S. cerevisiae) (PIF1), mRNA.
loc285989	0.494	-2.025	zinc finger protein 789 (ZNF789), transcript variant 2, mRNA.
znf446	0.494	-2.025	zinc finger protein 446 (ZNF446), mRNA.
myo1b	0.494	-2.025	myosin IB (MYO1B), mRNA.

mrpl36	0.494	-2.025	mitochondrial ribosomal protein L36 (MRPL36), nuclear gene encoding mitochondrial protein, mRNA.
cnot2	0.494	-2.025	CCR4-NOT transcription complex, subunit 2 (CNOT2), mRNA.
lepre1	0.494	-2.026	leucine proline-enriched proteoglycan (leprecan) 1 (LEPRE1), mRNA.
hs.568777	0.494	-2.026	AGENCOURT_8210259 NIH_MGC_112 cDNA clone IMAGE:6258046 5, mRNA sequence
timeless	0.494	-2.026	timeless homolog (Drosophila) (TIMELESS), mRNA.
sumo1p3	0.494	-2.026	SUMO1 pseudogene 3 (SUMO1P3) on chromosome 1.
skiv2l	0.493	-2.027	superkiller viralicidic activity 2-like (S. cerevisiae) (SKIV2L), mRNA.
loc285193	0.493	-2.027	dual specificity phosphatase 28 (DUSP28), mRNA.
c9orf111	0.493	-2.027	patatin-like phospholipase domain containing 7 (PNPLA7), mRNA.
tfip11	0.493	-2.027	tuftelin interacting protein 11 (TFIP11), transcript variant 1, mRNA.
flj12681	0.493	-2.027	hypothetical protein FLJ12681 (FLJ12681), mRNA.
loc643367	0.493	-2.028	PREDICTED: region containing SMA4; hypothetical protein LOC153561, transcript variant 2 (LOC643367), mRNA.
dbndd1	0.493	-2.028	dysbindin (dystrobrevin binding protein 1) domain containing 1 (DBNDD1), transcript variant 1, mRNA.
ccm2	0.493	-2.028	cerebral cavernous malformation 2 (CCM2), transcript variant 2, mRNA.
psmb4	0.493	-2.028	proteasome (prosome, macropain) subunit, beta type, 4 (PSMB4), mRNA.
c20orf24	0.493	-2.028	chromosome 20 open reading frame 24 (C20orf24), transcript variant 1, mRNA.
c5orf37	0.493	-2.029	chromosome 5 open reading frame 37 (C5orf37), mRNA.
fam113b	0.493	-2.029	family with sequence similarity 113, member B (FAM113B), mRNA.
cbx4	0.493	-2.029	chromobox homolog 4 (Pc class homolog, Drosophila) (CBX4), mRNA.
plrg1	0.493	-2.029	pleiotropic regulator 1 (PRL1 homolog, Arabidopsis) (PLRG1), mRNA.
bcs1l	0.493	-2.029	BCS1-like (yeast) (BCS1L), mRNA.
zc3hc1	0.493	-2.029	zinc finger, C3HC-type containing 1 (ZC3HC1), mRNA.
dph5	0.493	-2.03	DPH5 homolog (S. cerevisiae) (DPH5), transcript variant 3, mRNA.
kifc1	0.493	-2.03	kinesin family member C1 (KIFC1), mRNA.
wdr71	0.493	-2.03	WD repeat domain 71 (WDR71), mRNA.
zak	0.492	-2.031	sterile alpha motif and leucine zipper containing kinase AZK (ZAK), transcript variant 1, mRNA.
ptpn12	0.492	-2.031	protein tyrosine phosphatase, non-receptor type 12 (PTPN12), mRNA.
oraov1	0.492	-2.031	oral cancer overexpressed 1 (ORAOV1), mRNA.
mgc21830	0.492	-2.031	chromosome 16 open reading frame 79 (C16orf79), mRNA.
tmem101	0.492	-2.032	transmembrane protein 101 (TMEM101), mRNA.
clcn7	0.492	-2.032	chloride channel 7 (CLCN7), mRNA.
mrps24	0.492	-2.032	mitochondrial ribosomal protein S24 (MRPS24), nuclear gene encoding mitochondrial protein, mRNA.
pphln1	0.492	-2.032	periphilin 1 (PPHLN1), transcript variant 4, mRNA.
cutc	0.492	-2.033	cutC copper transporter homolog (E. coli) (CUTC), mRNA.
hspa1a	0.492	-2.033	heat shock 70kDa protein 1A (HSPA1A), mRNA.
btbd7	0.492	-2.033	BTB (POZ) domain containing 7 (BTBD7), transcript variant 2, mRNA.
myo10	0.492	-2.033	myosin X (MYO10), mRNA.
commd1	0.492	-2.034	copper metabolism (Murr1) domain containing 1 (COMMD1), mRNA.
mthfs	0.492	-2.034	5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase) (MTHFS), mRNA.
xrcc1	0.492	-2.034	X-ray repair complementing defective repair in Chinese hamster cells 1 (XRCC1), mRNA.
pfkp	0.492	-2.034	phosphofructokinase, platelet (PFKP), mRNA.
opa3	0.492	-2.034	optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia) (OPA3), transcript variant 2, mRNA.

2'-pde	0.492	-2.035	2'-phosphodiesterase (2'-PDE), mRNA.
ctps	0.491	-2.035	CTP synthase (CTPS), mRNA.
dhx29	0.491	-2.035	DEAH (Asp-Glu-Ala-His) box polypeptide 29 (DHX29), mRNA.
kdelc1	0.491	-2.035	KDEL (Lys-Asp-Glu-Leu) containing 1 (KDELC1), mRNA.
sergef	0.491	-2.035	secretion regulating guanine nucleotide exchange factor (SERGEF), mRNA.
xpo1	0.491	-2.036	exportin 1 (CRM1 homolog, yeast) (XPO1), mRNA.
plekha6	0.491	-2.036	pleckstrin homology domain containing, family A member 6 (PLEKHA6), mRNA.
gpr108	0.491	-2.036	G protein-coupled receptor 108 (GPR108), transcript variant 1, mRNA.
loc653188	0.491	-2.036	PREDICTED: similar to Beta-glucuronidase precursor, transcript variant 10 (LOC653188), mRNA.
banf1	0.491	-2.036	barrier to autointegration factor 1 (BANF1), mRNA.
oip5	0.491	-2.036	Opa interacting protein 5 (OIP5), mRNA.
sgpl1	0.491	-2.036	sphingosine-1-phosphate lyase 1 (SGPL1), mRNA.
c8orf33	0.491	-2.036	chromosome 8 open reading frame 33 (C8orf33), mRNA.
bbs4	0.491	-2.036	Bardet-Biedl syndrome 4 (BBS4), mRNA.
pbk	0.491	-2.036	PDZ binding kinase (PBK), mRNA.
scml1	0.491	-2.037	sex comb on midleg-like 1 (Drosophila) (SCML1), transcript variant 1, mRNA.
snrpd1	0.491	-2.037	small nuclear ribonucleoprotein D1 polypeptide 16kDa (SNRPD1), mRNA.
tbc1d2	0.491	-2.037	TBC1 domain family, member 2 (TBC1D2), mRNA.
ikbkb	0.491	-2.037	inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta (IKBKB), mRNA.
c8orf41	0.491	-2.037	chromosome 8 open reading frame 41 (C8orf41), mRNA.
mta1	0.491	-2.037	metastasis associated 1 (MTA1), mRNA.
cnn3	0.491	-2.037	calponin 3, acidic (CNN3), mRNA.
ror1	0.491	-2.038	receptor tyrosine kinase-like orphan receptor 1 (ROR1), mRNA.
c9orf9	0.491	-2.038	chromosome 9 open reading frame 9 (C9orf9), mRNA.
myo9b	0.491	-2.038	myosin IXB (MYO9B), mRNA.
znf11b	0.491	-2.038	zinc finger protein 33B (ZNF33B), mRNA.
smad4	0.491	-2.038	SMAD family member 4 (SMAD4), mRNA.
acsl3	0.491	-2.038	acyl-CoA synthetase long-chain family member 3 (ACSL3), transcript variant 1, mRNA.
qil1	0.491	-2.039	hypothetical protein P117 (P117), mRNA.
c13orf1	0.491	-2.039	chromosome 13 open reading frame 1 (C13orf1), mRNA.
csnk2a1	0.49	-2.039	casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript variant 1, mRNA.
agpat7	0.49	-2.039	1-acylglycerol-3-phosphate O-acyltransferase 7 (lysophosphatidic acid acyltransferase, eta) (AGPAT7), mRNA.
gpr115	0.49	-2.039	G protein-coupled receptor 115 (GPR115), mRNA.
polr2f	0.49	-2.039	polymerase (RNA) II (DNA directed) polypeptide F (POLR2F), mRNA.
tra2a	0.49	-2.039	transformer-2 alpha (TRA2A), mRNA.
hs.473191	0.49	-2.039	cDNA FLJ34428 fis, clone HLUNG2000761
cxorf40a	0.49	-2.039	chromosome X open reading frame 40A (CXorf40A), mRNA.
usp5	0.49	-2.039	ubiquitin specific peptidase 5 (isopeptidase T) (USP5), mRNA.
znf410	0.49	-2.04	zinc finger protein 410 (ZNF410), mRNA.
prkdc	0.49	-2.04	protein kinase, DNA-activated, catalytic polypeptide (PRKDC), transcript variant 1, mRNA.
stk19	0.49	-2.04	serine/threonine kinase 19 (STK19), transcript variant 2, mRNA.
akt1s1	0.49	-2.04	AKT1 substrate 1 (proline-rich) (AKT1S1), mRNA.

rps19bp1	0.49	-2.041	ribosomal protein S19 binding protein 1 (RPS19BP1), mRNA.
wdr50	0.49	-2.041	WD repeat domain 50 (WDR50), mRNA.
lyar	0.49	-2.041	Ly1 antibody reactive homolog (mouse) (LYAR), mRNA.
mgc2574	0.49	-2.041	coiled-coil domain containing 86 (CCDC86), mRNA.
fam98c	0.49	-2.041	family with sequence similarity 98, member C (FAM98C), mRNA.
sdccag3	0.49	-2.041	serologically defined colon cancer antigen 3 (SDCCAG3), mRNA.
c11orf24	0.49	-2.041	chromosome 11 open reading frame 24 (C11orf24), mRNA.
c7orf26	0.49	-2.041	chromosome 7 open reading frame 26 (C7orf26), mRNA.
hddc3	0.49	-2.042	HD domain containing 3 (HDDC3), mRNA.
loc649169	0.49	-2.042	PREDICTED: similar to WD-repeat protein 74 (NOP seven-associated protein 1), transcript variant 1 (LOC649169), mRNA.
bcl9	0.49	-2.043	B-cell CLL/lymphoma 9 (BCL9), mRNA.
pex19	0.49	-2.043	peroxisomal biogenesis factor 19 (PEX19), mRNA.
wnt3	0.489	-2.043	wingless-type MMTV integration site family, member 3 (WNT3), mRNA.
loc51035	0.489	-2.043	SAPK substrate protein 1 (LOC51035), mRNA.
nt5m	0.489	-2.043	5',3'-nucleotidase, mitochondrial (NT5M), nuclear gene encoding mitochondrial protein, mRNA.
actr2	0.489	-2.043	ARP2 actin-related protein 2 homolog (yeast) (ACTR2), transcript variant 1, mRNA.
nek11	0.489	-2.043	NIMA (never in mitosis gene a)- related kinase 11 (NEK11), transcript variant 2, mRNA.
tns4	0.489	-2.043	tensin 4 (TNS4), mRNA.
cdt1	0.489	-2.044	chromatin licensing and DNA replication factor 1 (CDT1), mRNA.
ta-pp2c	0.489	-2.044	PTC7 protein phosphatase homolog (S. cerevisiae) (PPTC7), mRNA.
c14orf32	0.489	-2.044	chromosome 14 open reading frame 32 (C14orf32), mRNA.
sphk1	0.489	-2.044	sphingosine kinase 1 (SPHK1), transcript variant 2, mRNA.
c20orf98	0.489	-2.044	neurensin 2 (NRSN2), mRNA.
rnut1	0.489	-2.044	RNA, U transporter 1 (RNUT1), mRNA.
cinp	0.489	-2.044	cyclin-dependent kinase 2-interacting protein (CINP), mRNA.
angel1	0.489	-2.045	angel homolog 1 (Drosophila) (ANGEL1), mRNA.
znf444	0.489	-2.045	zinc finger protein 444 (ZNF444), mRNA.
rabep1	0.489	-2.045	rabaptin, RAB GTPase binding effector protein 1 (RABEP1), mRNA.
hs.20255	0.489	-2.045	AV652851 GLC cDNA clone GLCDEG06 3, mRNA sequence
col1a1	0.489	-2.045	collagen, type I, alpha 1 (COL1A1), mRNA.
seh1l	0.489	-2.045	SEH1-like (S. cerevisiae) (SEH1L), transcript variant 1, mRNA.
blm	0.489	-2.046	Bloom syndrome (BLM), mRNA.
gspt1	0.489	-2.046	G1 to S phase transition 1 (GSPT1), mRNA.
jam3	0.489	-2.046	junctional adhesion molecule 3 (JAM3), mRNA.
znf694	0.489	-2.046	zinc finger with KRAB and SCAN domains 2 (ZKSCAN2), mRNA.
st6galnac6	0.489	-2.046	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1, 3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 6 (ST6GALNAC6), mRNA.
c1orf121	0.489	-2.047	chromosome 1 open reading frame 121 (C1orf121), mRNA.
nr3c1	0.489	-2.047	nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor) (NR3C1), transcript variant 4, mRNA.
hs.91389	0.489	-2.047	cDNA clone IMAGE:3079901
eme1	0.489	-2.047	essential meiotic endonuclease 1 homolog 1 (S. pombe) (EME1), mRNA.

zyg11bl	0.488	-2.047	zer-1 homolog (C. elegans) (ZER1), mRNA.
tbc1d14	0.488	-2.048	TBC1 domain family, member 14 (TBC1D14), mRNA.
asphd1	0.488	-2.048	aspartate beta-hydroxylase domain containing 1 (ASPHD1), mRNA.
rhobtb2	0.488	-2.049	Rho-related BTB domain containing 2 (RHOBTB2), mRNA.
bag5	0.488	-2.049	BCL2-associated athanogene 5 (BAG5), transcript variant 3, mRNA.
sez6l2	0.488	-2.049	seizure related 6 homolog (mouse)-like 2 (SEZ6L2), transcript variant 2, mRNA.
sbds	0.488	-2.049	Shwachman-Bodian-Diamond syndrome (SBDS), mRNA.
pde6d	0.488	-2.049	phosphodiesterase 6D, cGMP-specific, rod, delta (PDE6D), mRNA.
ibtk	0.488	-2.049	inhibitor of Bruton agammaglobulinemia tyrosine kinase (IBTK), mRNA.
abcb10	0.488	-2.05	ATP-binding cassette, sub-family B (MDR/TAP), member 10 (ABCB10), nuclear gene encoding mitochondrial protein, mRNA.
loc646900	0.488	-2.05	PREDICTED: similar to zinc finger and BTB domain containing 8 opposite strand (LOC646900), mRNA.
loc652162	0.488	-2.05	PREDICTED: similar to 60S ribosomal protein L32 (LOC652162), mRNA.
slc4a1ap	0.488	-2.05	solute carrier family 4 (anion exchanger), member 1, adaptor protein (SLC4A1AP), mRNA.
rkhd3	0.488	-2.05	mex-3 homolog B (C. elegans) (MEX3B), mRNA.
cnap1	0.488	-2.05	non-SMC condensin I complex, subunit D2 (NCAPD2), mRNA.
kpna6	0.488	-2.05	karyopherin alpha 6 (importin alpha 7) (KPNA6), mRNA.
lrdd	0.488	-2.05	leucine-rich repeats and death domain containing (LRDD), transcript variant 2, mRNA.
tinp1	0.488	-2.051	TGF beta-inducible nuclear protein 1 (TINP1), mRNA.
trim56	0.488	-2.051	tripartite motif-containing 56 (TRIM56), mRNA.
gpaa1	0.488	-2.051	GPAA1P anchor attachment protein 1 homolog (yeast) (GPAA1), mRNA.
prep	0.488	-2.051	prolyl endopeptidase (PREP), mRNA.
atg12	0.488	-2.051	ATG12 autophagy related 12 homolog (S. cerevisiae) (ATG12), mRNA.
znf195	0.487	-2.051	zinc finger protein 195 (ZNF195), mRNA.
uhmk1	0.487	-2.052	U2AF homology motif (UHM) kinase 1 (UHMK1), mRNA.
loc126208	0.487	-2.052	zinc finger protein 787 (ZNF787), mRNA.
met	0.487	-2.052	met proto-oncogene (hepatocyte growth factor receptor) (MET), mRNA.
hadhsc	0.487	-2.052	L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain (HADHSC), mRNA.
os9	0.487	-2.052	amplified in osteosarcoma (OS9), transcript variant 1, mRNA.
ifp38	0.487	-2.052	IFP38 (IFP38), mRNA.
pycr2	0.487	-2.052	pyrroline-5-carboxylate reductase family, member 2 (PYCR2), mRNA.
acy1	0.487	-2.053	aminoacylase 1 (ACY1), mRNA.
nubp1	0.487	-2.053	nucleotide binding protein 1 (MinD homolog, E. coli) (NUBP1), mRNA.
tp53ap1	0.487	-2.054	TP53 activated protein 1 (TP53AP1), mRNA.
mrpl33	0.487	-2.054	mitochondrial ribosomal protein L33 (MRPL33), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
flj21827	0.487	-2.054	hypothetical protein FLJ21827 (FLJ21827), mRNA.
loc139886	0.487	-2.054	hypothetical protein LOC139886 (LOC139886), mRNA.
ddefl1	0.487	-2.054	development and differentiation enhancing factor-like 1 (DDEFL1), mRNA.
fksg24	0.487	-2.054	hypothetical protein MGC12972 (FKSG24), mRNA.
arhgef17	0.487	-2.054	Rho guanine nucleotide exchange factor (GEF) 17 (ARHGEF17), mRNA.
znf42	0.487	-2.054	myeloid zinc finger 1 (MZF1), transcript variant 2, mRNA.

saal1	0.487	-2.055	serum amyloid A-like 1 (SAAL1), mRNA.
flj41423	0.487	-2.055	FLJ41423 protein (FLJ41423), mRNA.
hs.576106	0.487	-2.055	cDNA FLJ40058 fis, clone TCOLN1000180
alg3	0.487	-2.055	asparagine-linked glycosylation 3homolog(yeast, alpha-1,3-mannosyltransferase)(ALG3),trans. variant 3, mRNA.
rfxank	0.487	-2.055	regulatory factor X-associated ankyrin-containing protein (RFXANK), transcript variant 1, mRNA.
osgep	0.487	-2.055	O-sialoglycoprotein endopeptidase (OSGEP), mRNA.
tmem55b	0.487	-2.056	transmembrane protein 55B (TMEM55B), mRNA.
hs.98330	0.487	-2.056	PREDICTED: hypothetical LOC388227 (LOC388227), mRNA
fkbp3	0.487	-2.056	FK506 binding protein 3, 25kDa (FKBP3), mRNA.
polr3f	0.486	-2.056	polymerase (RNA) III (DNA directed) polypeptide F, 39 kDa (POLR3F), mRNA.
zhx3	0.486	-2.056	zinc fingers and homeoboxes 3 (ZHX3), mRNA.
c9orf21	0.486	-2.056	chromosome 9 open reading frame 21 (C9orf21), mRNA.
snd1	0.486	-2.056	staphylococcal nuclease and tudor domain containing 1 (SND1), mRNA.
rnu3ip2	0.486	-2.057	RNA, U3 small nucleolar interacting protein 2 (RNU3IP2), mRNA.
aurka	0.486	-2.057	aurora kinase A (AURKA), transcript variant 3, mRNA.
ankrd16	0.486	-2.057	ankyrin repeat domain 16 (ANKRD16), transcript variant 2, mRNA.
c1orf164	0.486	-2.057	chromosome 1 open reading frame 164 (C1orf164), mRNA.
lgals3bp	0.486	-2.057	lectin, galactoside-binding, soluble, 3 binding protein (LGALS3BP), mRNA.
aste1	0.486	-2.057	asteroid homolog 1 (Drosophila) (ASTE1), mRNA.
stx12	0.486	-2.057	syntaxin 12 (STX12), mRNA.
dusp23	0.486	-2.058	dual specificity phosphatase 23 (DUSP23), mRNA.
slc45a4	0.486	-2.058	solute carrier family 45, member 4 (SLC45A4), mRNA.
pigo	0.486	-2.058	phosphatidylinositol glycan anchor biosynthesis, class O (PIGO), transcript variant 1, mRNA.
gmeb1	0.486	-2.058	glucocorticoid modulatory element binding protein 1 (GMEB1), transcript variant 2, mRNA.
hoxd13	0.486	-2.058	homeobox D13 (HOXD13), mRNA.
map3k6	0.486	-2.058	mitogen-activated protein kinase kinase kinase 6 (MAP3K6), mRNA.
cmtm6	0.486	-2.058	CKLF-like MARVEL transmembrane domain containing 6 (CMTM6), mRNA.
rabl2b	0.486	-2.059	RAB, member of RAS oncogene family-like 2B (RABL2B), transcript variant 2, mRNA.
swap70	0.486	-2.06	SWAP-70 protein (SWAP70), mRNA.
tbk1	0.486	-2.06	TANK-binding kinase 1 (TBK1), mRNA.
ccdc47	0.486	-2.06	coiled-coil domain containing 47 (CCDC47), mRNA.
plxna1	0.485	-2.06	plexin A1 (PLXNA1), mRNA.
cpsf3l	0.485	-2.06	cleavage and polyadenylation specific factor 3-like (CPSF3L), transcript variant 1, mRNA.
ube2t	0.485	-2.06	ubiquitin-conjugating enzyme E2T (putative) (UBE2T), mRNA.
dcbld1	0.485	-2.06	discoidin, CUB and LCCL domain containing 1 (DCBLD1), mRNA.
katnal2	0.485	-2.06	katanin p60 subunit A-like 2 (KATNAL2), mRNA.
csad	0.485	-2.06	cysteine sulfinic acid decarboxylase (CSAD), mRNA.
c19orf24	0.485	-2.061	chromosome 19 open reading frame 24 (C19orf24), mRNA.
flj21901	0.485	-2.061	FAST kinase domains 1 (FASTKD1), mRNA.
vcpip1	0.485	-2.061	valosin containing protein (p97)/p47 complex interacting protein 1 (VCPIP1), mRNA.
znf655	0.485	-2.061	zinc finger protein 655 (ZNF655), transcript variant 6, mRNA.
loc645345	0.485	-2.061	PREDICTED: similar to Coiled-coil-helix-coiled-coil-helix domain containing protein 2

			(HCV NS2 trans-regulated protein) (NS2TP) (LOC645345), mRNA.
flj13236	0.485	-2.062	hypothetical protein FLJ13236 (FLJ13236), mRNA.
phf20	0.485	-2.062	PHD finger protein 20 (PHF20), mRNA.
dync2h1	0.485	-2.062	dynein, cytoplasmic 2, heavy chain 1 (DYNC2H1), mRNA.
loc650832	0.485	-2.062	PREDICTED: similar to mitogen-activated protein kinase kinase 3 isoform A (LOC650832), mRNA.
yrdc	0.485	-2.062	yrdC domain containing (E. coli) (YRDC), mRNA.
txndc14	0.485	-2.062	thioredoxin domain containing 14 (TXNDC14), mRNA.
lamc1	0.485	-2.063	laminin, gamma 1 (formerly LAMB2) (LAMC1), mRNA.
cice	0.485	-2.063	cell death-inducing DFFA-like effector c pseudogene (CIDECP) on chromosome 3.
kctd9	0.485	-2.063	potassium channel tetramerisation domain containing 9 (KCTD9), mRNA.
psarl	0.485	-2.063	presenilin associated, rhomboid-like (PSARL), mRNA.
ifit5	0.485	-2.063	interferon-induced protein with tetratricopeptide repeats 5 (IFIT5), mRNA.
ndufb11	0.485	-2.063	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11, 17.3kDa (NDUFB11), mRNA.
sacm1l	0.485	-2.063	SAC1 suppressor of actin mutations 1-like (yeast) (SACM1L), mRNA.
xrcc6	0.485	-2.063	X-ray repair complementing defective repair in Chinese hamster cells 6 (Ku autoantigen, 70kDa) (XRCC6), mRNA.
unc50	0.485	-2.063	unc-50 homolog (C. elegans) (UNC50), mRNA.
flj20272	0.485	-2.064	tetratricopeptide repeat domain 27 (TTC27), mRNA.
flj10661	0.485	-2.064	family with sequence similarity 86, member C (FAM86C), transcript variant 2, mRNA.
mon2	0.485	-2.064	MON2 homolog (yeast) (MON2), mRNA.
coq10a	0.485	-2.064	coenzyme Q10 homolog A (S. cerevisiae) (COQ10A), mRNA.
wdr46	0.485	-2.064	WD repeat domain 46 (WDR46), mRNA.
rpl27a	0.485	-2.064	ribosomal protein L27a (RPL27A), mRNA.
ssr1	0.484	-2.064	signal sequence receptor, alpha (translocon-associated protein alpha) (SSR1), mRNA.
mn1	0.484	-2.065	meningioma (disrupted in balanced translocation) 1 (MN1), mRNA.
tmbim1	0.484	-2.065	transmembrane BAX inhibitor motif containing 1 (TMBIM1), mRNA.
impa2	0.484	-2.065	inositol(myo)-1(or 4)-monophosphatase 2 (IMPA2), mRNA.
loc643997	0.484	-2.065	PREDICTED: similar to peptidylprolyl isomerase A isoform 1 (LOC643997), mRNA.
cda08	0.484	-2.065	T-cell immunomodulatory protein (CDA08), mRNA.
genx-3414	0.484	-2.066	genethonin 1 (GENX-3414), mRNA.
ptges3	0.484	-2.066	prostaglandin E synthase 3 (cytosolic) (PTGES3), mRNA.
snx19	0.484	-2.066	sorting nexin 19 (SNX19), mRNA.
dcc1	0.484	-2.066	defective in sister chromatid cohesion homolog 1 (S. cerevisiae) (DCC1), mRNA.
apoa1bp	0.484	-2.066	apolipoprotein A-I binding protein (APOA1BP), mRNA.
prr3	0.484	-2.066	proline rich 3 (PRR3), mRNA.
them2	0.484	-2.066	thioesterase superfamily member 2 (THEM2), mRNA.
twistnb	0.484	-2.067	TWIST neighbor (TWISTNB), mRNA.
hs.10862	0.484	-2.067	cDNA: FLJ23313 fis, clone HEP11919
kiaa0274	0.484	-2.067	FIG4 homolog (S. cerevisiae) (FIG4), mRNA.
fchsd2	0.484	-2.067	FCH and double SH3 domains 2 (FCHSD2), mRNA.
mapk11	0.484	-2.067	mitogen-activated protein kinase 11 (MAPK11), transcript variant 2, mRNA.
gtf3a	0.484	-2.067	general transcription factor IIIA (GTF3A), mRNA.
fdps	0.484	-2.067	farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase,

			dimethylallyltranstransferase, geranyltranstransferase) (FDPS), mRNA.
hs.556082	0.484	-2.067	cDNA clone MGC:13035 IMAGE:3613409, complete cds
tpd52l1	0.484	-2.067	tumor protein D52-like 1 (TPD52L1), transcript variant 4, mRNA.
kctd13	0.484	-2.067	potassium channel tetramerisation domain containing 13 (KCTD13), mRNA.
loc541469	0.484	-2.067	hypothetical protein LOC541469 (LOC541469), mRNA.
serpinb6	0.484	-2.067	serpin peptidase inhibitor, clade B (ovalbumin), member 6 (SERPINB6), mRNA.
c20orf52	0.484	-2.067	chromosome 20 open reading frame 52 (C20orf52), mRNA.
znhit3	0.484	-2.068	zinc finger, HIT type 3 (ZNHIT3), transcript variant 1, mRNA.
kif14	0.484	-2.068	kinesin family member 14 (KIF14), mRNA.
hs.553301	0.484	-2.068	AV737317 CB cDNA clone CBCAQH03 5, mRNA sequence
mrpl20	0.484	-2.068	mitochondrial ribosomal protein L20 (MRPL20), nuclear gene encoding mitochondrial protein, mRNA.
kiaa0240	0.484	-2.068	KIAA0240 (KIAA0240), mRNA.
bri3	0.484	-2.068	brain protein I3 (BRI3), mRNA.
tfap2a	0.483	-2.068	transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha) (TFAP2A), transcript variant 1, mRNA.
atxn3	0.483	-2.068	ataxin 3 (ATXN3), transcript variant 2, mRNA.
myo9a	0.483	-2.069	myosin IXA (MYO9A), mRNA.
nipsnap1	0.483	-2.069	nipsnap homolog 1 (C. elegans) (NIPSNAP1), mRNA.
oprs1	0.483	-2.069	opioid receptor, sigma 1 (OPRS1), transcript variant 5, mRNA.
tmem80	0.483	-2.069	transmembrane protein 80 (TMEM80), mRNA.
amd1	0.483	-2.069	adenosylmethionine decarboxylase 1 (AMD1), transcript variant 1, mRNA.
fcho1	0.483	-2.069	FCH domain only 1 (FCHO1), mRNA.
psmd13	0.483	-2.069	proteasome (prosome, macropain) 26S subunit, non-ATPase, 13 (PSMD13), transcript variant 1, mRNA.
vps39	0.483	-2.07	vacuolar protein sorting 39 (yeast) (VPS39), mRNA.
phf19	0.483	-2.07	PHD finger protein 19 (PHF19), transcript variant 2, mRNA.
lsm5	0.483	-2.07	LSM5 homolog, U6 small nuclear RNA associated (S. cerevisiae) (LSM5), mRNA.
tmco1	0.483	-2.07	transmembrane and coiled-coil domains 1 (TMCO1), mRNA.
mnab	0.483	-2.07	ring finger and CCCH-type zinc finger domains 2 (RC3H2), mRNA.
cntnap1	0.483	-2.07	contactin associated protein 1 (CNTNAP1), mRNA.
hnrph1	0.483	-2.07	heterogeneous nuclear ribonucleoprotein H1 (H) (HNRPH1), mRNA.
bsg	0.483	-2.071	basigin (Ok blood group) (BSG), transcript variant 1, mRNA.
sema3a	0.483	-2.071	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A (SEMA3A), mRNA.
tm4sf18	0.483	-2.071	transmembrane 4 L six family member 18 (TM4SF18), mRNA.
flj20551	0.483	-2.071	solute carrier family 25, member 38 (SLC25A38), mRNA.
c11orf1	0.483	-2.071	chromosome 11 open reading frame 1 (C11orf1), mRNA.
coq3	0.483	-2.071	coenzyme Q3 homolog, methyltransferase (yeast) (COQ3), mRNA.
loc90355	0.483	-2.071	chromosome 5 open reading frame 30 (C5orf30), mRNA.
vps45a	0.483	-2.071	vacuolar protein sorting 45A (yeast) (VPS45A), mRNA.
c9orf64	0.483	-2.071	chromosome 9 open reading frame 64 (C9orf64), mRNA.
ccnd1	0.483	-2.071	cyclin D1 (PRAD1: parathyroid adenomatosis 1) (CCND1), mRNA.
ubxd6	0.483	-2.071	UBX domain containing 6 (UBXD6), mRNA.
rnmtl1	0.483	-2.071	RNA methyltransferase like 1 (RNMTL1), mRNA.
tbc1d9	0.483	-2.072	TBC1 domain family, member 9 (with GRAM domain) (TBC1D9), mRNA.

flj31951	0.483	-2.072	ring finger protein 145 (RNF145), mRNA.
wdr37	0.483	-2.072	WD repeat domain 37 (WDR37), mRNA.
flj10099	0.483	-2.072	hypothetical protein FLJ10099 (FLJ10099), mRNA.
dctn1	0.483	-2.072	dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 2, mRNA.
kiaa0194	0.483	-2.072	PREDICTED: KIAA0194 protein (KIAA0194), mRNA.
myh4	0.483	-2.072	myosin, heavy polypeptide 4, skeletal muscle (MYH4), mRNA.
pak1	0.483	-2.073	p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast) (PAK1), mRNA.
c1orf198	0.483	-2.073	chromosome 1 open reading frame 198 (C1orf198), mRNA.
hla-e	0.482	-2.073	major histocompatibility complex, class I, E (HLA-E), mRNA.
atp6v0e1	0.482	-2.073	ATPase, H+ transporting, lysosomal 9kDa, V0 subunit e1 (ATP6V0E1), mRNA.
pms2	0.482	-2.073	PMS2 postmeiotic segregation increased 2 (S. cerevisiae) (PMS2), transcript variant 1, mRNA.
cldnd2	0.482	-2.073	claudin domain containing 2 (CLDND2), mRNA.
ppox	0.482	-2.073	protoporphyrinogen oxidase (PPOX), nuclear gene encoding mitochondrial protein, mRNA.
gnb1	0.482	-2.073	guanine nucleotide binding protein (G protein), beta polypeptide 1 (GNB1), mRNA.
ddx1	0.482	-2.073	DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 (DDX1), mRNA.
tusc4	0.482	-2.073	tumor suppressor candidate 4 (TUSC4), mRNA.
nte	0.482	-2.073	neuropathy target esterase (NTE), mRNA.
cldn2	0.482	-2.074	claudin 2 (CLDN2), mRNA.
ccdc16	0.482	-2.074	coiled-coil domain containing 16 (CCDC16), mRNA.
plxnb2	0.482	-2.074	PREDICTED: plexin B2 (PLXNB2), mRNA.
pex6	0.482	-2.074	peroxisomal biogenesis factor 6 (PEX6), mRNA.
pigv	0.482	-2.074	phosphatidylinositol glycan anchor biosynthesis, class V (PIGV), mRNA.
cast	0.482	-2.074	calpastatin (CAST), transcript variant 8, mRNA.
tp53inp2	0.482	-2.074	tumor protein p53 inducible nuclear protein 2 (TP53INP2), mRNA.
erich1	0.482	-2.074	glutamate-rich 1 (ERICH1), mRNA.
dcun1d5	0.482	-2.074	DCN1, defective in cullin neddylation 1, domain containing 5 (S. cerevisiae) (DCUN1D5), mRNA.
ash2l	0.482	-2.074	ash2 (absent, small, or homeotic)-like (Drosophila) (ASH2L), mRNA.
pmpcb	0.482	-2.075	peptidase (mitochondrial processing) beta (PMPCB), mRNA.
ostm1	0.482	-2.075	osteopetrosis associated transmembrane protein 1 (OSTM1), mRNA.
etv4	0.482	-2.075	ets variant gene 4 (E1A enhancer binding protein, E1AF) (ETV4), mRNA.
zfx	0.482	-2.075	zinc finger protein, X-linked (ZFX), mRNA.
hspc023	0.482	-2.076	HSPC023 protein (HSPC023), mRNA.
paqr7	0.482	-2.076	progestin and adipoQ receptor family member VII (PAQR7), mRNA.
flj23322	0.482	-2.076	hypothetical protein FLJ23322 (FLJ23322), mRNA.
nip7	0.482	-2.077	nuclear import 7 homolog (S. cerevisiae) (NIP7), mRNA.
myo1c	0.482	-2.077	myosin IC (MYO1C), mRNA.
dlst	0.481	-2.077	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) (DLST), mRNA.
slc25a28	0.481	-2.077	solute carrier family 25, member 28 (SLC25A28), mRNA.
cdc42bpa	0.481	-2.077	CDC42 binding protein kinase alpha (DMPK-like) (CDC42BPA), transcript variant B, mRNA.
dgcr8	0.481	-2.078	DiGeorge syndrome critical region gene 8 (DGCR8), mRNA.
arcn1	0.481	-2.078	archain 1 (ARCN1), mRNA.
rw1	0.481	-2.078	PREDICTED: RW1 protein, transcript variant 1 (RW1), mRNA.

nthl1	0.481	-2.078	nth endonuclease III-like 1 (E. coli) (NTHL1), mRNA.
golga1	0.481	-2.078	golgi autoantigen, golgin subfamily a, 1 (GOLGA1), mRNA.
c6orf148	0.481	-2.078	chromosome 6 open reading frame 148 (C6orf148), mRNA.
usp24	0.481	-2.078	PREDICTED: ubiquitin specific peptidase 24, transcript variant 6 (USP24), mRNA.
cep63	0.481	-2.079	centrosomal protein 63kDa (CEP63), transcript variant 2, mRNA.
rpl11	0.481	-2.079	ribosomal protein L11 (RPL11), mRNA.
rbm17	0.481	-2.079	RNA binding motif protein 17 (RBM17), mRNA.
smyd3	0.481	-2.079	SET and MYND domain containing 3 (SMYD3), mRNA.
rad51c	0.481	-2.08	RAD51 homolog C (S. cerevisiae) (RAD51C), transcript variant 1, mRNA.
mrpl51	0.481	-2.08	mitochondrial ribosomal protein L51 (MRPL51), nuclear gene encoding mitochondrial protein, mRNA.
trpm4	0.481	-2.08	transient receptor potential cation channel, subfamily M, member 4 (TRPM4), mRNA.
nenf	0.481	-2.08	neuron derived neurotrophic factor (NENF), mRNA.
mgc11257	0.481	-2.08	hypothetical protein MGC11257 (MGC11257), mRNA.
ids	0.481	-2.08	iduronate 2-sulfatase (Hunter syndrome) (IDS), transcript variant 2, mRNA.
mrpl55	0.481	-2.08	mitochondrial ribosomal protein L55 (MRPL55), nuclear gene encoding mitochondrial protein, transcript variant 5, mRNA.
cntrob	0.481	-2.08	centrobin, centrosomal BRCA2 interacting protein (CNTROB), transcript variant 1, mRNA.
smarca4	0.481	-2.08	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 (SMARCA4), mRNA.
loc284393	0.481	-2.081	PREDICTED: similar to ribosomal protein L10, transcript variant 1 (LOC284393), mRNA.
acp6	0.481	-2.081	acid phosphatase 6, lysophosphatidic (ACP6), mRNA.
lgmn	0.481	-2.081	legumain (LGMN), transcript variant 1, mRNA.
c1orf50	0.481	-2.081	chromosome 1 open reading frame 50 (C1orf50), mRNA.
tmed5	0.48	-2.081	transmembrane emp24 protein transport domain containing 5 (TMED5), mRNA.
gemin4	0.48	-2.081	gem (nuclear organelle) associated protein 4 (GEMIN4), mRNA.
ercc3	0.48	-2.082	excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing) (ERCC3), mRNA.
srpk2	0.48	-2.082	SFRS protein kinase 2 (SRPK2), transcript variant 2, mRNA.
pcid1	0.48	-2.082	eukaryotic translation initiation factor 3, subunit M (EIF3M), mRNA.
cdc23	0.48	-2.082	CDC23 (cell division cycle 23, yeast, homolog) (CDC23), mRNA.
hccs	0.48	-2.082	holocytochrome c synthase (cytochrome c heme-lyase) (HCCS), mRNA.
dnaptp6	0.48	-2.082	DNA polymerase-transactivated protein 6 (DNAPTP6), mRNA.
mask	0.48	-2.082	Mst3 and SOK1-related kinase (MASK), mRNA.
nmb	0.48	-2.082	neuromedin B (NMB), transcript variant 1, mRNA.
itm2c	0.48	-2.082	integral membrane protein 2C (ITM2C), transcript variant 2, mRNA.
cep164	0.48	-2.082	centrosomal protein 164kDa (CEP164), mRNA.
arv1	0.48	-2.082	ARV1 homolog (S. cerevisiae) (ARV1), mRNA.
pdcd6ip	0.48	-2.083	programmed cell death 6 interacting protein (PDCD6IP), mRNA.
rnaseh2a	0.48	-2.083	ribonuclease H2, subunit A (RNASEH2A), mRNA.
loc653974	0.48	-2.083	PREDICTED: similar to ribosomal protein L15, transcript variant 3 (LOC653974), mRNA.
h2afz	0.48	-2.084	H2A histone family, member Z (H2AFZ), mRNA.
dhrsx	0.48	-2.084	dehydrogenase/reductase (SDR family) X-linked (DHRSX), mRNA.

wwp1	0.48	-2.084	WW domain containing E3 ubiquitin protein ligase 1 (WWP1), mRNA.
pros1	0.48	-2.084	protein S (alpha) (PROS1), mRNA.
sec6l1	0.48	-2.084	SEC6-like 1 (S. cerevisiae) (SEC6L1), mRNA.
nsun5b	0.48	-2.085	NOL1/NOP2/Sun domain family, member 5B (NSUN5B), transcript variant 2, mRNA.
crk	0.48	-2.085	v-crk sarcoma virus CT10 oncogene homolog (avian) (CRK), transcript variant II, mRNA.
diaph1	0.48	-2.085	diaphanous homolog 1 (Drosophila) (DIAPH1), transcript variant 1, mRNA.
hspa14	0.48	-2.085	heat shock 70kDa protein 14 (HSPA14), mRNA.
rpap1	0.48	-2.085	RNA polymerase II associated protein 1 (RPAP1), mRNA.
timm22	0.48	-2.086	translocase of inner mitochondrial membrane 22 homolog (yeast) (TIMM22), mRNA.
pcnt2	0.479	-2.086	pericentrin 2 (kendrin) (PCNT2), mRNA.
ranbp6	0.479	-2.086	RAN binding protein 6 (RANBP6), mRNA.
flj25476	0.479	-2.086	FLJ25476 protein (FLJ25476), mRNA.
flj90652	0.479	-2.086	coiled-coil domain containing 95 (CCDC95), mRNA.
sepx1	0.479	-2.087	selenoprotein X, 1 (SEPX1), mRNA.
loc643657	0.479	-2.087	PREDICTED: similar to Nonhistone chromosomal protein HMG-17 (High-mobility group nucleosome binding domain 2) (LOC643657), mRNA.
tchp	0.479	-2.087	trichoplein, keratin filament binding (TCHP), mRNA.
iscu	0.479	-2.087	iron-sulfur cluster scaffold homolog (E. coli) (ISCU), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA
c14orf124	0.479	-2.087	chromosome 14 open reading frame 124 (C14orf124), mRNA.
galm	0.479	-2.087	galactose mutarotase (aldose 1-epimerase) (GALM), mRNA.
fhl2	0.479	-2.087	four and a half LIM domains 2 (FHL2), transcript variant 4, mRNA.
pole4	0.479	-2.087	polymerase (DNA-directed), epsilon 4 (p12 subunit) (POLE4), mRNA.
mrpl43	0.479	-2.087	mitochondrial ribosomal protein L43 (MRPL43), nuclear gene encoding mitochondrial protein, transcript variant 4, mRNA.
c16orf80	0.479	-2.088	chromosome 16 open reading frame 80 (C16orf80), mRNA.
lancl2	0.479	-2.088	LanC lantibiotic synthetase component C-like 2 (bacterial) (LANCL2), mRNA.
nt5c3	0.479	-2.088	5'-nucleotidase, cytosolic III (NT5C3), transcript variant 3, mRNA.
ubqln4	0.479	-2.088	ubiquilin 4 (UBQLN4), mRNA.
zdhhc12	0.479	-2.088	zinc finger, DHHC-type containing 12 (ZDHHC12), mRNA.
wdr26	0.479	-2.088	WD repeat domain 26 (WDR26), mRNA.
rragd	0.479	-2.088	Ras-related GTP binding D (RRAGD), mRNA.
tmem115	0.479	-2.088	transmembrane protein 115 (TMEM115), mRNA.
rnf44	0.479	-2.088	ring finger protein 44 (RNF44), mRNA.
loc647727	0.479	-2.088	PREDICTED: similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein)(hnRNP core protein A1)(HDP-1)(Topoisomerase-inhibitor suppressed)(LOC647727),mRNA
cdk9	0.479	-2.088	cyclin-dependent kinase 9 (CDC2-related kinase) (CDK9), mRNA.
c21orf45	0.479	-2.089	chromosome 21 open reading frame 45 (C21orf45), mRNA.
ptprm	0.479	-2.09	protein tyrosine phosphatase, receptor type, M (PTPRM), mRNA.
dhx38	0.479	-2.09	DEAH (Asp-Glu-Ala-His) box polypeptide 38 (DHX38), mRNA.
ruvbl2	0.478	-2.09	RuvB-like 2 (E. coli) (RUVBL2), mRNA.
loc198437	0.478	-2.09	bA299N6.3 (LOC198437), mRNA.

loc647389	0.478	-2.09	PREDICTED: hypothetical protein LOC647389 (LOC647389), mRNA.
ap4e1	0.478	-2.09	adaptor-related protein complex 4, epsilon 1 subunit (AP4E1), mRNA.
tegt	0.478	-2.09	testis enhanced gene transcript (BAX inhibitor 1) (TEGT), mRNA.
loc643284	0.478	-2.09	PREDICTED: similar to 40S ribosomal protein S29 (LOC643284), mRNA.
fkbp11	0.478	-2.091	FK506 binding protein 11, 19 kDa (FKBP11), mRNA.
sfn	0.478	-2.091	stratifin (SFN), mRNA.
galgt	0.478	-2.091	UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)- galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T) (GALGT), mRNA.
nup188	0.478	-2.091	nucleoporin 188kDa (NUP188), mRNA.
pctp	0.478	-2.091	phosphatidylcholine transfer protein (PCTP), mRNA.
kiaa0179	0.478	-2.092	KIAA0179 (KIAA0179), mRNA.
rwdd4a	0.478	-2.092	RWD domain containing 4A (RWDD4A), mRNA.
loc221955	0.478	-2.092	diacylglycerol lipase, beta (DAGLB), mRNA.
mtmr9	0.478	-2.092	myotubularin related protein 9 (MTMR9), mRNA.
ttk	0.478	-2.092	TTK protein kinase (TTK), mRNA.
dus2l	0.478	-2.092	dihydrouridine synthase 2-like, SMM1 homolog (S. cerevisiae) (DUS2L), mRNA.
pdss1	0.478	-2.092	prenyl (decaprenyl) diphosphate synthase, subunit 1 (PDSS1), mRNA.
gfod1	0.478	-2.092	glucose-fructose oxidoreductase domain containing 1 (GFOD1), mRNA.
aadac	0.478	-2.092	arylacetamide deacetylase (esterase) (AADAC), mRNA.
loc651894	0.478	-2.092	PREDICTED: similar to ribosomal protein S12 (LOC651894), mRNA.
tmed10p	0.478	-2.093	transmembrane emp24-like trafficking protein 10 (yeast) pseudogene (TMED10P) on chromosome 8.
nptn	0.478	-2.093	neuroplastin (NPTN), transcript variant alpha, mRNA.
eif3s2	0.478	-2.093	eukaryotic translation initiation factor 3, subunit 2 beta, 36kDa (EIF3S2), mRNA.
twsg1	0.478	-2.093	twisted gastrulation homolog 1 (Drosophila) (TWSG1), mRNA.
ppp1r3e	0.478	-2.093	PREDICTED: protein phosphatase 1, regulatory (inhibitor) subunit 3E (PPP1R3E), mRNA.
wdr39	0.478	-2.093	WD repeat domain 39 (WDR39), mRNA.
ptprk	0.478	-2.093	protein tyrosine phosphatase, receptor type, K (PTPRK), mRNA.
loc646786	0.478	-2.093	PREDICTED: similar to Afadin (AF-6 protein) (LOC646786), mRNA.
oxa1l	0.478	-2.094	oxidase (cytochrome c) assembly 1-like (OXA1L), mRNA.
ppp2r2a	0.478	-2.094	protein phosphatase 2 (formerly 2A), regulatory subunit B, alpha isoform (PPP2R2A), mRNA.
dab2	0.478	-2.094	disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) (DAB2), mRNA.
atp5f1	0.477	-2.094	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit b, isoform 1 (ATP5F1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
tnfaip6	0.477	-2.095	tumor necrosis factor, alpha-induced protein 6 (TNFAIP6), mRNA.
fam44b	0.477	-2.095	family with sequence similarity 44, member B (FAM44B), mRNA.
znf621	0.477	-2.095	zinc finger protein 621 (ZNF621), mRNA.
ndufb2	0.477	-2.096	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2, 8kDa (NDUFB2), nuclear gene encoding mitochondrial protein, mRNA.
ptdss1	0.477	-2.096	phosphatidylserine synthase 1 (PTDSS1), mRNA.
c12orf52	0.477	-2.096	chromosome 12 open reading frame 52 (C12orf52), mRNA.
atg10	0.477	-2.096	ATG10 autophagy related 10 homolog (S. cerevisiae) (ATG10), mRNA.
ilkap	0.477	-2.096	integrin-linked kinase-associated serine/threonine phosphatase 2C (ILKAP), transcript variant 2, mRNA.

c20orf155	0.477	-2.097	cardiolipin synthase 1 (CRLS1), mRNA.
c14orf2	0.477	-2.097	chromosome 14 open reading frame 2 (C14orf2), mRNA.
dhrs4	0.477	-2.097	dehydrogenase/reductase (SDR family) member 4 (DHRS4), mRNA.
dlg4	0.477	-2.097	discs, large homolog 4 (Drosophila) (DLG4), mRNA.
rpl23ap7	0.477	-2.098	ribosomal protein L23a pseudogene 7 (RPL23AP7) on chromosome 2.
mtpn	0.477	-2.098	myotrophin (MTPN), mRNA.
yaf2	0.477	-2.098	YY1 associated factor 2 (YAF2), transcript variant 2, mRNA.
lass5	0.477	-2.098	LAG1 homolog, ceramide synthase 5 (S. cerevisiae) (LASS5), mRNA.
slc33a1	0.477	-2.098	solute carrier family 33 (acetyl-CoA transporter), member 1 (SLC33A1), mRNA.
cstf3	0.477	-2.099	cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kDa (CSTF3), transcript variant 2, mRNA.
raver1	0.477	-2.099	RAVER1 (RAVER1), mRNA.
utp15	0.476	-2.099	UTP15, U3 small nucleolar ribonucleoprotein, homolog (S. cerevisiae) (UTP15), mRNA.
pafah2	0.476	-2.099	platelet-activating factor acetylhydrolase 2, 40kDa (PAFAH2), mRNA.
c18orf24	0.476	-2.099	chromosome 18 open reading frame 24 (C18orf24), mRNA.
c4orf14	0.476	-2.099	chromosome 4 open reading frame 14 (C4orf14), mRNA.
ggh	0.476	-2.1	gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl hydrolase) (GGH), mRNA.
dedd	0.476	-2.1	death effector domain containing (DEDD), transcript variant 2, mRNA.
mgc3196	0.476	-2.1	PREDICTED: hypothetical protein MGC3196 (MGC3196), mRNA.
c12orf43	0.476	-2.1	chromosome 12 open reading frame 43 (C12orf43), mRNA.
s100a16	0.476	-2.1	S100 calcium binding protein A16 (S100A16), mRNA.
ergic2	0.476	-2.101	ERGIC and golgi 2 (ERGIC2), mRNA.
lars	0.476	-2.101	leucyl-tRNA synthetase (LARS), mRNA.
emd	0.476	-2.101	emerin (Emery-Dreifuss muscular dystrophy) (EMD), mRNA.
plau	0.476	-2.101	plasminogen activator, urokinase (PLAU), mRNA.
man2a1	0.476	-2.102	mannosidase, alpha, class 2A, member 1 (MAN2A1), mRNA.
afap1l2	0.476	-2.102	actin filament associated protein 1-like 2 (AFAP1L2), transcript variant 2, mRNA.
map2k4	0.476	-2.102	mitogen-activated protein kinase kinase 4 (MAP2K4), mRNA.
atp9b	0.476	-2.102	ATPase, Class II, type 9B (ATP9B), mRNA.
ncl	0.476	-2.102	nucleolin (NCL), mRNA.
nr1h2	0.476	-2.103	nuclear receptor subfamily 1, group H, member 2 (NR1H2), mRNA.
mrps22	0.475	-2.103	mitochondrial ribosomal protein S22 (MRPS22), nuclear gene encoding mitochondrial protein, mRNA.
paqr5	0.475	-2.104	progestin and adipoQ receptor family member V (PAQR5), mRNA.
c14orf131	0.475	-2.104	chromosome 14 open reading frame 131 (C14orf131), mRNA.
znf297b	0.475	-2.104	zinc finger and BTB domain containing 43 (ZBTB43), mRNA.
slc9a8	0.475	-2.104	solute carrier family 9 (sodium/hydrogen exchanger), member 8 (SLC9A8), mRNA.
ap3b1	0.475	-2.104	adaptor-related protein complex 3, beta 1 subunit (AP3B1), mRNA.
habp4	0.475	-2.104	hyaluronan binding protein 4 (HABP4), mRNA.
accn3	0.475	-2.104	amiloride-sensitive cation channel 3 (ACCN3), transcript variant 2, mRNA.
vps4b	0.475	-2.105	vacuolar protein sorting 4B (yeast) (VPS4B), mRNA.
rps6	0.475	-2.105	ribosomal protein S6 (RPS6), mRNA.
lamp1	0.475	-2.105	lysosomal-associated membrane protein 1 (LAMP1), mRNA.
rpl21	0.475	-2.105	ribosomal protein L21 (RPL21), mRNA.

prkaa1	0.475	-2.106	protein kinase, AMP-activated, alpha 1 catalytic subunit (PRKAA1), transcript variant 2, mRNA.
ahi1	0.475	-2.106	Abelson helper integration site 1 (AHI1), mRNA.
lama5	0.475	-2.106	laminin, alpha 5 (LAMA5), mRNA.
loc648314	0.475	-2.106	PREDICTED: similar to ribosomal protein L4, transcript variant 2 (LOC648314), mRNA.
gstm2	0.475	-2.106	glutathione S-transferase M2 (muscle) (GSTM2), mRNA.
zdhhc16	0.475	-2.106	zinc finger, DHHC-type containing 16 (ZDHHC16), transcript variant 3, mRNA.
loc129285	0.475	-2.106	coiled-coil domain containing 128 (CCDC128), mRNA.
thyn1	0.475	-2.107	thymocyte nuclear protein 1 (THYN1), transcript variant 2, mRNA.
arl2bp	0.475	-2.107	ADP-ribosylation factor-like 2 binding protein (ARL2BP), mRNA.
atp5j2	0.475	-2.107	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F2 (ATP5J2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
actn1	0.475	-2.107	actinin, alpha 1 (ACTN1), mRNA.
hdac3	0.475	-2.107	histone deacetylase 3 (HDAC3), mRNA.
btd	0.475	-2.107	biotinidase (BTD), mRNA.
chmp6	0.475	-2.107	chromatin modifying protein 6 (CHMP6), mRNA.
smarca1	0.475	-2.107	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 (SMARCA1), transcript variant 1, mRNA.
loc650495	0.475	-2.107	PREDICTED: hypothetical LOC650495 (LOC650495), mRNA.
vps25	0.475	-2.107	vacuolar protein sorting 25 homolog (S. cerevisiae) (VPS25), mRNA.
ube2l3	0.474	-2.108	ubiquitin-conjugating enzyme E2L 3 (UBE2L3), transcript variant 2, mRNA.
cbx6	0.474	-2.108	chromobox homolog 6 (CBX6), mRNA.
kiaa0286	0.474	-2.108	KIAA0286 protein (KIAA0286), mRNA.
svep1	0.474	-2.108	sushi, von Willebrand factor type A, EGF and pentraxin domain containing 1 (SVEP1), mRNA.
arfip1	0.474	-2.108	ADP-ribosylation factor interacting protein 1 (arfaptin 1) (ARFIP1), transcript variant 1, mRNA.
pim1	0.474	-2.108	pim-1 oncogene (PIM1), mRNA.
ndufa9	0.474	-2.108	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9, 39kDa (NDUFA9), mRNA.
klk3	0.474	-2.109	kallikrein-related peptidase 3 (KLK3), transcript variant 6, mRNA.
hist1h2ac	0.474	-2.109	histone cluster 1, H2ac (HIST1H2AC), mRNA.
loc127295	0.474	-2.109	PREDICTED: similar to ribosomal protein L36 (LOC127295), mRNA.
bcas4	0.474	-2.11	breast carcinoma amplified sequence 4 (BCAS4), transcript variant 1, mRNA.
raph1	0.474	-2.11	Ras association (RalGDS/AF-6) and pleckstrin homology domains 1 (RAPH1), transcript variant 1, mRNA.
tfrc	0.474	-2.11	transferrin receptor (p90, CD71) (TFRC), mRNA.
znf669	0.474	-2.11	zinc finger protein 669 (ZNF669), mRNA.
pnpo	0.474	-2.11	pyridoxine 5'-phosphate oxidase (PNPO), mRNA.
sco1	0.474	-2.11	SCO cytochrome oxidase deficient homolog 1 (yeast) (SCO1), nuclear gene encoding mitochondrial protein, mRNA
upf2	0.474	-2.111	UPF2 regulator of nonsense transcripts homolog (yeast) (UPF2), transcript variant 1, mRNA.
adrb2	0.474	-2.111	adrenergic, beta-2-, receptor, surface (ADRB2), mRNA.
ugp2	0.474	-2.111	UDP-glucose pyrophosphorylase 2 (UGP2), transcript variant 2, mRNA.
ubl4a	0.474	-2.111	ubiquitin-like 4A (UBL4A), mRNA.
m6pr	0.474	-2.111	mannose-6-phosphate receptor (cation dependent) (M6PR), mRNA.
dhx15	0.474	-2.111	DEAH (Asp-Glu-Ala-His) box polypeptide 15 (DHX15), mRNA.
cfd	0.473	-2.112	complement factor D (adipsin) (CFD), mRNA.

leo1	0.473	-2.112	Leo1, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae) (LEO1), mRNA.
mgc33648	0.473	-2.112	chromosome 5 open reading frame 35 (C5orf35), mRNA.
mgc16385	0.473	-2.112	hypothetical protein MGC16385 (MGC16385), mRNA.
clptm1	0.473	-2.113	cleft lip and palate associated transmembrane protein 1 (CLPTM1), mRNA.
wee1	0.473	-2.113	WEE1 homolog (S. pombe) (WEE1), mRNA.
slc19a1	0.473	-2.113	solute carrier family 19 (folate transporter), member 1 (SLC19A1), transcript variant 1, mRNA.
rplp2	0.473	-2.113	ribosomal protein, large, P2 (RPLP2), mRNA.
rbm14	0.473	-2.113	RNA binding motif protein 14 (RBM14), mRNA.
hsd3b7	0.473	-2.114	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7 (HSD3B7), mRNA.
cyb5r3	0.473	-2.114	cytochrome b5 reductase 3 (CYB5R3), transcript variant S, mRNA.
hs.535360	0.473	-2.114	cDNA clone IMAGE:6250506
sltm	0.473	-2.114	SAFB-like, transcription modulator (SLTM), transcript variant 1, mRNA.
echs1	0.473	-2.115	enoyl Coenzyme A hydratase, short chain, 1, mitochondrial (ECHS1), nuclear gene encoding mitochondrial protein, mRNA.
tspan6	0.473	-2.115	tetraspanin 6 (TSPAN6), mRNA.
c2orf18	0.473	-2.115	chromosome 2 open reading frame 18 (C2orf18), mRNA.
odz3	0.473	-2.115	odz, odd Oz/ten-m homolog 3 (Drosophila) (ODZ3), mRNA.
tmem42	0.473	-2.115	transmembrane protein 42 (TMEM42), mRNA.
hs.40289	0.473	-2.115	mRNA; cDNA DKFZp686I23208 (from clone DKFZp686I23208)
csnk2a2	0.473	-2.115	casein kinase 2, alpha prime polypeptide (CSNK2A2), mRNA.
mycbp2	0.473	-2.115	MYC binding protein 2 (MYCBP2), mRNA.
samm50	0.473	-2.115	sorting and assembly machinery component 50 homolog (S. cerevisiae) (SAMM50), mRNA.
ube2d4	0.473	-2.116	ubiquitin-conjugating enzyme E2D 4 (putative) (UBE2D4), mRNA.
rb1cc1	0.473	-2.116	RB1-inducible coiled-coil 1 (RB1CC1), mRNA.
rbm5	0.473	-2.116	RNA binding motif protein 5 (RBM5), mRNA.
dync1li2	0.472	-2.116	dynein, cytoplasmic 1, light intermediate chain 2 (DYNC1LI2), mRNA.
herc6	0.472	-2.117	hect domain and RLD 6 (HERC6), transcript variant 4, mRNA.
xrcc5	0.472	-2.117	X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kDa) (XRCC5), mRNA.
sfxn4	0.472	-2.117	sideroflexin 4 (SFXN4), transcript variant 2, mRNA.
sh3bgrl3	0.472	-2.117	SH3 domain binding glutamic acid-rich protein like 3 (SH3BGRL3), mRNA.
aftph	0.472	-2.117	aftiphilin (AFTPH), transcript variant 2, mRNA.
styxl1	0.472	-2.117	serine/threonine/tyrosine interacting-like 1 (STYXL1), mRNA.
loc221710	0.472	-2.117	PREDICTED: hypothetical protein LOC221710 (LOC221710), mRNA.
dhtkd1	0.472	-2.117	dehydrogenase E1 and transketolase domain containing 1 (DHTKD1), mRNA.
raxl1	0.472	-2.118	retina and anterior neural fold homeobox like 1 (RAXL1), mRNA.
dus3l	0.472	-2.118	dihydrouridine synthase 3-like (S. cerevisiae) (DUS3L), mRNA.
mcm8	0.472	-2.118	minichromosome maintenance complex component 8 (MCM8), transcript variant 1, mRNA.
eftud2	0.472	-2.118	elongation factor Tu GTP binding domain containing 2 (EFTUD2), mRNA.
hdhd1a	0.472	-2.118	haloacid dehalogenase-like hydrolase domain containing 1A (HDHD1A), mRNA.
abc1	0.472	-2.119	amplified in breast cancer 1 (ABC1), mRNA.
gmps	0.472	-2.119	guanine monphosphate synthetase (GMPS), mRNA.

dchs1	0.472	-2.119	dachsous 1 (Drosophila) (DCHS1), mRNA.
loc150223	0.472	-2.119	hypothetical protein LOC150223 (LOC150223), transcript variant 2, mRNA.
hs.535044	0.472	-2.119	PREDICTED: similar to FLJ00290 protein (LOC441310), mRNA
mkln1	0.472	-2.119	muskelin 1, intracellular mediator containing kelch motifs (MKLN1), mRNA.
zmym1	0.472	-2.119	zinc finger, MYM-type 1 (ZMYM1), mRNA.
glyctk	0.472	-2.119	glycerate kinase (GLYCTK), mRNA.
dock1	0.472	-2.12	dedicator of cytokinesis 1 (DOCK1), mRNA.
atf6	0.472	-2.12	activating transcription factor 6 (ATF6), mRNA.
ankrd38	0.472	-2.12	ankyrin repeat domain 38 (ANKRD38), mRNA.
metrn	0.472	-2.12	meteorin, glial cell differentiation regulator (METRN), mRNA.
spats2	0.472	-2.12	spermatogenesis associated, serine-rich 2 (SPATS2), mRNA.
bok	0.472	-2.12	BCL2-related ovarian killer (BOK), mRNA.
c1orf48	0.472	-2.121	chromosome 1 open reading frame 48 (C1orf48), mRNA.
pkm2	0.471	-2.121	pyruvate kinase, muscle (PKM2), transcript variant 1, mRNA.
srp46	0.471	-2.122	Splicing factor, arginine/serine-rich, 46kD (SRP46), mRNA.
mgc12981	0.471	-2.122	coiled-coil domain containing 115 (CCDC115), mRNA.
arhgef19	0.471	-2.122	Rho guanine nucleotide exchange factor (GEF) 19 (ARHGEF19), mRNA.
abcc4	0.471	-2.122	ATP-binding cassette, sub-family C (CFTR/MRP), member 4 (ABCC4), mRNA.
loc644979	0.471	-2.122	PREDICTED: hypothetical LOC644979 (LOC644979), mRNA.
chchd1	0.471	-2.122	coiled-coil-helix-coiled-coil-helix domain containing 1 (CHCHD1), mRNA.
cdc2l6	0.471	-2.122	cell division cycle 2-like 6 (CDK8-like) (CDC2L6), mRNA.
loc646630	0.471	-2.123	PREDICTED: similar to Coiled-coil-helix-coiled-coil-helix domain containing protein 2 (HCV NS2 trans-regulated protein) (NS2TP) (LOC646630), mRNA.
hs.485155	0.471	-2.123	cDNA: FLJ22515 fis, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence
flj45557	0.471	-2.123	hypothetical protein LOC642938 (FLJ45557), mRNA.
psmc3	0.471	-2.123	proteasome (prosome, macropain) 26S subunit, ATPase, 3 (PSMC3), mRNA.
chst10	0.471	-2.123	carbohydrate sulfotransferase 10 (CHST10), mRNA.
agr2	0.471	-2.123	anterior gradient homolog 2 (Xenopus laevis) (AGR2), mRNA.
pro0149	0.471	-2.123	chromosome 16 open reading frame 72 (C16orf72), mRNA.
otub2	0.471	-2.123	OTU domain, ubiquitin aldehyde binding 2 (OTUB2), mRNA.
cyp20a1	0.471	-2.123	cytochrome P450, family 20, subfamily A, polypeptide 1 (CYP20A1), transcript variant 1, mRNA.
loc652846	0.471	-2.124	PREDICTED: similar to Annexin A8 (Annexin VIII) (Vascular anticoagulant-beta) (VAC-beta) (LOC652846), mRNA.
entpd6	0.471	-2.124	ectonucleoside triphosphate diphosphohydrolase 6 (putative function) (ENTPD6), mRNA.
prdx4	0.471	-2.124	peroxiredoxin 4 (PRDX4), mRNA.
stip1	0.471	-2.124	stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein) (STIP1), mRNA.
fam38a	0.471	-2.124	family with sequence similarity 38, member A (FAM38A), mRNA.
npip	0.471	-2.124	nuclear pore complex interacting protein (NPIP), mRNA.
tmem29	0.471	-2.124	transmembrane protein 29 (TMEM29), mRNA.
rnaseh1	0.471	-2.125	ribonuclease H1 (RNASEH1), mRNA.
fcrlm2	0.471	-2.125	Fc receptor-like B (FCRLB), mRNA.
loc648210	0.471	-2.125	PREDICTED: similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein) (hnRNP core protein A1) (HDP-1) (Topoisomerase-inhibitor suppressed)

			, transcript variant 1 (LOC648210), mRNA.
hs.22907	0.471	-2.125	mRNA; cDNA DKFZp686P0492 (from clone DKFZp686P0492)
saps3	0.471	-2.125	SAPS domain family, member 3 (SAPS3), mRNA.
evc	0.471	-2.125	Ellis van Creveld syndrome (EVC), transcript variant 2, mRNA.
brms1l	0.471	-2.125	breast cancer metastasis-suppressor 1-like (BRMS1L), mRNA.
tmsb10	0.471	-2.125	thymosin, beta 10 (TMSB10), mRNA.
tspan9	0.471	-2.125	tetraspanin 9 (TSPAN9), mRNA.
slc2a4rg	0.471	-2.125	SLC2A4 regulator (SLC2A4RG), mRNA.
pin1	0.47	-2.126	protein (peptidylprolyl cis/trans isomerase) NIMA-interacting 1 (PIN1), mRNA.
rtcd1	0.47	-2.126	RNA terminal phosphate cyclase domain 1 (RTCD1), mRNA.
znf419	0.47	-2.126	zinc finger protein 419 (ZNF419), mRNA.
kif18a	0.47	-2.126	kinesin family member 18A (KIF18A), mRNA.
rps13	0.47	-2.127	ribosomal protein S13 (RPS13), mRNA.
exosc7	0.47	-2.127	exosome component 7 (EXOSC7), mRNA.
pepd	0.47	-2.127	peptidase D (PEPD), mRNA.
gnb1l	0.47	-2.127	guanine nucleotide binding protein (G protein), beta polypeptide 1-like (GNB1L), mRNA.
sec24c	0.47	-2.128	SEC24 related gene family, member C (S. cerevisiae) (SEC24C), transcript variant 1, mRNA.
et	0.47	-2.128	major facilitator superfamily domain containing 11 (MFSD11), mRNA.
c1orf123	0.47	-2.128	chromosome 1 open reading frame 123 (C1orf123), mRNA.
loc338758	0.47	-2.128	PREDICTED: hypothetical protein LOC338758 (LOC338758), mRNA.
rpl3	0.47	-2.128	ribosomal protein L3 (RPL3), transcript variant 2, mRNA.
tuba1	0.47	-2.129	tubulin, alpha 4a (TUBA4A), mRNA.
mimitin	0.47	-2.129	NDUFA12-like (NDUFA12L), mRNA.
gcat	0.47	-2.129	glycine C-acetyltransferase (2-amino-3-ketobutyrate coenzyme A ligase) (GCAT), nuclear gene encoding mitochondrial protein, mRNA.
cd2ap	0.47	-2.129	CD2-associated protein (CD2AP), mRNA.
gamt	0.47	-2.13	guanidinoacetate N-methyltransferase (GAMT), transcript variant 1, mRNA.
lysmd2	0.469	-2.13	LysM, putative peptidoglycan-binding, domain containing 2 (LYSMD2), mRNA.
polr2l	0.469	-2.13	polymerase (RNA) II (DNA directed) polypeptide L, 7.6kDa (POLR2L), mRNA.
tmem183b	0.469	-2.13	transmembrane protein 183B (TMEM183B), mRNA.
loc643319	0.469	-2.13	PREDICTED: similar to Transgelin-2 (LOC643319), mRNA.
hs.134650	0.469	-2.131	AGENCOURT_13567612 NIH_MGC_184 cDNA clone IMAGE:30326575 5, mRNA sequence
dlnb14	0.469	-2.131	coiled-coil domain containing 84 (CCDC84), mRNA.
fam96b	0.469	-2.131	family with sequence similarity 96, member B (FAM96B), mRNA.
kpnb1	0.469	-2.131	karyopherin (importin) beta 1 (KPNB1), mRNA.
atp6v1e2	0.469	-2.132	ATPase, H+ transporting, lysosomal 31kDa, V1 subunit E2 (ATP6V1E2), mRNA.
rab1b	0.469	-2.132	RAB1B, member RAS oncogene family (RAB1B), mRNA.
stard7	0.469	-2.132	START domain containing 7 (STARD7), transcript variant 1, mRNA.
plcxd1	0.469	-2.132	phosphatidylinositol-specific phospholipase C, X domain containing 1 (PLCXD1), mRNA.
praf2	0.469	-2.132	PRA1 domain family, member 2 (PRAF2), mRNA.
polr1e	0.469	-2.132	polymerase (RNA) I polypeptide E, 53kDa (POLR1E), mRNA.
ccs	0.469	-2.132	PREDICTED: copper chaperone for superoxide dismutase (CCS), mRNA.

loc148915	0.469	-2.133	PREDICTED: similar to Nonhistone chromosomal protein HMG-17 (High-mobility group nucleosome binding domain 2) (LOC148915), mRNA.
loc440567	0.469	-2.133	PREDICTED: similar to Ubiquinol-cytochrome c reductase complex 11 kDa protein, mitochondrial precursor (Mitochondrial hinge protein) (Cytochrome C1, nonheme 11 kDa protein) (Complex III subunit VIII), transcript variant 3 (LOC440567), mRNA.
slc39a3	0.469	-2.133	solute carrier family 39 (zinc transporter), member 3 (SLC39A3), transcript variant 2, mRNA.
fkrp	0.469	-2.133	fukutin related protein (FKRP), mRNA.
mta2	0.469	-2.133	metastasis associated 1 family, member 2 (MTA2), mRNA.
gpd1l	0.469	-2.133	glycerol-3-phosphate dehydrogenase 1-like (GPD1L), mRNA.
lactb	0.469	-2.133	lactamase, beta (LACTB), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
camk2n1	0.469	-2.134	calcium/calmodulin-dependent protein kinase II inhibitor 1 (CAMK2N1), mRNA.
itgb4bp	0.469	-2.134	integrin beta 4 binding protein (ITGB4BP), transcript variant 3, mRNA.
usp8	0.469	-2.134	ubiquitin specific peptidase 8 (USP8), mRNA.
11-Sep	0.469	-2.134	septin 11 (SEPT11), mRNA.
stambp	0.469	-2.134	STAM binding protein (STAMBP), transcript variant 1, mRNA.
mocos	0.469	-2.134	molybdenum cofactor sulfurase (MOCOS), mRNA.
dynlt1	0.469	-2.134	dynein, light chain, Tctex-type 1 (DYNLT1), mRNA.
slc44a1	0.468	-2.135	solute carrier family 44, member 1 (SLC44A1), transcript variant 1, mRNA.
gnpda1	0.468	-2.135	glucosamine-6-phosphate deaminase 1 (GNPDA1), mRNA.
loc654194	0.468	-2.135	PREDICTED: similar to ribosomal protein S27 (LOC654194), mRNA.
osbpl1a	0.468	-2.135	oxysterol binding protein-like 1A (OSBPL1A), transcript variant OSBPL1B, mRNA.
vps37a	0.468	-2.135	vacuolar protein sorting 37 homolog A (S. cerevisiae) (VPS37A), mRNA.
adh5	0.468	-2.135	alcohol dehydrogenase 5 (class III), chi polypeptide (ADH5), mRNA.
hist2h4a	0.468	-2.135	histone cluster 2, H4a (HIST2H4A), mRNA.
hoxc9	0.468	-2.135	homeobox C9 (HOXC9), mRNA.
mta3	0.468	-2.135	metastasis associated 1 family, member 3 (MTA3), mRNA.
kiaa1212	0.468	-2.135	coiled-coil domain containing 88A (CCDC88A), mRNA.
b3gat3	0.468	-2.136	beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I) (B3GAT3), mRNA.
armcx5	0.468	-2.136	armadillo repeat containing, X-linked 5 (ARMCX5), mRNA.
flj39827	0.468	-2.136	hypothetical protein FLJ39827 (FLJ39827), mRNA.
loc653506	0.468	-2.136	PREDICTED: similar to meteorin, glial cell differentiation regulator-like (LOC653506), mRNA.
polr3a	0.468	-2.136	polymerase (RNA) III (DNA directed) polypeptide A, 155kDa (POLR3A), mRNA.
btf3l4	0.468	-2.136	basic transcription factor 3-like 4 (BTF3L4), mRNA.
gng5	0.468	-2.136	guanine nucleotide binding protein (G protein), gamma 5 (GNG5), mRNA.
c3orf10	0.468	-2.136	chromosome 3 open reading frame 10 (C3orf10), mRNA.
loc642395	0.468	-2.136	PREDICTED: similar to paraspeckle protein 1, transcript variant 2 (LOC642395), mRNA.
mosc2	0.468	-2.137	MOCO sulphurase C-terminal domain containing 2 (MOSC2), mRNA.
prdm4	0.468	-2.137	PR domain containing 4 (PRDM4), mRNA.
tex264	0.468	-2.137	testis expressed 264 (TEX264), mRNA.
pgea1	0.468	-2.137	chibby homolog 1 (Drosophila) (CBY1), transcript variant 2, mRNA.
loc441408	0.468	-2.137	PREDICTED: hypothetical LOC441408, transcript variant 2 (LOC441408), mRNA.
tef	0.468	-2.137	thyrotrophic embryonic factor (TEF), mRNA.

mgc34830	0.468	-2.137	chromosome 11 open reading frame 69 (C11orf69), mRNA.
znf403	0.468	-2.138	zinc finger protein 403 (ZNF403), mRNA.
zfp91	0.468	-2.138	zinc finger protein 91 homolog (mouse) (ZFP91), transcript variant 2, mRNA.
dkfzp434b03	0.468	-2.138	DKFZP434B0335 protein (DKFZP434B0335), mRNA.
inpp5e	0.468	-2.138	inositol polyphosphate-5-phosphatase, 72 kDa (INPP5E), mRNA.
c14orf58	0.468	-2.138	feline leukemia virus subgroup C cellular receptor family, member 2 (FLVCR2), mRNA.
pdlim7	0.468	-2.138	PDZ and LIM domain 7 (enigma) (PDLIM7), transcript variant 4, mRNA.
lima1	0.468	-2.139	LIM domain and actin binding 1 (LIMA1), mRNA.
map2k1ip1	0.468	-2.139	mitogen-activated protein kinase kinase 1 interacting protein 1 (MAP2K1IP1), mRNA.
phkb	0.467	-2.139	phosphorylase kinase, beta (PHKB), transcript variant 2, mRNA.
trg20	0.467	-2.14	mediator complex subunit 10 (MED10), mRNA.
egln1	0.467	-2.14	egl nine homolog 1 (C. elegans) (EGLN1), mRNA.
dock11	0.467	-2.14	dedicator of cytokinesis 11 (DOCK11), mRNA.
polq	0.467	-2.14	polymerase (DNA directed), theta (POLQ), mRNA.
skp2	0.467	-2.14	S-phase kinase-associated protein 2 (p45) (SKP2), transcript variant 1, mRNA.
pde4b	0.467	-2.14	phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila) (PDE4B), transcript variant d, mRNA.
klhl24	0.467	-2.14	kelch-like 24 (Drosophila) (KLHL24), mRNA.
hs.294603	0.467	-2.14	cDNA FLJ12874 fis, clone NT2RP2003769
xpc	0.467	-2.14	xeroderma pigmentosum, complementation group C (XPC), mRNA.
loc647673	0.467	-2.141	PREDICTED: similar to Translationally-controlled tumor protein (TCTP) (p23) (Histamine-releasing factor) (HRF) (Fortilin) (LOC647673), mRNA.
il6	0.467	-2.141	interleukin 6 (interferon, beta 2) (IL6), mRNA.
egfr	0.467	-2.141	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian) (EGFR), transcript variant 3, mRNA.
psmd3	0.467	-2.141	proteasome (prosome, macropain) 26S subunit, non-ATPase, 3 (PSMD3), mRNA.
uckl1	0.467	-2.141	uridine-cytidine kinase 1-like 1 (UCKL1), mRNA.
ndufs8	0.467	-2.141	NADH dehydrogenase (ubiquinone) Fe-S protein 8, 23kDa (NADH-coenzyme Q reductase) (NDUFS8), mRNA.
rab15	0.467	-2.142	RAB15, member RAS onocogene family (RAB15), mRNA.
sidt2	0.467	-2.142	SID1 transmembrane family, member 2 (SIDT2), mRNA.
elmod2	0.467	-2.142	ELMO/CED-12 domain containing 2 (ELMOD2), mRNA.
hs.107418	0.467	-2.142	cDNA FLJ41853 fis, clone NT2RI3004161
mr1	0.467	-2.142	major histocompatibility complex, class I-related (MR1), mRNA.
znf562	0.467	-2.143	zinc finger protein 562 (ZNF562), mRNA.
loc402116	0.467	-2.143	PREDICTED: similar to plakophilin 4 (LOC402116), mRNA.
sh3kbp1	0.467	-2.143	SH3-domain kinase binding protein 1 (SH3KBP1), transcript variant 2, mRNA.
c13orf3	0.467	-2.143	chromosome 13 open reading frame 3 (C13orf3), mRNA.
bc002942	0.467	-2.143	transmembrane protein 112B (TMEM112B), mRNA.
ranbp5	0.467	-2.143	RAN binding protein 5 (RANBP5), mRNA.
cdca8	0.467	-2.143	cell division cycle associated 8 (CDCA8), mRNA.
tnks1bp1	0.467	-2.144	tankyrase 1 binding protein 1, 182kDa (TNKS1BP1), mRNA.
ddx41	0.466	-2.144	DEAD (Asp-Glu-Ala-Asp) box polypeptide 41 (DDX41), mRNA.

hs.13262	0.466	-2.144	PREDICTED: hypothetical gene supported by AY007155 (LOC439949), mRNA
c1orf77	0.466	-2.144	chromosome 1 open reading frame 77 (C1orf77), mRNA.
cart1	0.466	-2.144	cartilage paired-class homeoprotein 1 (CART1), mRNA.
hs.444290	0.466	-2.144	cDNA FLJ42786 fis, clone BRAWH3006761
c1orf53	0.466	-2.144	chromosome 1 open reading frame 53 (C1orf53), mRNA.
capn2	0.466	-2.144	calpain 2, (m/II) large subunit (CAPN2), mRNA.
dnaja2	0.466	-2.145	DnaJ (Hsp40) homolog, subfamily A, member 2 (DNAJA2), mRNA.
susd2	0.466	-2.145	sushi domain containing 2 (SUSD2), mRNA.
kiaa1815	0.466	-2.145	KIAA1815 (KIAA1815), mRNA.
sppl2a	0.466	-2.145	signal peptide peptidase-like 2A (SPPL2A), mRNA.
rnf126	0.466	-2.145	ring finger protein 126 (RNF126), mRNA.
nars2	0.466	-2.146	asparaginyl-tRNA synthetase 2, mitochondrial (putative) (NARS2), mRNA.
flj20699	0.466	-2.146	hypothetical protein FLJ20699 (FLJ20699), mRNA.
gspt2	0.466	-2.146	G1 to S phase transition 2 (GSPT2), mRNA.
fam62b	0.466	-2.146	family with sequence similarity 62 (C2 domain containing) member B (FAM62B), mRNA.
ap1m2	0.466	-2.147	adaptor-related protein complex 1, mu 2 subunit (AP1M2), mRNA.
loc440055	0.466	-2.147	PREDICTED: similar to ribosomal protein S12 (LOC440055), mRNA.
med9	0.466	-2.147	mediator of RNA polymerase II transcription, subunit 9 homolog (S. cerevisiae) (MED9), mRNA.
fam50b	0.466	-2.147	family with sequence similarity 50, member B (FAM50B), mRNA.
kif3b	0.466	-2.147	kinesin family member 3B (KIF3B), mRNA.
scand1	0.466	-2.148	SCAN domain containing 1 (SCAND1), transcript variant 2, mRNA.
ccdc23	0.466	-2.148	coiled-coil domain containing 23 (CCDC23), mRNA.
tpr	0.466	-2.148	translocated promoter region (to activated MET oncogene) (TPR), mRNA.
clasp1	0.466	-2.148	cytoplasmic linker associated protein 1 (CLASP1), mRNA.
hs.127310	0.465	-2.149	mRNA; cDNA DKFZp434C1613 (from clone DKFZp434C1613)
asahl	0.465	-2.149	N-acylsphingosine amidohydrolase (acid ceramidase)-like (ASAHL), mRNA.
dpysl3	0.465	-2.149	dihydropyrimidinase-like 3 (DPYSL3), mRNA.
psmc3ip	0.465	-2.149	PSMC3 interacting protein (PSMC3IP), transcript variant 2, mRNA.
mbnl1	0.465	-2.149	muscleblind-like (Drosophila) (MBNL1), transcript variant 3, mRNA.
map4	0.465	-2.149	microtubule-associated protein 4 (MAP4), transcript variant 2, mRNA.
kiaa1797	0.465	-2.149	KIAA1797 (KIAA1797), mRNA.
uap1	0.465	-2.149	UDP-N-acteylglucosamine pyrophosphorylase 1 (UAP1), mRNA.
ptk9l	0.465	-2.149	twinfilin, actin-binding protein, homolog 2 (Drosophila) (TWF2), mRNA.
lypla2	0.465	-2.15	lysophospholipase II (LYPLA2), mRNA.
f12	0.465	-2.15	coagulation factor XII (Hageman factor) (F12), mRNA.
loc389672	0.465	-2.15	PREDICTED: similar to 40S ribosomal protein SA (p40) (34/67 kDa laminin receptor) (Colon carcinoma laminin-binding protein) (NEM/1CHD4) (Multidrug resistance-associated protein MGr1-Ag) , transcript variant 3 (LOC389672), mRNA.
setdb1	0.465	-2.15	SET domain, bifurcated 1 (SETDB1), mRNA.
golph3	0.465	-2.15	golgi phosphoprotein 3 (coat-protein) (GOLPH3), mRNA.
c12orf32	0.465	-2.151	chromosome 12 open reading frame 32 (C12orf32), mRNA.
ncoa4	0.465	-2.151	nuclear receptor coactivator 4 (NCOA4), mRNA.

ebp	0.465	-2.151	emopamil binding protein (sterol isomerase) (EBP), mRNA.
hs.475334	0.465	-2.151	Human mRNA for KIAA0280 gene, partial cds
prickle2	0.465	-2.152	prickle-like 2 (Drosophila) (PRICKLE2), mRNA.
cxorf53	0.465	-2.152	BRCA1/BRCA2-containing complex, subunit 3 (BRCC3), transcript variant 1, mRNA.
prkar2a	0.465	-2.152	protein kinase, cAMP-dependent, regulatory, type II, alpha (PRKAR2A), mRNA.
pex14	0.465	-2.152	peroxisomal biogenesis factor 14 (PEX14), mRNA.
mrpl54	0.465	-2.152	mitochondrial ribosomal protein L54 (MRPL54), nuclear gene encoding mitochondrial protein, mRNA.
flnb	0.464	-2.153	filamin B, beta (actin binding protein 278) (FLNB), mRNA.
lrfn4	0.464	-2.153	leucine rich repeat and fibronectin type III domain containing 4 (LRFN4), mRNA.
hsd17b8	0.464	-2.153	hydroxysteroid (17-beta) dehydrogenase 8 (HSD17B8), mRNA.
cherp	0.464	-2.153	calcium homeostasis endoplasmic reticulum protein (CHERP), mRNA.
c16orf33	0.464	-2.154	chromosome 16 open reading frame 33 (C16orf33), mRNA.
eid3	0.464	-2.154	EP300 interacting inhibitor of differentiation 3 (EID3), mRNA.
gtf2f1	0.464	-2.154	general transcription factor IIF, polypeptide 1, 74kDa (GTF2F1), mRNA.
emp1	0.464	-2.154	epithelial membrane protein 1 (EMP1), mRNA.
c14orf104	0.464	-2.155	chromosome 14 open reading frame 104 (C14orf104), mRNA.
znfx1	0.464	-2.155	zinc finger, NFX1-type containing 1 (ZNFX1), mRNA.
hs.551538	0.464	-2.155	mRNA; cDNA DKFZp761E1721 (from clone DKFZp761E1721)
kiaa0460	0.464	-2.155	KIAA0460 (KIAA0460), mRNA.
mgc24665	0.464	-2.156	chromosome 16 open reading frame 75 (C16orf75), mRNA.
hs.430851	0.464	-2.156	ns68e03.y5 NCI_CGAP_Pr2 cDNA clone IMAGE:1188796, mRNA sequence
kiaa0133	0.464	-2.157	KIAA0133 (KIAA0133), mRNA.
gng4	0.464	-2.157	guanine nucleotide binding protein (G protein), gamma 4 (GNG4), mRNA.
mrps5	0.464	-2.157	mitochondrial ribosomal protein S5 (MRPS5), nuclear gene encoding mitochondrial protein, mRNA.
loc651436	0.464	-2.157	PREDICTED: similar to ribosomal protein L9 (LOC651436), mRNA.
bst1	0.464	-2.157	bone marrow stromal cell antigen 1 (BST1), mRNA.
icam3	0.464	-2.158	intercellular adhesion molecule 3 (ICAM3), mRNA.
slc35b4	0.463	-2.158	solute carrier family 35, member B4 (SLC35B4), mRNA.
gramd3	0.463	-2.158	GRAM domain containing 3 (GRAMD3), mRNA.
scap	0.463	-2.158	SREBP cleavage-activating protein (SCAP), mRNA.
nup153	0.463	-2.158	nucleoporin 153kDa (NUP153), mRNA.
hspa12a	0.463	-2.158	PREDICTED: heat shock 70kDa protein 12A, transcript variant 2 (HSPA12A), mRNA.
lsm2	0.463	-2.158	LSM2 homolog, U6 small nuclear RNA associated (S. cerevisiae) (LSM2), mRNA.
c13orf8	0.463	-2.158	chromosome 13 open reading frame 8 (C13orf8), mRNA.
c14orf149	0.463	-2.158	chromosome 14 open reading frame 149 (C14orf149), mRNA.
c14orf159	0.463	-2.158	chromosome 14 open reading frame 159 (C14orf159), mRNA.
c12orf44	0.463	-2.158	chromosome 12 open reading frame 44 (C12orf44), mRNA.
csnk1g1	0.463	-2.159	casein kinase 1, gamma 1 (CSNK1G1), transcript variant 2, mRNA.
sdccag1	0.463	-2.159	serologically defined colon cancer antigen 1 (SDCCAG1), mRNA.
c8orf40	0.463	-2.159	chromosome 8 open reading frame 40 (C8orf40), mRNA.
loc647037	0.463	-2.16	PREDICTED: similar to Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 (Rotamase Pin4) (PPIase Pin4) (Parvulin 14) (Par14) (Peptidyl-prolyl cis/trans isomerase EPVH)

			(hPar14) (LOC647037), mRNA.
gluld1	0.463	-2.16	glutamate-ammonia ligase (glutamine synthetase) domain containing 1 (GLULD1), mRNA.
itgb2	0.463	-2.16	integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit) (ITGB2), mRNA.
ang	0.463	-2.161	angiogenin, ribonuclease, RNase A family, 5 (ANG), mRNA.
loc645385	0.463	-2.161	PREDICTED: similar to heterogeneous nuclear ribonucleoprotein A1 (LOC645385), mRNA.
hs.175465	0.463	-2.161	cDNA FLJ36847 fis, clone ASTRO2013671
txnl5	0.463	-2.161	thioredoxin-like 5 (TXNL5), mRNA.
mov10	0.463	-2.161	Mov10, Moloney leukemia virus 10, homolog (mouse) (MOV10), mRNA.
rabl4	0.463	-2.161	RAB, member of RAS oncogene family-like 4 (RABL4), mRNA.
c17orf49	0.463	-2.161	chromosome 17 open reading frame 49 (C17orf49), mRNA.
pex16	0.463	-2.161	peroxisomal biogenesis factor 16 (PEX16), transcript variant 1, mRNA.
loc400464	0.463	-2.161	similar to FLJ43276 protein (LOC400464), mRNA.
pmpca	0.463	-2.161	peptidase (mitochondrial processing) alpha (PMPCA), nuclear gene encoding mitochondrial protein, mRNA.
ywhaq	0.463	-2.161	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide (YWHAQ), mRNA.
rfc5	0.463	-2.161	replication factor C (activator 1) 5, 36.5kDa (RFC5), transcript variant 2, mRNA.
loc642934	0.463	-2.161	PREDICTED: hypothetical protein LOC642934 (LOC642934), mRNA.
zfpl1	0.463	-2.162	zinc finger protein-like 1 (ZFPL1), mRNA.
asnsd1	0.463	-2.162	asparagine synthetase domain containing 1 (ASNSD1), mRNA.
pcm1	0.462	-2.162	pericentriolar material 1 (PCM1), mRNA.
prmt7	0.462	-2.162	protein arginine methyltransferase 7 (PRMT7), mRNA.
gfm1	0.462	-2.162	G elongation factor, mitochondrial 1 (GFM1), nuclear gene encoding mitochondrial protein, mRNA.
cdc5l	0.462	-2.162	CDC5 cell division cycle 5-like (S. pombe) (CDC5L), mRNA.
anapc5	0.462	-2.163	anaphase promoting complex subunit 5 (ANAPC5), mRNA.
acaa1	0.462	-2.163	acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase) (ACAA1), nuclear gene encoding mitochondrial protein, mRNA.
prpf4	0.462	-2.163	PRP4 pre-mRNA processing factor 4 homolog (yeast) (PRPF4), mRNA.
mtx1	0.462	-2.163	metaxin 1 (MTX1), transcript variant 1, mRNA.
ifnar1	0.462	-2.163	interferon (alpha, beta and omega) receptor 1 (IFNAR1), mRNA.
prpsap2	0.462	-2.163	phosphoribosyl pyrophosphate synthetase-associated protein 2 (PRPSAP2), mRNA.
tnrc6b	0.462	-2.164	trinucleotide repeat containing 6B (TNRC6B), transcript variant 1, mRNA.
srf	0.462	-2.164	serum response factor (c-fos serum response element-binding transcription factor) (SRF), mRNA.
cabin1	0.462	-2.164	calcineurin binding protein 1 (CABIN1), mRNA.
phldb1	0.462	-2.164	pleckstrin homology-like domain, family B, member 1 (PHLDB1), mRNA.
reep5	0.462	-2.164	receptor accessory protein 5 (REEP5), mRNA.
c10orf22	0.462	-2.165	chromosome 10 open reading frame 22 (C10orf22), mRNA.
kiaa1287	0.462	-2.165	integrator complex subunit 2 (INTS2), mRNA.
nktr	0.462	-2.165	natural killer-tumor recognition sequence (NKTR), mRNA.
gcs1	0.462	-2.165	glucosidase I (GCS1), mRNA.
comt	0.462	-2.165	catechol-O-methyltransferase (COMT), transcript variant MB-COMT, mRNA.
dkfzp686o24	0.462	-2.165	hypothetical protein DKFZp686O24166 (DKFZp686O24166), mRNA.
znf598	0.462	-2.165	zinc finger protein 598 (ZNF598), mRNA.

loc400879	0.462	-2.166	PREDICTED: hypothetical LOC400879, transcript variant 1 (LOC400879), mRNA.
raver2	0.462	-2.166	ribonucleoprotein, PTB-binding 2 (RAVER2), mRNA.
bat1	0.462	-2.166	HLA-B associated transcript 1 (BAT1), transcript variant 2, mRNA.
loc144363	0.462	-2.166	hypothetical protein LOC144363 (LOC144363), mRNA.
b3gntl1	0.462	-2.166	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase-like 1 (B3GNTL1), mRNA.
fbxo21	0.462	-2.166	F-box protein 21 (FBXO21), transcript variant 2, mRNA.
rere	0.462	-2.166	arginine-glutamic acid dipeptide (RE) repeats (RERE), transcript variant 1, mRNA.
b3gtl	0.462	-2.167	beta 3-glycosyltransferase-like (B3GTL), mRNA.
ndfip1	0.461	-2.167	Nedd4 family interacting protein 1 (NDFIP1), mRNA.
pthlh	0.461	-2.167	parathyroid hormone-like hormone (PTHLH), transcript variant 1, mRNA.
pmm2	0.461	-2.167	phosphomannomutase 2 (PMM2), mRNA.
rgs11	0.461	-2.167	regulator of G-protein signalling 11 (RGS11), transcript variant 1, mRNA.
cep70	0.461	-2.167	centrosomal protein 70kDa (CEP70), mRNA.
nmt1	0.461	-2.168	N-myristoyltransferase 1 (NMT1), mRNA.
dad1	0.461	-2.168	defender against cell death 1 (DAD1), mRNA.
znf689	0.461	-2.168	zinc finger protein 689 (ZNF689), mRNA.
rapsn	0.461	-2.168	receptor-associated protein of the synapse (RAPSN), transcript variant 1, mRNA.
hs.4892	0.461	-2.168	clone 24841 mRNA sequence
triap1	0.461	-2.168	TP53 regulated inhibitor of apoptosis 1 (TRIAP1), mRNA.
maf1	0.461	-2.169	MAF1 homolog (S. cerevisiae) (MAF1), mRNA.
samd4b	0.461	-2.169	sterile alpha motif domain containing 4B (SAMD4B), mRNA.
mef2d	0.461	-2.169	myocyte enhancer factor 2D (MEF2D), mRNA.
idh3b	0.461	-2.17	isocitrate dehydrogenase 3 (NAD+) beta (IDH3B), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
loc642829	0.461	-2.17	PREDICTED: prostate-specific P704P (P704P), mRNA.
itga11	0.461	-2.17	integrin, alpha 11 (ITGA11), transcript variant 2, mRNA.
loc653890	0.461	-2.17	PREDICTED: similar to serine/threonine/tyrosine interacting protein (LOC653890), mRNA.
foxd4	0.461	-2.17	forkhead box D4 (FOXD4), mRNA.
supt16h	0.461	-2.17	suppressor of Ty 16 homolog (S. cerevisiae) (SUPT16H), mRNA.
plxna3	0.461	-2.17	plexin A3 (PLXNA3), mRNA.
tgfb1i1	0.461	-2.17	transforming growth factor beta 1 induced transcript 1 (TGFB1I1), transcript variant 2, mRNA.
gk2	0.461	-2.17	glycerol kinase 2 (GK2), mRNA.
cldn12	0.461	-2.171	claudin 12 (CLDN12), mRNA.
ndufs5	0.461	-2.171	NADH dehydrogenase (ubiquinone) Fe-S protein 5, 15kDa (NADH-coenzyme Q reductase) (NDUFS5), mRNA.
cd99	0.461	-2.171	CD99 molecule (CD99), mRNA.
mrpl12	0.46	-2.172	mitochondrial ribosomal protein L12 (MRPL12), nuclear gene encoding mitochondrial protein, mRNA.
hs1bp3	0.46	-2.172	HCLS1 binding protein 3 (HS1BP3), mRNA.
c16orf7	0.46	-2.172	chromosome 16 open reading frame 7 (C16orf7), mRNA.
rab5b	0.46	-2.172	RAB5B, member RAS oncogene family (RAB5B), mRNA.
flj39378	0.46	-2.172	hypothetical protein FLJ39378 (FLJ39378), mRNA.
hs.355933	0.46	-2.172	cDNA FLJ41921 fis, clone PERIC2002766
kiaa1698	0.46	-2.173	integrator complex subunit 5 (INTS5), mRNA.

sfrs8	0.46	-2.173	splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila) (SFRS8), mRNA.
nme6	0.46	-2.174	non-metastatic cells 6, protein expressed in (nucleoside-diphosphate kinase) (NME6), mRNA.
dcamkl2	0.46	-2.174	doublecortin and CaM kinase-like 2 (DCAMKL2), mRNA.
ell2	0.46	-2.175	elongation factor, RNA polymerase II, 2 (ELL2), mRNA.
slc39a11	0.46	-2.175	solute carrier family 39 (metal ion transporter), member 11 (SLC39A11), mRNA.
sec22l1	0.46	-2.175	SEC22 vesicle trafficking protein-like 1 (S. cerevisiae) (SEC22L1), mRNA.
ormdl3	0.46	-2.175	ORM1-like 3 (S. cerevisiae) (ORMDL3), mRNA.
aes	0.46	-2.175	amino-terminal enhancer of split (AES), transcript variant 1, mRNA.
suclg1	0.459	-2.177	succinate-CoA ligase, GDP-forming, alpha subunit (SUCLG1), mRNA.
brd9	0.459	-2.177	bromodomain containing 9 (BRD9), transcript variant 1, mRNA.
magee1	0.459	-2.177	melanoma antigen family E, 1 (MAGEE1), mRNA.
loc650472	0.459	-2.177	PREDICTED: similar to 60S ribosomal protein L23a (LOC650472), mRNA.
znf524	0.459	-2.177	zinc finger protein 524 (ZNF524), mRNA.
ctdp1	0.459	-2.177	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1 (CTDP1), transcript variant FCP1a, mRNA.
rnpep	0.459	-2.178	arginyl aminopeptidase (aminopeptidase B) (RNPEP), mRNA.
hs.547712	0.459	-2.178	mRNA full length insert cDNA clone EUROIMAGE 1090207
cpt2	0.459	-2.178	carnitine palmitoyltransferase II (CPT2), nuclear gene encoding mitochondrial protein, mRNA.
mif	0.459	-2.178	macrophage migration inhibitory factor (glycosylation-inhibiting factor) (MIF), mRNA.
snap29	0.459	-2.178	synaptosomal-associated protein, 29kDa (SNAP29), mRNA.
malt1	0.459	-2.178	mucosa associated lymphoid tissue lymphoma translocation gene 1 (MALT1), transcript variant 1, mRNA.
eif3s1	0.459	-2.178	eukaryotic translation initiation factor 3, subunit 1 alpha, 35kDa (EIF3S1), mRNA.
snx6	0.459	-2.178	sorting nexin 6 (SNX6), transcript variant 1, mRNA.
ppp1r12c	0.459	-2.179	protein phosphatase 1, regulatory (inhibitor) subunit 12C (PPP1R12C), mRNA.
josd1	0.459	-2.179	Josephin domain containing 1 (JOSD1), mRNA.
pinx1	0.459	-2.179	PIN2-interacting protein 1 (PINX1), mRNA.
atp5g1	0.459	-2.179	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit C1 (subunit 9) (ATP5G1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
shb	0.459	-2.179	Src homology 2 domain containing adaptor protein B (SHB), mRNA.
eya4	0.459	-2.179	eyes absent homolog 4 (Drosophila) (EYA4), transcript variant 4, mRNA.
edn1	0.459	-2.179	endothelin 1 (EDN1), mRNA.
hivep1	0.459	-2.179	human immunodeficiency virus type I enhancer binding protein 1 (HIVEP1), mRNA.
vrk2	0.459	-2.18	vaccinia related kinase 2 (VRK2), mRNA.
churc1	0.459	-2.18	churchill domain containing 1 (CHURC1), mRNA.
fam57a	0.459	-2.18	family with sequence similarity 57, member A (FAM57A), mRNA.
tmem99	0.459	-2.18	transmembrane protein 99 (TMEM99), mRNA.
zfp106	0.459	-2.18	zinc finger protein 106 homolog (mouse) (ZFP106), mRNA.
erf	0.459	-2.18	Ets2 repressor factor (ERF), mRNA.
rhobtb3	0.459	-2.181	Rho-related BTB domain containing 3 (RHOBTB3), mRNA.
gars	0.459	-2.181	glycyl-tRNA synthetase (GARS), mRNA.
hkdc1	0.458	-2.181	hexokinase domain containing 1 (HKDC1), mRNA.
elovl5	0.458	-2.181	ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) (ELOVL5), mRNA.

ik	0.458	-2.181	IK cytokine, down-regulator of HLA II (IK), mRNA.
flj46072	0.458	-2.182	FLJ46072 protein (FLJ46072), mRNA.
pfdn1	0.458	-2.183	prefoldin 1 (PFDN1), mRNA.
pacsin3	0.458	-2.183	protein kinase C and casein kinase substrate in neurons 3 (PACSIN3), mRNA.
loc654260	0.458	-2.183	PREDICTED: similar to beta-tubulin cofactor D isoform 1 (LOC654260), mRNA.
flj20628	0.458	-2.183	hypothetical protein FLJ20628 (FLJ20628), mRNA.
dpy19l4	0.458	-2.183	dpy-19-like 4 (C. elegans) (DPY19L4), mRNA.
musk	0.458	-2.183	muscle, skeletal, receptor tyrosine kinase (MUSK), mRNA.
rab3il1	0.458	-2.183	RAB3A interacting protein (rabin3)-like 1 (RAB3IL1), mRNA.
gli2	0.458	-2.183	GLI-Kruppel family member GLI2 (GLI2), transcript variant 4, mRNA.
cspp1	0.458	-2.183	centrosome and spindle pole associated protein 1 (CSPP1), mRNA.
hs.556018	0.458	-2.183	mRNA; cDNA DKFZp779F0411 (from clone DKFZp779F0411)
orc5l	0.458	-2.184	origin recognition complex, subunit 5-like (yeast) (ORC5L), transcript variant 2, mRNA.
lmtk3	0.458	-2.184	PREDICTED: lemur tyrosine kinase 3 (LMTK3), mRNA.
fam122b	0.458	-2.184	family with sequence similarity 122B (FAM122B), mRNA.
lctl	0.458	-2.184	lactase-like (LCTL), mRNA.
wdr55	0.458	-2.184	WD repeat domain 55 (WDR55), mRNA.
rabggtb	0.458	-2.184	Rab geranylgeranyltransferase, beta subunit (RABGGTB), mRNA.
ccdc50	0.458	-2.184	coiled-coil domain containing 50 (CCDC50), transcript variant 2, mRNA.
ttyh3	0.458	-2.184	tweety homolog 3 (Drosophila) (TTYH3), mRNA.
dusp12	0.458	-2.184	dual specificity phosphatase 12 (DUSP12), mRNA.
tmem14b	0.458	-2.184	transmembrane protein 14B (TMEM14B), mRNA.
srprb	0.458	-2.184	signal recognition particle receptor, B subunit (SRPRB), mRNA.
rai14	0.458	-2.184	retinoic acid induced 14 (RAI14), mRNA.
kpna1	0.458	-2.185	karyopherin alpha 1 (importin alpha 5) (KPNA1), mRNA.
htr2a	0.458	-2.185	5-hydroxytryptamine (serotonin) receptor 2A (HTR2A), mRNA.
c14orf135	0.458	-2.185	chromosome 14 open reading frame 135 (C14orf135), mRNA.
fam86a	0.458	-2.185	family with sequence similarity 86, member A (FAM86A), transcript variant 2, mRNA.
hs.579631	0.458	-2.185	AGENCOURT_10229596 NIH_MGC_141 cDNA clone IMAGE:6563923 5, mRNA sequence
cpt1c	0.458	-2.185	carnitine palmitoyltransferase 1C (CPT1C), mRNA.
zgpat	0.458	-2.185	zinc finger, CCCH-type with G patch domain (ZGPAT), transcript variant 2, mRNA.
ppic	0.458	-2.185	peptidylprolyl isomerase C (cyclophilin C) (PPIC), mRNA.
pdia6	0.458	-2.185	protein disulfide isomerase family A, member 6 (PDIA6), mRNA.
rdx	0.458	-2.186	radixin (RDX), mRNA.
slc38a2	0.458	-2.186	solute carrier family 38, member 2 (SLC38A2), mRNA.
shkbp1	0.457	-2.186	SH3KBP1 binding protein 1 (SHKBP1), mRNA.
flj14640	0.457	-2.186	coiled-coil domain containing 123 (CCDC123), mRNA.
pold3	0.457	-2.186	polymerase (DNA-directed), delta 3, accessory subunit (POLD3), mRNA.
smarca5	0.457	-2.186	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 (SMARCA5), mRNA.
ctsl	0.457	-2.187	cathepsin L (CTSL), transcript variant 1, mRNA.
loc283951	0.457	-2.187	hypothetical protein LOC283951 (LOC283951), mRNA.

acad11	0.457	-2.187	acyl-Coenzyme A dehydrogenase family, member 11 (ACAD11), mRNA.
serinc3	0.457	-2.187	serine incorporator 3 (SERINC3), transcript variant 2, mRNA.
coq9	0.457	-2.187	coenzyme Q9 homolog (S. cerevisiae) (COQ9), mRNA.
tomm34	0.457	-2.187	translocase of outer mitochondrial membrane 34 (TOMM34), nuclear gene encoding mitochondrial protein, mRNA
nipa1	0.457	-2.188	non imprinted in Prader-Willi/Angelman syndrome 1 (NIPA1), mRNA.
ifi35	0.457	-2.188	interferon-induced protein 35 (IFI35), mRNA.
dnajc12	0.457	-2.188	DnaJ (Hsp40) homolog, subfamily C, member 12 (DNAJC12), transcript variant 2, mRNA.
c9orf142	0.457	-2.188	chromosome 9 open reading frame 142 (C9orf142), mRNA.
mrpl9	0.457	-2.189	mitochondrial ribosomal protein L9 (MRPL9), nuclear gene encoding mitochondrial protein, mRNA.
txnrd2	0.457	-2.189	thioredoxin reductase 2 (TXNRD2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
psmd9	0.457	-2.189	proteasome (prosome, macropain) 26S subunit, non-ATPase, 9 (PSMD9), mRNA.
c16orf9	0.457	-2.19	integrin alpha FG-GAP repeat containing 3 (ITFG3), mRNA.
coasy	0.457	-2.19	Coenzyme A synthase (COASY), mRNA.
c3orf64	0.457	-2.19	chromosome 3 open reading frame 64 (C3orf64), mRNA.
plekhc1	0.456	-2.191	pleckstrin homology domain containing, family C (with FERM domain) member 1 (PLEKHC1), mRNA.
ocrl	0.456	-2.191	oculocerebrorenal syndrome of Lowe (OCRL), transcript variant b, mRNA.
centd3	0.456	-2.191	centaurin, delta 3 (CENTD3), mRNA.
tmem39b	0.456	-2.191	transmembrane protein 39B (TMEM39B), mRNA.
uba2	0.456	-2.192	SUMO1 activating enzyme subunit 2 (SAE2), mRNA.
dyrk4	0.456	-2.192	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 4 (DYRK4), mRNA.
otud5	0.456	-2.192	OTU domain containing 5 (OTUD5), mRNA.
xab2	0.456	-2.192	XPA binding protein 2 (XAB2), mRNA.
mrps6	0.456	-2.192	mitochondrial ribosomal protein S6 (MRPS6), nuclear gene encoding mitochondrial protein, mRNA.
scg5	0.456	-2.193	secretogranin V (7B2 protein) (SCG5), mRNA.
cybrd1	0.456	-2.193	cytochrome b reductase 1 (CYBRD1), mRNA.
mrps33	0.456	-2.193	mitochondrial ribosomal protein S33 (MRPS33), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
mrps18a	0.456	-2.193	mitochondrial ribosomal protein S18A (MRPS18A), nuclear gene encoding mitochondrial protein, mRNA.
kiaa0913	0.456	-2.193	KIAA0913 (KIAA0913), mRNA.
loc644743	0.456	-2.193	PREDICTED: hypothetical protein LOC644743 (LOC644743), mRNA.
loc642019	0.456	-2.193	PREDICTED: similar to nucleolar protein 5A (LOC642019), mRNA.
tmem103	0.456	-2.193	transmembrane protein 103 (TMEM103), transcript variant 1, mRNA.
suz12	0.456	-2.194	suppressor of zeste 12 homolog (Drosophila) (SUZ12), mRNA.
pgcp	0.456	-2.194	plasma glutamate carboxypeptidase (PGCP), mRNA.
homer3	0.456	-2.194	homer homolog 3 (Drosophila) (HOMER3), mRNA.
ninj1	0.456	-2.194	ninjurin 1 (NINJ1), mRNA.
wdr45	0.456	-2.195	WD repeat domain 45 (WDR45), transcript variant 2, mRNA.
mat2a	0.456	-2.195	methionine adenosyltransferase II, alpha (MAT2A), mRNA.
cpvl	0.456	-2.195	carboxypeptidase, vitellogenic-like (CPVL), transcript variant 1, mRNA.
ufc1	0.456	-2.195	ubiquitin-fold modifier conjugating enzyme 1 (UFC1), mRNA.
stx6	0.456	-2.195	syntaxin 6 (STX6), mRNA.
zbp1	0.456	-2.195	Z-DNA binding protein 1 (ZBP1), mRNA.

prcc	0.456	-2.195	papillary renal cell carcinoma (translocation-associated) (PRCC), transcript variant 2, mRNA.
zcsl2	0.455	-2.195	DPH3, KTI11 homolog (S. cerevisiae) (DPH3), transcript variant 1, mRNA.
ldoc1l	0.455	-2.196	leucine zipper, down-regulated in cancer 1-like (LDOC1L), mRNA.
zdhhc17	0.455	-2.196	zinc finger, DHHC-type containing 17 (ZDHHC17), mRNA.
cxx1	0.455	-2.196	CAAX box 1 (CXX1), mRNA.
ripk1	0.455	-2.197	receptor (TNFRSF)-interacting serine-threonine kinase 1 (RIPK1), mRNA.
loc90639	0.455	-2.197	hypothetical protein LOC90639 (LOC90639), mRNA.
cep55	0.455	-2.197	centrosomal protein 55kDa (CEP55), mRNA.
tob1	0.455	-2.198	transducer of ERBB2, 1 (TOB1), mRNA.
cbx1	0.455	-2.198	chromobox homolog 1 (HP1 beta homolog Drosophila) (CBX1), mRNA.
adrm1	0.455	-2.198	adhesion regulating molecule 1 (ADRM1), transcript variant 1, mRNA.
sfrs16	0.455	-2.198	splicing factor, arginine/serine-rich 16 (SFRS16), mRNA.
tubb	0.455	-2.198	tubulin, beta (TUBB), mRNA.
abhd5	0.455	-2.199	abhydrolase domain containing 5 (ABHD5), mRNA.
sertad2	0.455	-2.199	SERTA domain containing 2 (SERTAD2), mRNA.
gnpat	0.455	-2.199	glyceronephosphate O-acyltransferase (GNPAT), mRNA.
loc347292	0.455	-2.199	PREDICTED: similar to ribosomal protein L36 (LOC347292), mRNA.
mrps16	0.455	-2.199	mitochondrial ribosomal protein S16 (MRPS16), nuclear gene encoding mitochondrial protein, mRNA.
mgc4562	0.455	-2.199	DIS3 mitotic control homolog (S. cerevisiae)-like (DIS3L), mRNA.
sipa1l2	0.455	-2.199	signal-induced proliferation-associated 1 like 2 (SIPA1L2), mRNA.
bysl	0.455	-2.199	bystin-like (BYSL), mRNA.
cdkn3	0.455	-2.199	cyclin-dependent kinase inhibitor 3 (CDK2-associated dual specificity phosphatase) (CDKN3), mRNA.
josd2	0.455	-2.2	Josephin domain containing 2 (JOSD2), mRNA.
tnpo2	0.455	-2.2	transportin 2 (importin 3, karyopherin beta 2b) (TNPO2), mRNA.
cbara1	0.455	-2.2	calcium binding atopy-related autoantigen 1 (CBARA1), mRNA.
bcl7b	0.455	-2.2	B-cell CLL/lymphoma 7B (BCL7B), transcript variant 2, mRNA.
loc128439	0.455	-2.2	hypothetical gene LOC128439 (LOC128439), mRNA.
bscl2	0.454	-2.2	Bernardinelli-Seip congenital lipodystrophy 2 (seipin) (BSCL2), mRNA.
flj20534	0.454	-2.201	chromosome 4 open reading frame 27 (C4orf27), mRNA.
c6orf141	0.454	-2.201	chromosome 6 open reading frame 141 (C6orf141), mRNA.
usp13	0.454	-2.201	ubiquitin specific peptidase 13 (isopeptidase T-3) (USP13), mRNA.
grb2	0.454	-2.201	growth factor receptor-bound protein 2 (GRB2), transcript variant 1, mRNA.
flj10006	0.454	-2.201	IWS1 homolog (S. cerevisiae) (IWS1), mRNA.
serf1b	0.454	-2.201	small EDRK-rich factor 1B (centromeric) (SERF1B), mRNA.
insl4	0.454	-2.201	insulin-like 4 (placenta) (INSL4), mRNA.
gtf2ip1	0.454	-2.201	general transcription factor II, i, pseudogene 1 (GTF2IP1) on chromosome 7.
c8orf32	0.454	-2.202	chromosome 8 open reading frame 32 (C8orf32), mRNA.
ift172	0.454	-2.202	intraflagellar transport 172 homolog (Chlamydomonas) (IFT172), mRNA.
rbm39	0.454	-2.202	RNA binding motif protein 39 (RBM39), transcript variant 1, mRNA.
rab21	0.454	-2.202	RAB21, member RAS oncogene family (RAB21), mRNA.
cox7a2	0.454	-2.202	cytochrome c oxidase subunit VIIa polypeptide 2 (liver) (COX7A2), mRNA.
fbxo5	0.454	-2.203	F-box protein 5 (FBXO5), mRNA.

eif5a2	0.454	-2.203	eukaryotic translation initiation factor 5A2 (EIF5A2), mRNA.
impa1	0.454	-2.203	inositol(myo)-1(or 4)-monophosphatase 1 (IMPA1), mRNA.
scamp5	0.454	-2.203	secretory carrier membrane protein 5 (SCAMP5), mRNA.
tacc2	0.454	-2.204	transforming, acidic coiled-coil containing protein 2 (TACC2), transcript variant 2, mRNA.
hs.279842	0.454	-2.204	HSPC157 protein, mRNA (cDNA clone IMAGE:6672800), partial cds
sh3bp5l	0.454	-2.204	SH3-binding domain protein 5-like (SH3BP5L), mRNA.
c9orf23	0.454	-2.204	chromosome 9 open reading frame 23 (C9orf23), transcript variant 1, mRNA.
nars	0.454	-2.204	asparaginyl-tRNA synthetase (NARS), mRNA.
magmas	0.454	-2.204	mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction (Magmas), nuclear gene encoding mitochondrial protein, mRNA.
sult2b1	0.454	-2.204	sulfotransferase family, cytosolic, 2B, member 1 (SULT2B1), transcript variant 1, mRNA.
gja1	0.453	-2.205	gap junction protein, alpha 1, 43kDa (connexin 43) (GJA1), mRNA.
galnt4	0.453	-2.206	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4) (GALNT4), mRNA.
wdr70	0.453	-2.206	WD repeat domain 70 (WDR70), mRNA.
efha1	0.453	-2.206	EF-hand domain family, member A1 (EFHA1), mRNA.
loc389137	0.453	-2.206	PREDICTED: similar to SVH protein (LOC389137), mRNA.
loc654121	0.453	-2.206	PREDICTED: similar to C49H3.3 (LOC654121), mRNA.
drg1	0.453	-2.207	developmentally regulated GTP binding protein 1 (DRG1), mRNA.
cops3	0.453	-2.207	COP9 constitutive photomorphogenic homolog subunit 3 (Arabidopsis) (COPS3), mRNA.
ddx24	0.453	-2.208	DEAD (Asp-Glu-Ala-Asp) box polypeptide 24 (DDX24), mRNA.
yif1a	0.453	-2.208	Yip1 interacting factor homolog A (S. cerevisiae) (YIF1A), mRNA.
tyms	0.453	-2.208	thymidylate synthetase (TYMS), mRNA.
hdhd2	0.453	-2.208	haloacid dehalogenase-like hydrolase domain containing 2 (HDHD2), mRNA.
ndufa10	0.453	-2.208	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10, 42kDa (NDUFA10), nuclear gene encoding mitochondrial protein, mRNA.
calm1	0.453	-2.208	calmodulin 1 (phosphorylase kinase, delta) (CALM1), mRNA.
med28	0.453	-2.208	mediator complex subunit 28 (MED28), mRNA.
c10orf33	0.452	-2.21	chromosome 10 open reading frame 33 (C10orf33), mRNA.
cat	0.452	-2.21	catalase (CAT), mRNA.
anxa5	0.452	-2.21	annexin A5 (ANXA5), mRNA.
aip	0.452	-2.21	aryl hydrocarbon receptor interacting protein (AIP), mRNA.
psme3	0.452	-2.211	proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki) (PSME3), transcript variant 1, mRNA.
tns3	0.452	-2.212	tensin 3 (TNS3), mRNA.
mgc10911	0.452	-2.213	chromosome 7 open reading frame 48 (C7orf48), mRNA.
polrmt	0.452	-2.213	polymerase (RNA) mitochondrial (DNA directed) (POLRMT), nuclear gene encoding mitochondrial protein, mRNA.
slc6a15	0.452	-2.213	solute carrier family 6, member 15 (SLC6A15), transcript variant 2, mRNA.
kctd3	0.452	-2.213	potassium channel tetramerisation domain containing 3 (KCTD3), mRNA.
sec23ip	0.452	-2.214	SEC23 interacting protein (SEC23IP), mRNA.
cbr1	0.452	-2.214	carbonyl reductase 1 (CBR1), mRNA.
ankrd12	0.452	-2.214	ankyrin repeat domain 12 (ANKRD12), mRNA.
clec11a	0.452	-2.214	C-type lectin domain family 11, member A (CLEC11A), mRNA.

cap2	0.452	-2.214	CAP, adenylate cyclase-associated protein, 2 (yeast) (CAP2), mRNA.
ttl	0.452	-2.215	tubulin tyrosine ligase (TTL), mRNA.
thada	0.452	-2.215	thyroid adenoma associated (THADA), transcript variant 1, mRNA.
loc284988	0.452	-2.215	PREDICTED: hypothetical LOC284988 (LOC284988), mRNA.
armc1	0.452	-2.215	armadillo repeat containing 1 (ARMC1), mRNA.
b4galt5	0.451	-2.215	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5 (B4GALT5), mRNA.
hs.522924	0.451	-2.215	cDNA FLJ43160 fis, clone FCBBF2000199
mrps34	0.451	-2.215	mitochondrial ribosomal protein S34 (MRPS34), nuclear gene encoding mitochondrial protein, mRNA.
flj38717	0.451	-2.215	FLJ38717 protein (FLJ38717), mRNA.
brpf3	0.451	-2.215	bromodomain and PHD finger containing, 3 (BRPF3), mRNA.
c16orf45	0.451	-2.216	chromosome 16 open reading frame 45 (C16orf45), mRNA.
loc653226	0.451	-2.216	PREDICTED: similar to Signal recognition particle 9 kDa protein (SRP9) (LOC653226), mRNA.
rbm13	0.451	-2.217	RNA binding motif protein 13 (RBM13), mRNA.
narfl	0.451	-2.217	nuclear prelamin A recognition factor-like (NARFL), mRNA.
lpxn	0.451	-2.217	leupaxin (LPXN), mRNA.
cpe	0.451	-2.217	carboxypeptidase E (CPE), mRNA.
stxbp2	0.451	-2.217	syntaxin binding protein 2 (STXBP2), mRNA.
galnt14	0.451	-2.217	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 14 (GalNAc-T14) (GALNT14), mRNA.
taf1b	0.451	-2.218	TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63kDa (TAF1B), mRNA.
emp3	0.451	-2.218	epithelial membrane protein 3 (EMP3), mRNA.
aldh2	0.451	-2.219	aldehyde dehydrogenase 2 family (mitochondrial) (ALDH2), nuclear gene encoding mitochondrial protein, mRNA.
wbp5	0.451	-2.219	WW domain binding protein 5 (WBP5), transcript variant 4, mRNA.
srp9	0.451	-2.219	signal recognition particle 9kDa (SRP9), mRNA.
klf11	0.451	-2.219	PREDICTED: Kruppel-like factor 11 (KLF11), mRNA.
klf13	0.451	-2.219	Kruppel-like factor 13 (KLF13), mRNA.
clpp	0.451	-2.22	ClpP caseinolytic peptidase, ATP-dependent, proteolytic subunit homolog (E. coli) (CLPP), nuclear gene encoding mitochondrial protein, mRNA.
vps36	0.45	-2.22	vacuolar protein sorting 36 homolog (S. cerevisiae) (VPS36), mRNA.
flj12886	0.45	-2.221	hypothetical protein FLJ12886 (FLJ12886), mRNA.
pet112l	0.45	-2.221	PET112-like (yeast) (PET112L), mRNA.
bcat1	0.45	-2.221	branched chain aminotransferase 1, cytosolic (BCAT1), mRNA.
gys1	0.45	-2.221	glycogen synthase 1 (muscle) (GYS1), mRNA.
c20orf111	0.45	-2.221	chromosome 20 open reading frame 111 (C20orf111), mRNA.
c1orf86	0.45	-2.221	chromosome 1 open reading frame 86 (C1orf86), mRNA.
pomzp3	0.45	-2.221	POM (POM121 homolog, rat) and ZP3 fusion (POMZP3), transcript variant 1, mRNA.
fam100a	0.45	-2.221	family with sequence similarity 100, member A (FAM100A), mRNA.
akr7a3	0.45	-2.222	aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase) (AKR7A3), mRNA.
loc51255	0.45	-2.222	ring finger protein 181 (RNF181), mRNA.
cited4	0.45	-2.222	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 4 (CITED4), mRNA.
c6orf26	0.45	-2.222	chromosome 6 open reading frame 26 (C6orf26), mRNA.
tmem50b	0.45	-2.222	transmembrane protein 50B (TMEM50B), mRNA.

kiaa0674	0.45	-2.222	PREDICTED: KIAA0674 (KIAA0674), mRNA.
rps27a	0.45	-2.222	ribosomal protein S27a (RPS27A), mRNA.
me1	0.45	-2.222	malic enzyme 1, NADP(+)-dependent, cytosolic (ME1), mRNA.
slc39a6	0.45	-2.222	solute carrier family 39 (zinc transporter), member 6 (SLC39A6), mRNA.
rpl5	0.45	-2.223	ribosomal protein L5 (RPL5), mRNA.
brd2	0.45	-2.223	bromodomain containing 2 (BRD2), mRNA.
sulf2	0.45	-2.223	sulfatase 2 (SULF2), transcript variant 1, mRNA.
mgat2	0.45	-2.223	mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT2), transcript variant 1, mRNA.
loc650518	0.45	-2.223	PREDICTED: similar to Proteasome subunit alpha type 6 (Proteasome iota chain) (Macropain iota chain) (Multicatalytic endopeptidase complex iota chain) (LOC650518), mRNA.
smg1	0.45	-2.224	PI-3-kinase-related kinase SMG-1 (SMG1), mRNA.
mkks	0.45	-2.224	McKusick-Kaufman syndrome (MKKS), transcript variant 1, mRNA.
tm9sf2	0.45	-2.224	transmembrane 9 superfamily member 2 (TM9SF2), mRNA.
hyls1	0.45	-2.224	hydrolethalus syndrome 1 (HYLS1), mRNA.
hmgn4	0.45	-2.224	high mobility group nucleosomal binding domain 4 (HMGN4), mRNA.
il13ra1	0.45	-2.224	interleukin 13 receptor, alpha 1 (IL13RA1), mRNA.
idua	0.45	-2.225	iduronidase, alpha-L- (IDUA), mRNA.
cox7b	0.449	-2.225	cytochrome c oxidase subunit VIIb (COX7B), nuclear gene encoding mitochondrial protein, mRNA.
guk1	0.449	-2.225	guanylate kinase 1 (GUK1), mRNA.
kiaa0922	0.449	-2.226	KIAA0922 (KIAA0922), mRNA.
aph1a	0.449	-2.226	anterior pharynx defective 1 homolog A (C. elegans) (APH1A), transcript variant 1, mRNA.
acd	0.449	-2.226	adrenocortical dysplasia homolog (mouse) (ACD), mRNA.
c9orf86	0.449	-2.226	chromosome 9 open reading frame 86 (C9orf86), mRNA.
mus81	0.449	-2.227	MUS81 endonuclease homolog (S. cerevisiae) (MUS81), mRNA.
rpusd1	0.449	-2.227	RNA pseudouridylate synthase domain containing 1 (RPUSD1), mRNA.
phf23	0.449	-2.227	PHD finger protein 23 (PHF23), mRNA.
atp6v1d	0.449	-2.228	ATPase, H+ transporting, lysosomal 34kDa, V1 subunit D (ATP6V1D), mRNA.
tufm	0.449	-2.228	Tu translation elongation factor, mitochondrial (TUFM), mRNA.
farslb	0.449	-2.228	phenylalanine-tRNA synthetase-like, beta subunit (FARSLB), mRNA.
hs.535392	0.449	-2.228	BX090843 Soares_testis_NHT cDNA clone IMAGp998C074110 ; IMAGE:1620414, mRNA sequence
tmem16k	0.449	-2.228	transmembrane protein 16K (TMEM16K), mRNA.
sdf2l1	0.449	-2.228	stromal cell-derived factor 2-like 1 (SDF2L1), mRNA.
polr1c	0.449	-2.229	polymerase (RNA) I polypeptide C, 30kDa (POLR1C), transcript variant 2, mRNA.
ifitm2	0.449	-2.229	interferon induced transmembrane protein 2 (1-8D) (IFITM2), mRNA.
gadd45a	0.449	-2.229	growth arrest and DNA-damage-inducible, alpha (GADD45A), mRNA.
hs.296031	0.449	-2.229	cDNA clone IMAGE:5262734
ppp1r2	0.449	-2.23	protein phosphatase 1, regulatory (inhibitor) subunit 2 (PPP1R2), mRNA.
wdr6	0.449	-2.23	WD repeat domain 6 (WDR6), mRNA.
slc40a1	0.448	-2.23	solute carrier family 40 (iron-regulated transporter), member 1 (SLC40A1), mRNA.
timm17b	0.448	-2.23	translocase of inner mitochondrial membrane 17 homolog B (yeast) (TIMM17B), mRNA.
gla	0.448	-2.23	galactosidase, alpha (GLA), mRNA.
mettl5	0.448	-2.231	methyltransferase like 5 (METTL5), mRNA.

prim1	0.448	-2.231	primase, DNA, polypeptide 1 (49kDa) (PRIM1), mRNA.
leprotl1	0.448	-2.231	leptin receptor overlapping transcript-like 1 (LEPROTL1), mRNA.
faim3	0.448	-2.231	Fas apoptotic inhibitory molecule 3 (FAIM3), mRNA.
ndufs2	0.448	-2.231	NADH dehydrogenase (ubiquinone) Fe-S protein 2, 49kDa (NADH-coenzyme Q reductase) (NDUFS2), mRNA.
psmb10	0.448	-2.232	proteasome (prosome, macropain) subunit, beta type, 10 (PSMB10), mRNA.
sntb2	0.448	-2.232	syntrophin, b2 (dystrophin-associated protein A1, 59kDa, basic component 2) (SNTB2), trans.variant 1, mRNA.
kiaa1893	0.448	-2.232	G protein-regulated inducer of neurite outgrowth 1 (KIAA1893), mRNA.
tomm40l	0.448	-2.232	translocase of outer mitochondrial membrane 40 homolog-like (yeast) (TOMM40L), mRNA.
tbc1d4	0.448	-2.232	TBC1 domain family, member 4 (TBC1D4), mRNA.
ppme1	0.448	-2.233	protein phosphatase methylesterase 1 (PPME1), mRNA.
mrpl53	0.448	-2.233	mitochondrial ribosomal protein L53 (MRPL53), nuclear gene encoding mitochondrial protein, mRNA.
fam82c	0.448	-2.233	family with sequence similarity 82, member C (FAM82C), mRNA.
lsm12	0.448	-2.233	LSM12 homolog (S. cerevisiae) (LSM12), mRNA.
trappc5	0.448	-2.233	trafficking protein particle complex 5 (TRAPPC5), transcript variant 3, mRNA.
bat2d1	0.448	-2.233	BAT2 domain containing 1 (BAT2D1), mRNA.
c10orf86	0.448	-2.233	non-SMC element 4 homolog A (S. cerevisiae) (NSMCE4A), mRNA.
clstn1	0.448	-2.233	calsyntenin 1 (CLSTN1), transcript variant 1, mRNA.
scap2	0.448	-2.234	src family associated phosphoprotein 2 (SCAP2), mRNA.
loc653158	0.448	-2.234	PREDICTED: similar to hypothetical protein MGC40405, transcript variant 1 (LOC653158), mRNA.
tmem41b	0.448	-2.234	transmembrane protein 41B (TMEM41B), mRNA.
mapre1	0.448	-2.234	microtubule-associated protein, RP/EB family, member 1 (MAPRE1), mRNA.
foxm1	0.448	-2.235	forkhead box M1 (FOXM1), transcript variant 1, mRNA.
stat5a	0.448	-2.235	signal transducer and activator of transcription 5A (STAT5A), mRNA.
rag1ap1	0.447	-2.235	recombination activating gene 1 activating protein 1 (RAG1AP1), mRNA.
mgc16824	0.447	-2.235	esophageal cancer associated protein (MGC16824), mRNA.
mgc13125	0.447	-2.235	BUD13 homolog (S. cerevisiae) (BUD13), mRNA.
pacs1	0.447	-2.235	phosphofurin acidic cluster sorting protein 1 (PACS1), mRNA.
aig1	0.447	-2.236	androgen-induced 1 (AIG1), mRNA.
dhx32	0.447	-2.236	DEAH (Asp-Glu-Ala-His) box polypeptide 32 (DHX32), mRNA.
havcr1	0.447	-2.236	hepatitis A virus cellular receptor 1 (HAVCR1), mRNA.
qtrt1	0.447	-2.236	queuine tRNA-ribosyltransferase 1 (tRNA-guanine transglycosylase) (QTRT1), mRNA.
flj13909	0.447	-2.237	hypothetical protein FLJ13909 (FLJ13909), mRNA.
tjp1	0.447	-2.237	tight junction protein 1 (zona occludens 1) (TJP1), transcript variant 1, mRNA.
lsmd1	0.447	-2.237	LSM domain containing 1 (LSMD1), mRNA.
syde1	0.447	-2.238	synapse defective 1, Rho GTPase, homolog 1 (C. elegans) (SYDE1), mRNA.
ppp1r7	0.447	-2.238	protein phosphatase 1, regulatory (inhibitor) subunit 7 (PPP1R7), mRNA.
sf3b2	0.447	-2.238	splicing factor 3b, subunit 2, 145kDa (SF3B2), mRNA.
frat2	0.447	-2.238	frequently rearranged in advanced T-cell lymphomas 2 (FRAT2), mRNA.
ddx5	0.447	-2.239	DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 (DDX5), mRNA.
orc3l	0.447	-2.239	origin recognition complex, subunit 3-like (yeast) (ORC3L), transcript variant 1, mRNA.
hs.4988	0.447	-2.239	mRNA; cDNA DKFZp686B24166 (from clone DKFZp686B24166)
znf358	0.447	-2.239	zinc finger protein 358 (ZNF358), mRNA.

ube2j1	0.447	-2.239	ubiquitin-conjugating enzyme E2, J1 (UBC6 homolog, yeast) (UBE2J1), mRNA.
ahctf1	0.447	-2.239	PREDICTED: AT hook containing transcription factor 1 (AHCTF1), mRNA.
scfd1	0.447	-2.239	sec1 family domain containing 1 (SCFD1), transcript variant 1, mRNA.
tnfrsf1a	0.447	-2.24	tumor necrosis factor receptor superfamily, member 1A (TNFRSF1A), mRNA.
gnl2	0.447	-2.24	guanine nucleotide binding protein-like 2 (nucleolar) (GNL2), mRNA.
loc645058	0.446	-2.24	PREDICTED: similar to hepatitis B virus x-interacting protein (LOC645058), mRNA.
phf11	0.446	-2.24	PHD finger protein 11 (PHF11), transcript variant 1, mRNA.
smu1	0.446	-2.241	smu-1 suppressor of mec-8 and unc-52 homolog (C. elegans) (SMU1), mRNA.
esd	0.446	-2.241	esterase D/formylglutathione hydrolase (ESD), mRNA.
ayp1p1	0.446	-2.241	PREDICTED: AYP1 pseudogene 1 (AYP1p1), misc RNA.
chsy1	0.446	-2.241	carbohydrate (chondroitin) synthase 1 (CHSY1), mRNA.
cd109	0.446	-2.241	CD109 antigen (Gov platelet alloantigens) (CD109), mRNA.
lmo1	0.446	-2.241	LIM domain only 1 (rhombotin 1) (LMO1), mRNA.
rce1	0.446	-2.241	RCE1 homolog, prenyl protein peptidase (S. cerevisiae) (RCE1), transcript variant 2, mRNA.
loc389787	0.446	-2.241	PREDICTED: similar to Translationally-controlled tumor protein (TCTP) (p23) (Histamine-releasing factor) (HRF) (Fortilin) (LOC389787), mRNA.
hs.483906	0.446	-2.242	mRNA; cDNA DKFZp667L2214 (from clone DKFZp667L2214)
u2af1l3	0.446	-2.242	U2(RNU2) small nuclear RNA auxiliary factor 1-like 3 (U2AF1L3), mRNA.
eml1	0.446	-2.242	echinoderm microtubule associated protein like 1 (EML1), transcript variant 2, mRNA.
rpl36	0.446	-2.242	ribosomal protein L36 (RPL36), transcript variant 2, mRNA.
sma4	0.446	-2.243	SMA4 (SMA4), mRNA.
hs.163752	0.446	-2.244	zr87e09.r1 NCI_CGAP_GCB1 cDNA clone IMAGE:682696 5, mRNA sequence
hmgb2	0.445	-2.245	high-mobility group box 2 (HMGB2), mRNA.
vbp1	0.445	-2.245	von Hippel-Lindau binding protein 1 (VBP1), mRNA.
fgfrl1	0.445	-2.245	fibroblast growth factor receptor-like 1 (FGFRL1), transcript variant 2, mRNA.
isgf3g	0.445	-2.245	interferon-stimulated transcription factor 3, gamma 48kDa (ISGF3G), mRNA.
mlph	0.445	-2.245	melanophilin (MLPH), mRNA.
znf499	0.445	-2.245	zinc finger and BTB domain containing 45 (ZBTB45), mRNA.
loc440157	0.445	-2.245	hypothetical gene supported by AK096951; BC066547 (LOC440157), mRNA.
gabarap	0.445	-2.246	GABA(A) receptor-associated protein (GABARAP), mRNA.
esrra	0.445	-2.246	estrogen-related receptor alpha (ESRRA), mRNA.
c11orf10	0.445	-2.246	chromosome 11 open reading frame 10 (C11orf10), mRNA.
toe1	0.445	-2.246	target of EGR1, member 1 (nuclear) (TOE1), mRNA.
rhbdl2	0.445	-2.247	rhomboid, veinlet-like 2 (Drosophila) (RHBDL2), mRNA.
psme4	0.445	-2.247	proteasome (prosome, macropain) activator subunit 4 (PSME4), mRNA.
cno	0.445	-2.247	cappuccino homolog (mouse) (CNO), mRNA.
mt1a	0.445	-2.247	metallothionein 1A (MT1A), mRNA.
kiaa0406	0.445	-2.248	KIAA0406 (KIAA0406), mRNA.
hnrpa2b1	0.445	-2.248	heterogeneous nuclear ribonucleoprotein A2/B1 (HNRPA2B1), transcript variant B1, mRNA.
pcbp2	0.445	-2.248	poly(rC) binding protein 2 (PCBP2), transcript variant 2, mRNA.
spc24	0.445	-2.248	SPC24, NDC80 kinetochore complex component, homolog (S. cerevisiae) (SPC24), mRNA.
hsa9761	0.445	-2.248	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae) (DIMT1L), mRNA.

c18orf55	0.445	-2.249	chromosome 18 open reading frame 55 (C18orf55), mRNA.
pcmtd1	0.445	-2.249	protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1 (PCMTD1), mRNA.
sars	0.445	-2.249	seryl-tRNA synthetase (SARS), mRNA.
flj20625	0.445	-2.249	chromosome 11 open reading frame 59 (C11orf59), mRNA.
marcksl1	0.444	-2.25	MARCKS-like 1 (MARCKSL1), mRNA.
c7orf21	0.444	-2.25	transmembrane and ubiquitin-like domain containing 1 (TMUB1), mRNA.
il27ra	0.444	-2.25	interleukin 27 receptor, alpha (IL27RA), mRNA.
c1orf31	0.444	-2.25	chromosome 1 open reading frame 31 (C1orf31), mRNA.
vps26a	0.444	-2.251	vacuolar protein sorting 26 homolog A (S. pombe) (VPS26A), transcript variant 2, mRNA.
scarb2	0.444	-2.251	scavenger receptor class B, member 2 (SCARB2), mRNA.
psma5	0.444	-2.251	proteasome (prosome, macropain) subunit, alpha type, 5 (PSMA5), mRNA.
hsph1	0.444	-2.251	heat shock 105kDa/110kDa protein 1 (HSPH1), mRNA.
tbc1d8	0.444	-2.251	TBC1 domain family, member 8 (with GRAM domain) (TBC1D8), mRNA.
kiaa0090	0.444	-2.251	KIAA0090 (KIAA0090), mRNA.
wdr59	0.444	-2.253	WD repeat domain 59 (WDR59), mRNA.
scyl1	0.444	-2.253	SCY1-like 1 (S. cerevisiae) (SCYL1), mRNA.
ift52	0.444	-2.253	intraflagellar transport 52 homolog (Chlamydomonas) (IFT52), mRNA.
snx12	0.444	-2.253	sorting nexin 12 (SNX12), mRNA.
poldip3	0.444	-2.253	polymerase (DNA-directed), delta interacting protein 3 (POLDIP3), transcript variant 1, mRNA.
cops4	0.444	-2.253	COP9 constitutive photomorphogenic homolog subunit 4 (Arabidopsis) (COPS4), mRNA.
znf174	0.444	-2.254	zinc finger protein 174 (ZNF174), transcript variant 2, mRNA.
foxk1	0.444	-2.254	forkhead box K1 (FOXK1), mRNA.
aspscr1	0.444	-2.254	alveolar soft part sarcoma chromosome region, candidate 1 (ASPSCR1), mRNA.
pip5k1c	0.444	-2.254	phosphatidylinositol-4-phosphate 5-kinase, type I, gamma (PIP5K1C), mRNA.
pir	0.444	-2.254	pirin (iron-binding nuclear protein) (PIR), transcript variant 2, mRNA.
slc35a2	0.444	-2.254	solute carrier family 35 (UDP-galactose transporter), member A2 (SLC35A2), transcript variant 1, mRNA.
hibch	0.444	-2.254	3-hydroxyisobutyryl-Coenzyme A hydrolase (HIBCH), transcript variant 1, mRNA.
sephs1	0.444	-2.255	selenophosphate synthetase 1 (SEPHS1), mRNA.
hs.145049	0.444	-2.255	full-length cDNA clone CS0DF005YI08 of Fetal brain of (human)
exoc2	0.443	-2.255	exocyst complex component 2 (EXOC2), mRNA.
gnptab	0.443	-2.256	N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits (GNPTAB), mRNA.
tmem126b	0.443	-2.256	transmembrane protein 126B (TMEM126B), mRNA.
p4ha2	0.443	-2.256	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II (P4HA2), transcript variant 3, mRNA.
ccnk	0.443	-2.256	cyclin K (CCNK), mRNA.
c1orf171	0.443	-2.256	tRNA-yW synthesizing protein 3 homolog (S. cerevisiae) (TYW3), mRNA.
larp5	0.443	-2.256	La ribonucleoprotein domain family, member 5 (LARP5), mRNA.
slc27a5	0.443	-2.257	solute carrier family 27 (fatty acid transporter), member 5 (SLC27A5), mRNA.
parva	0.443	-2.257	parvin, alpha (PARVA), mRNA.
hs.374278	0.443	-2.257	cDNA FLJ38388 fis, clone FEBRA2004485
lat1-3tm	0.443	-2.257	PREDICTED: SLC7A5 pseudogene, transcript variant 2 (LAT1-3TM), misc RNA.
cd2bp2	0.443	-2.257	CD2 (cytoplasmic tail) binding protein 2 (CD2BP2), mRNA.

psenen	0.443	-2.258	presenilin enhancer 2 homolog (C. elegans) (PSENEN), mRNA.
loc114984	0.443	-2.258	FLYWCH family member 2 (FLYWCH2), mRNA.
ptpre	0.443	-2.258	protein tyrosine phosphatase, receptor type, E (PTPRE), transcript variant 2, mRNA.
loc642989	0.443	-2.258	PREDICTED: similar to 40S ribosomal protein S25 (LOC642989), mRNA.
baz2b	0.443	-2.258	bromodomain adjacent to zinc finger domain, 2B (BAZ2B), mRNA.
hook2	0.443	-2.258	hook homolog 2 (Drosophila) (HOOK2), mRNA.
ifrd1	0.443	-2.259	interferon-related developmental regulator 1 (IFRD1), transcript variant 2, mRNA.
prmt2	0.443	-2.259	protein arginine methyltransferase 2 (PRMT2), transcript variant 1, mRNA.
mrpl32	0.443	-2.259	mitochondrial ribosomal protein L32 (MRPL32), nuclear gene encoding mitochondrial protein, mRNA.
athl1	0.443	-2.259	ATH1, acid trehalase-like 1 (yeast) (ATHL1), mRNA.
cse1l	0.443	-2.259	CSE1 chromosome segregation 1-like (yeast) (CSE1L), transcript variant 2, mRNA.
e2f3	0.443	-2.259	E2F transcription factor 3 (E2F3), mRNA.
mylip	0.443	-2.26	myosin regulatory light chain interacting protein (MYLIP), mRNA.
arl6ip4	0.442	-2.26	ADP-ribosylation-like factor 6 interacting protein 4 (ARL6IP4), transcript variant 3, mRNA.
rnd3	0.442	-2.26	Rho family GTPase 3 (RND3), mRNA.
ddx19a	0.442	-2.26	DEAD (Asp-Glu-Ala-As) box polypeptide 19A (DDX19A), mRNA.
rnaset2	0.442	-2.26	ribonuclease T2 (RNASET2), mRNA.
lypla3	0.442	-2.261	lysophospholipase 3 (lysosomal phospholipase A2) (LYPLA3), mRNA.
fras1	0.442	-2.261	Fraser syndrome 1 (FRAS1), transcript variant 2, mRNA.
trim11	0.442	-2.261	tripartite motif-containing 11 (TRIM11), mRNA.
pin4	0.442	-2.262	protein (peptidylprolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin) (PIN4), mRNA.
loc347376	0.442	-2.262	PREDICTED: similar to H3 histone, family 3B (LOC347376), mRNA.
c20orf44	0.442	-2.262	chromosome 20 open reading frame 44 (C20orf44), transcript variant 3, mRNA.
rbm38	0.442	-2.262	RNA binding motif protein 38 (RBM38), transcript variant 2, mRNA.
nfyc	0.442	-2.262	nuclear transcription factor Y, gamma (NFYC), mRNA.
dtl	0.442	-2.262	denticleless homolog (Drosophila) (DTL), mRNA.
loc651831	0.442	-2.263	PREDICTED: similar to CG31232-PA, isoform A (LOC651831), mRNA.
map1lc3b	0.442	-2.263	microtubule-associated protein 1 light chain 3 beta (MAP1LC3B), mRNA.
obfc2b	0.442	-2.264	oligonucleotide/oligosaccharide-binding fold containing 2B (OBFC2B), mRNA.
cbr3	0.442	-2.264	carbonyl reductase 3 (CBR3), mRNA.
rarres1	0.442	-2.264	retinoic acid receptor responder (tazarotene induced) 1 (RARRES1), transcript variant 2, mRNA.
kiaa1598	0.442	-2.264	KIAA1598 (KIAA1598), mRNA.
pnrc2	0.442	-2.264	proline-rich nuclear receptor coactivator 2 (PNRC2), mRNA.
syncrip	0.442	-2.264	synaptotagmin binding, cytoplasmic RNA interacting protein (SYNCRIP), mRNA.
eif1	0.442	-2.265	eukaryotic translation initiation factor 1 (EIF1), mRNA.
mcm3ap	0.442	-2.265	MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) associated protein (MCM3AP), mRNA.
pprc1	0.442	-2.265	peroxisome proliferator-activated receptor gamma, coactivator-related 1 (PPRC1), mRNA.
c12orf24	0.441	-2.265	chromosome 12 open reading frame 24 (C12orf24), mRNA.
adpgk	0.441	-2.265	ADP-dependent glucokinase (ADPGK), mRNA.
mgc2752	0.441	-2.265	hypothetical protein MGC2752 (MGC2752), mRNA.
cic	0.441	-2.266	capicua homolog (Drosophila) (CIC), mRNA.
znf622	0.441	-2.266	zinc finger protein 622 (ZNF622), mRNA.

nono	0.441	-2.266	non-POU domain containing, octamer-binding (NONO), mRNA.
ephb2	0.441	-2.266	EPH receptor B2 (EPHB2), transcript variant 1, mRNA.
tada1l	0.441	-2.267	transcriptional adaptor 1 (HFI1 homolog, yeast)-like (TADA1L), mRNA.
fadd	0.441	-2.267	Fas (TNFRSF6)-associated via death domain (FADD), mRNA.
trrap	0.441	-2.267	transformation/transcription domain-associated protein (TRRAP), mRNA.
c14orf172	0.441	-2.267	chromosome 14 open reading frame 172 (C14orf172), mRNA.
loc129138	0.441	-2.267	ankyrin repeat domain 54 (ANKRD54), mRNA.
hif1a	0.441	-2.267	hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor) (HIF1A), transcript variant 1, mRNA.
arhgap19	0.441	-2.268	Rho GTPase activating protein 19 (ARHGAP19), mRNA.
hs.505676	0.441	-2.268	cDNA FLJ33772 fis, clone BRSSN2000175
rcor3	0.441	-2.268	REST corepressor 3 (RCOR3), mRNA.
ddt	0.441	-2.268	D-dopachrome tautomerase (DDT), mRNA.
srebf1	0.441	-2.268	sterol regulatory element binding transcription factor 1 (SREBF1), transcript variant 2, mRNA.
ells1	0.441	-2.268	hypothetical protein Ells1 (Ells1), mRNA.
kctd10	0.441	-2.269	potassium channel tetramerisation domain containing 10 (KCTD10), mRNA.
c14orf80	0.441	-2.269	chromosome 14 open reading frame 80 (C14orf80), mRNA.
crot	0.441	-2.269	carnitine O-octanoyltransferase (CROT), mRNA.
ccdc53	0.441	-2.269	coiled-coil domain containing 53 (CCDC53), mRNA.
srm	0.441	-2.269	spermidine synthase (SRM), mRNA.
isoc1	0.441	-2.269	isochorismatase domain containing 1 (ISOC1), mRNA.
mgc27345	0.441	-2.269	PREDICTED: hypothetical protein MGC27345 (MGC27345), misc RNA.
c19orf7	0.441	-2.269	chromosome 19 open reading frame 7 (C19orf7), mRNA.
h2afx	0.441	-2.269	H2A histone family, member X (H2AFX), mRNA.
c12orf47	0.441	-2.27	chromosome 12 open reading frame 47 (C12orf47), mRNA.
mpp5	0.441	-2.27	membrane protein, palmitoylated 5 (MAGUK p55 subfamily member 5) (MPP5), mRNA.
cav1	0.44	-2.27	caveolin 1, caveolae protein, 22kDa (CAV1), mRNA.
wibg	0.44	-2.271	within bgcn homolog (Drosophila) (WIBG), mRNA.
slc35e3	0.44	-2.271	solute carrier family 35, member E3 (SLC35E3), mRNA.
fh	0.44	-2.271	fumarate hydratase (FH), nuclear gene encoding mitochondrial protein, mRNA.
eif4ebp2	0.44	-2.271	eukaryotic translation initiation factor 4E binding protein 2 (EIF4EBP2), mRNA.
hebp2	0.44	-2.272	heme binding protein 2 (HEBP2), mRNA.
igf2bp3	0.44	-2.272	insulin-like growth factor 2 mRNA binding protein 3 (IGF2BP3), mRNA.
cstb	0.44	-2.273	cystatin B (stefin B) (CSTB), mRNA.
rpa1	0.44	-2.273	replication protein A1, 70kDa (RPA1), mRNA.
irs1	0.44	-2.274	insulin receptor substrate 1 (IRS1), mRNA.
atox1	0.44	-2.274	ATX1 antioxidant protein 1 homolog (yeast) (ATOX1), mRNA.
sema4f	0.44	-2.274	sema domain, immunoglobulin domain (Ig), transmembrane domain ™ and short cytoplasmic domain, (semaphorin) 4F (SEMA4F), mRNA.
spin1	0.44	-2.274	spinster homolog 1 (Drosophila) (SPNS1), mRNA.
tsc2	0.44	-2.275	tuberous sclerosis 2 (TSC2), transcript variant 1, mRNA.
idh2	0.44	-2.275	isocitrate dehydrogenase 2 (NADP+), mitochondrial (IDH2), mRNA.

tm2d2	0.44	-2.275	TM2 domain containing 2 (TM2D2), transcript variant 1, mRNA.
uxt	0.44	-2.275	ubiquitously-expressed transcript (UXT), transcript variant 1, mRNA.
arl1	0.44	-2.275	ADP-ribosylation factor-like 1 (ARL1), mRNA.
c6orf61	0.44	-2.275	PREDICTED: chromosome 6 open reading frame 61 (C6orf61), mRNA.
fvt1	0.439	-2.276	follicular lymphoma variant translocation 1 (FVT1), mRNA.
c1orf69	0.439	-2.276	chromosome 1 open reading frame 69 (C1orf69), mRNA.
lmo4	0.439	-2.276	LIM domain only 4 (LMO4), mRNA.
derl1	0.439	-2.276	Der1-like domain family, member 1 (DERL1), mRNA.
slc24a6	0.439	-2.276	solute carrier family 24 (sodium/potassium/calcium exchanger), member 6 (SLC24A6), mRNA.
alkbh8	0.439	-2.276	alkB, alkylation repair homolog 8 (E. coli) (ALKBH8), mRNA.
mt1g	0.439	-2.276	metallothionein 1G (MT1G), mRNA.
ablim3	0.439	-2.277	actin binding LIM protein family, member 3 (ABLIM3), mRNA.
prkcq	0.439	-2.277	protein kinase C, theta (PRKCQ), mRNA.
loc402251	0.439	-2.277	PREDICTED: similar to eukaryotic translation elongation factor 1 alpha 2 (LOC402251), mRNA.
med19	0.439	-2.277	mediator complex subunit 19 (MED19), mRNA.
bivm	0.439	-2.277	basic, immunoglobulin-like variable motif containing (BIVM), mRNA.
kiaa0753	0.439	-2.278	KIAA0753 (KIAA0753), mRNA.
wsb2	0.439	-2.278	WD repeat and SOCS box-containing 2 (WSB2), mRNA.
p4hb	0.439	-2.278	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (P4HB), mRNA.
gnai2	0.439	-2.278	guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 2 (GNAI2), mRNA.
gtf3c2	0.439	-2.278	general transcription factor IIIC, polypeptide 2, beta 110kDa (GTF3C2), transcript variant 2, mRNA.
foxred1	0.439	-2.278	FAD-dependent oxidoreductase domain containing 1 (FOXRED1), mRNA.
smug1	0.439	-2.278	single-strand-selective monofunctional uracil-DNA glycosylase 1 (SMUG1), mRNA.
prkab2	0.439	-2.279	protein kinase, AMP-activated, beta 2 non-catalytic subunit (PRKAB2), mRNA.
syvn1	0.439	-2.279	synovial apoptosis inhibitor 1, synoviolin (SYVN1), transcript variant 2, mRNA.
flj10986	0.439	-2.279	hypothetical protein FLJ10986 (FLJ10986), mRNA.
exosc10	0.439	-2.279	exosome component 10 (EXOSC10), transcript variant 1, mRNA.
dbi	0.439	-2.279	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein) (DBI), mRNA.
kdelr1	0.439	-2.28	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1 (KDELR1), mRNA.
htatip2	0.439	-2.28	HIV-1 Tat interactive protein 2, 30kDa (HTATIP2), mRNA.
flj20558	0.439	-2.28	chromosome 2 open reading frame 42 (C2orf42), mRNA.
slc10a3	0.439	-2.28	solute carrier family 10 (sodium/bile acid cotransporter family), member 3 (SLC10A3), mRNA.
spred1	0.439	-2.28	sprouty-related, EVH1 domain containing 1 (SPRED1), mRNA.
lhfp	0.439	-2.28	lipoma HMGIC fusion partner (LHFP), mRNA.
sdf2	0.439	-2.28	stromal cell-derived factor 2 (SDF2), mRNA.
tp53rk	0.439	-2.28	TP53 regulating kinase (TP53RK), mRNA.
enosf1	0.439	-2.28	enolase superfamily member 1 (ENOSF1), mRNA.
sult1a4	0.439	-2.281	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 4 (SULT1A4), transcript variant 3, mRNA.
gpr172a	0.439	-2.281	G protein-coupled receptor 172A (GPR172A), mRNA.
fen1	0.439	-2.281	flap structure-specific endonuclease 1 (FEN1), mRNA.
rxra	0.438	-2.281	retinoid X receptor, alpha (RXRA), mRNA.
kiaa0494	0.438	-2.281	KIAA0494 (KIAA0494), mRNA.

mettl2a	0.438	-2.281	methyltransferase like 2A (METTL2A), mRNA.
ccdc71	0.438	-2.281	coiled-coil domain containing 71 (CCDC71), mRNA.
ttc32	0.438	-2.281	tetratricopeptide repeat domain 32 (TTC32), mRNA.
c16orf63	0.438	-2.281	chromosome 16 open reading frame 63 (C16orf63), mRNA.
zmpste24	0.438	-2.282	zinc metallopeptidase (STE24 homolog, yeast) (ZMPSTE24), mRNA.
rbm15b	0.438	-2.282	RNA binding motif protein 15B (RBM15B), mRNA.
tmem185a	0.438	-2.282	transmembrane protein 185A (TMEM185A), mRNA.
tceal3	0.438	-2.282	transcription elongation factor A (SII)-like 3 (TCEAL3), transcript variant 1, mRNA.
slc30a7	0.438	-2.283	solute carrier family 30 (zinc transporter), member 7 (SLC30A7), mRNA.
gtf2ird1	0.438	-2.283	GTF2I repeat domain containing 1 (GTF2IRD1), transcript variant 1, mRNA.
mapkapk5	0.438	-2.283	mitogen-activated protein kinase-activated protein kinase 5 (MAPKAPK5), transcript variant 2, mRNA.
mrpl35	0.438	-2.284	mitochondrial ribosomal protein L35 (MRPL35), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
loc168850	0.438	-2.284	hypothetical protein LOC168850 (LOC168850), mRNA.
nupr1	0.438	-2.284	nuclear protein 1 (NUPR1), transcript variant 1, mRNA.
loc400657	0.438	-2.284	hypothetical gene supported by BC036588 (LOC400657), mRNA.
gng11	0.438	-2.284	guanine nucleotide binding protein (G protein), gamma 11 (GNG11), mRNA.
phacs	0.438	-2.284	1-aminocyclopropane-1-carboxylate synthase (PHACS), mRNA.
c14orf143	0.438	-2.285	chromosome 14 open reading frame 143 (C14orf143), mRNA.
nrbp2	0.438	-2.285	nuclear receptor binding protein 2 (NRBP2), mRNA.
cpsf3	0.438	-2.285	cleavage and polyadenylation specific factor 3, 73kDa (CPSF3), mRNA.
tubb2c	0.438	-2.285	tubulin, beta 2C (TUBB2C), mRNA.
efhc1	0.437	-2.286	EF-hand domain (C-terminal) containing 1 (EFHC1), mRNA.
tmem15	0.437	-2.286	transmembrane protein 15 (TMEM15), mRNA.
pomp	0.437	-2.287	proteasome maturation protein (POMP), mRNA.
perld1	0.437	-2.287	per1-like domain containing 1 (PERLD1), mRNA.
nup205	0.437	-2.287	nucleoporin 205kDa (NUP205), mRNA.
wbscr27	0.437	-2.287	Williams Beuren syndrome chromosome region 27 (WBSCR27), mRNA.
dhx30	0.437	-2.287	DEAH (Asp-Glu-Ala-His) box polypeptide 30 (DHX30), transcript variant 2, mRNA.
loc650276	0.437	-2.287	PREDICTED: similar to 60S ribosomal protein L7 (LOC650276), mRNA.
scamp4	0.437	-2.288	secretory carrier membrane protein 4 (SCAMP4), mRNA.
iars2	0.437	-2.288	isoleucyl-tRNA synthetase 2, mitochondrial (IARS2), mRNA.
c10orf6	0.437	-2.289	chromosome 10 open reading frame 6 (C10orf6), mRNA.
oact5	0.437	-2.289	membrane bound O-acyltransferase domain containing 5 (MBOAT5), mRNA.
tmem43	0.437	-2.289	transmembrane protein 43 (TMEM43), mRNA.
prkra	0.437	-2.289	protein kinase, interferon-inducible double stranded RNA dependent activator (PRKRA), mRNA.
zcchc11	0.437	-2.289	zinc finger, CCHC domain containing 11 (ZCCHC11), transcript variant 2, mRNA.
maged1	0.437	-2.289	melanoma antigen family D, 1 (MAGED1), transcript variant 1, mRNA.
nfe2l2	0.437	-2.29	nuclear factor (erythroid-derived 2)-like 2 (NFE2L2), mRNA.
cnih	0.437	-2.29	cornichon homolog (Drosophila) (CNIH), transcript variant 2, mRNA.
mgc71993	0.437	-2.29	similar to DNA segment, Chr 11, Brigham & Womens Genetics 0434 expressed (MGC71993), mRNA.
loc440341	0.437	-2.29	PREDICTED: similar to nuclear pore complex interacting protein, transcript variant 4 (LOC440341), mRNA.
eny2	0.437	-2.29	enhancer of yellow 2 homolog (Drosophila) (ENY2), mRNA.

loc284230	0.436	-2.291	PREDICTED: similar to large subunit ribosomal protein L36a (LOC284230), mRNA.
pgls	0.436	-2.291	6-phosphogluconolactonase (PGLS), mRNA.
hyal3	0.436	-2.291	hyaluronoglucosaminidase 3 (HYAL3), mRNA.
rnu70	0.436	-2.291	small nucleolar RNA, H/ACA box 70 (SNORA70) on chromosome X.
diaph2	0.436	-2.292	diaphanous homolog 2 (Drosophila) (DIAPH2), transcript variant 12C, mRNA.
tspan14	0.436	-2.292	tetraspanin 14 (TSPAN14), mRNA.
utp11l	0.436	-2.292	UTP11-like, U3 small nucleolar ribonucleoprotein, (yeast) (UTP11L), mRNA.
zfpm1	0.436	-2.292	zinc finger protein, multitype 1 (ZFPM1), mRNA.
ifi6	0.436	-2.293	interferon, alpha-inducible protein 6 (IFI6), transcript variant 2, mRNA.
mras	0.436	-2.293	muscle RAS oncogene homolog (MRAS), mRNA.
epc1	0.436	-2.293	enhancer of polycomb homolog 1 (Drosophila) (EPC1), mRNA.
c1orf60	0.436	-2.293	integrator complex subunit 3 (INTS3), mRNA.
hes4	0.436	-2.294	hairy and enhancer of split 4 (Drosophila) (HES4), mRNA.
smarcd2	0.436	-2.294	SWI/SNF related, matrix associated, actin dependent regulator of chromatin , subfamily d, member 2 (SMARCD2), mRNA.
taok1	0.436	-2.294	TAO kinase 1 (TAOK1), mRNA.
plaur	0.436	-2.294	plasminogen activator, urokinase receptor (PLAUR), transcript variant 2, mRNA.
loc645367	0.436	-2.295	PREDICTED: similar to Gamma-glutamyltranspeptidase 1 precursor (Gamma-glutamyltransferase 1) (CD224 antigen), transcript variant 2 (LOC645367), mRNA.
mtif2	0.436	-2.295	mitochondrial translational initiation factor 2 (MTIF2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
nicn1	0.436	-2.295	nicolin 1 (NICN1), mRNA.
glg1	0.436	-2.295	golgi apparatus protein 1 (GLG1), mRNA.
atp6v0d1	0.436	-2.295	ATPase, H+ transporting, lysosomal 38kDa, V0 subunit d isoform 1 (ATP6V0D1), mRNA.
c22orf5	0.436	-2.295	transmembrane protein 184B (TMEM184B), mRNA.
rpl35	0.435	-2.296	ribosomal protein L35 (RPL35), mRNA.
oprl1	0.435	-2.297	opiate receptor-like 1 (OPRL1), transcript variant 1, mRNA.
c14orf120	0.435	-2.297	PREDICTED: chromosome 14 open reading frame 120, transcript variant 11 (C14orf120), mRNA.
hectd1	0.435	-2.297	HECT domain containing 1 (HECTD1), mRNA.
rabgap1	0.435	-2.297	RAB GTPase activating protein 1 (RABGAP1), mRNA.
c1orf128	0.435	-2.297	chromosome 1 open reading frame 128 (C1orf128), mRNA.
vps28	0.435	-2.297	vacuolar protein sorting 28 homolog (S. cerevisiae) (VPS28), transcript variant 1, mRNA.
rbms2	0.435	-2.297	RNA binding motif, single stranded interacting protein 2 (RBMS2), mRNA.
stc1	0.435	-2.298	stanniocalcin 1 (STC1), mRNA.
c11orf70	0.435	-2.298	chromosome 11 open reading frame 70 (C11orf70), mRNA.
loc642033	0.435	-2.298	PREDICTED: similar to ATP-binding cassette, sub-family F, member 1 isoform b (LOC642033), mRNA.
znf148	0.435	-2.298	zinc finger protein 148 (ZNF148), mRNA.
gbl	0.435	-2.298	G protein beta subunit-like (GBL), mRNA.
dap3	0.435	-2.299	death associated protein 3 (DAP3), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
ngly1	0.435	-2.299	N-glycanase 1 (NGLY1), mRNA.
cep68	0.435	-2.299	centrosomal protein 68kDa (CEP68), mRNA.
nol6	0.435	-2.299	nucleolar protein family 6 (RNA-associated) (NOL6), transcript variant alpha, mRNA.

snw1	0.435	-2.299	SNW domain containing 1 (SNW1), mRNA.
cul5	0.435	-2.299	cullin 5 (CUL5), mRNA.
ifnar2	0.435	-2.3	interferon (alpha, beta and omega) receptor 2 (IFNAR2), transcript variant 3, mRNA.
gnb2	0.435	-2.3	guanine nucleotide binding protein (G protein), beta polypeptide 2 (GNB2), mRNA.
ndufs4	0.435	-2.3	NADH dehydrogenase (ubiquinone) Fe-S protein 4, 18kDa (NADH-coenzyme Q reductase) (NDUFS4), mRNA.
c1qdc1	0.435	-2.3	caprin family member 2 (CAPRIN2), transcript variant 2, mRNA.
ttc15	0.435	-2.3	tetratricopeptide repeat domain 15 (TTC15), mRNA.
tpst1	0.435	-2.3	tyrosylprotein sulfotransferase 1 (TPST1), mRNA.
map3k2	0.435	-2.3	PREDICTED: mitogen-activated protein kinase kinase kinase 2, transcript variant 3 (MAP3K2), mRNA.
pdcd6	0.435	-2.3	programmed cell death 6 (PDCD6), mRNA.
terf1	0.435	-2.301	telomeric repeat binding factor (NIMA-interacting) 1 (TERF1), transcript variant 2, mRNA.
osbpl7	0.435	-2.301	oxysterol binding protein-like 7 (OSBPL7), transcript variant 1, mRNA.
mtp18	0.435	-2.301	mitochondrial protein 18 kDa (MTP18), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
bcar1	0.435	-2.301	breast cancer anti-estrogen resistance 1 (BCAR1), mRNA.
iqsec1	0.434	-2.302	IQ motif and Sec7 domain 1 (IQSEC1), mRNA.
f8a1	0.434	-2.302	coagulation factor VIII-associated (intronic transcript) 1 (F8A1), mRNA.
c12orf23	0.434	-2.302	chromosome 12 open reading frame 23 (C12orf23), mRNA.
abcb6	0.434	-2.302	ATP-binding cassette, sub-family B (MDR/TAP), member 6 (ABCB6), nuclear gene encoding mitochondrial protein, mRNA.
gng10	0.434	-2.303	guanine nucleotide binding protein (G protein), gamma 10 (GNG10), mRNA.
kdelr3	0.434	-2.303	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3 (KDELR3), transcript variant 2, mRNA.
loc441019	0.434	-2.303	PREDICTED: hypothetical LOC441019 (LOC441019), mRNA.
ddx23	0.434	-2.304	DEAD (Asp-Glu-Ala-Asp) box polypeptide 23 (DDX23), mRNA.
eef1e1	0.434	-2.304	eukaryotic translation elongation factor 1 epsilon 1 (EEF1E1), mRNA.
hs.348514	0.434	-2.304	Homo sapiens, clone IMAGE:4052238, mRNA, partial cds
sfrs5	0.434	-2.304	splicing factor, arginine/serine-rich 5 (SFRS5), transcript variant 2, mRNA.
znf217	0.434	-2.305	zinc finger protein 217 (ZNF217), mRNA.
il11	0.434	-2.305	interleukin 11 (IL11), mRNA.
jtv1	0.434	-2.305	JTV1 gene (JTV1), mRNA.
rnase4	0.434	-2.306	ribonuclease, RNase A family, 4 (RNASE4), transcript variant 1, mRNA.
pdhx	0.434	-2.306	pyruvate dehydrogenase complex, component X (PDHX), mRNA.
abhd14a	0.434	-2.306	abhydrolase domain containing 14A (ABHD14A), mRNA.
flna	0.434	-2.306	filamin A, alpha (actin binding protein 280) (FLNA), mRNA.
wdr40a	0.433	-2.307	WD repeat domain 40A (WDR40A), mRNA.
loc644033	0.433	-2.307	PREDICTED: similar to similar to RPL23AP7 protein (LOC644033), mRNA.
cdc42bpb	0.433	-2.308	CDC42 binding protein kinase beta (DMPK-like) (CDC42BPB), mRNA.
rad23a	0.433	-2.308	RAD23 homolog A (S. cerevisiae) (RAD23A), mRNA.
acin1	0.433	-2.308	apoptotic chromatin condensation inducer 1 (ACIN1), mRNA.
ptd008	0.433	-2.309	chromosome 19 open reading frame 56 (C19orf56), mRNA.
ap2b1	0.433	-2.309	adaptor-related protein complex 2, beta 1 subunit (AP2B1), transcript variant 2, mRNA.
hnrpc	0.433	-2.309	heterogeneous nuclear ribonucleoprotein C (C1/C2) (HNRPC), transcript variant 3, mRNA.
g3bp1	0.433	-2.309	GTPase activating protein (SH3 domain) binding protein 1 (G3BP1), transcript variant 2, mRNA.

loc652864	0.433	-2.309	PREDICTED: similar to Mitochondrial import inner membrane translocase subunit Tim23 (LOC652864), mRNA.
mccc1	0.433	-2.31	methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) (MCCC1), mRNA.
dmap1	0.433	-2.31	DNA methyltransferase 1 associated protein 1 (DMAP1), transcript variant 3, mRNA.
mrps7	0.433	-2.31	mitochondrial ribosomal protein S7 (MRPS7), nuclear gene encoding mitochondrial protein, mRNA.
ier3ip1	0.433	-2.31	immediate early response 3 interacting protein 1 (IER3IP1), mRNA.
gfpt1	0.433	-2.31	glutamine-fructose-6-phosphate transaminase 1 (GFPT1), mRNA.
usp10	0.433	-2.31	ubiquitin specific peptidase 10 (USP10), mRNA.
thnsl1	0.433	-2.311	threonine synthase-like 1 (bacterial) (THNSL1), mRNA.
slc25a37	0.433	-2.311	solute carrier family 25, member 37 (SLC25A37), transcript variant 1, mRNA.
pdia5	0.433	-2.312	protein disulfide isomerase family A, member 5 (PDIA5), mRNA.
wnk1	0.432	-2.313	WNK lysine deficient protein kinase 1 (WNK1), mRNA.
hspc171	0.432	-2.313	HSPC171 protein (HSPC171), mRNA.
vps33b	0.432	-2.313	vacuolar protein sorting 33B (yeast) (VPS33B), mRNA.
flj22709	0.432	-2.314	occludin/ELL domain containing 1 (OCEL1), mRNA.
upf3a	0.432	-2.314	UPF3 regulator of nonsense transcripts homolog A (yeast) (UPF3A), transcript variant 2, mRNA.
pkmyt1	0.432	-2.315	protein kinase, membrane associated tyrosine/threonine 1 (PKMYT1), transcript variant 2, mRNA.
msh6	0.432	-2.315	mutS homolog 6 (E. coli) (MSH6), mRNA.
git1	0.432	-2.315	G protein-coupled receptor kinase interactor 1 (GIT1), mRNA.
hs.334831	0.432	-2.315	mRNA; cDNA DKFZp686I05206 (from clone DKFZp686I05206)
slc41a1	0.432	-2.316	solute carrier family 41, member 1 (SLC41A1), mRNA.
naca	0.432	-2.316	nascent-polypeptide-associated complex alpha polypeptide (NACA), mRNA.
rexo4	0.432	-2.316	REX4, RNA exonuclease 4 homolog (S. cerevisiae) (REXO4), mRNA.
rnf26	0.432	-2.316	ring finger protein 26 (RNF26), mRNA.
ccdc25	0.432	-2.316	coiled-coil domain containing 25 (CCDC25), transcript variant 2, mRNA.
nupl2	0.432	-2.317	nucleoporin like 2 (NUPL2), mRNA.
loc133619	0.432	-2.317	proline-rich coiled-coil 1 (PRRC1), mRNA.
commd6	0.432	-2.317	COMM domain containing 6 (COMMD6), transcript variant 1, mRNA.
afg3l1	0.432	-2.317	AFG3 ATPase family gene 3-like 1 (S. cerevisiae) (AFG3L1) on chromosome 16.
mea1	0.432	-2.317	male-enhanced antigen 1 (MEA1), mRNA.
xylt2	0.432	-2.317	xylosyltransferase II (XYLT2), mRNA.
loc400566	0.432	-2.317	hypothetical gene supported by AK128660 (LOC400566), mRNA.
sfrs14	0.432	-2.317	splicing factor, arginine/serine-rich 14 (SFRS14), transcript variant 1, mRNA.
ppt1	0.432	-2.318	palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile) (PPT1), mRNA.
tnfsf12	0.431	-2.318	tumor necrosis factor (ligand) superfamily, member 12 (TNFSF12), transcript variant 2, mRNA.
dctn5	0.431	-2.318	dynactin 5 (p25) (DCTN5), mRNA.
cyb561d2	0.431	-2.318	cytochrome b-561 domain containing 2 (CYB561D2), mRNA.
znf329	0.431	-2.318	zinc finger protein 329 (ZNF329), mRNA.
bxdc5	0.431	-2.318	brix domain containing 5 (BXDC5), mRNA.
kiaa0195	0.431	-2.318	KIAA0195 (KIAA0195), mRNA.
traip	0.431	-2.318	TRAF interacting protein (TRAIP), mRNA.
tnfsf9	0.431	-2.318	tumor necrosis factor (ligand) superfamily, member 9 (TNFSF9), mRNA.
kua	0.431	-2.319	ubiquitin-conjugating enzyme variant Kua (Kua), mRNA.

lig1	0.431	-2.319	ligase I, DNA, ATP-dependent (LIG1), mRNA.
ccnh	0.431	-2.319	cyclin H (CCNH), mRNA.
rbm4	0.431	-2.319	PREDICTED: RNA binding motif protein 4, transcript variant 3 (RBM4), mRNA.
cdca3	0.431	-2.319	cell division cycle associated 3 (CDCA3), mRNA.
mgc21881	0.431	-2.319	hypothetical protein MGC21881 (MGC21881), mRNA.
nbpf11	0.431	-2.32	neuroblastoma breakpoint family, member 11 (NBPF11), mRNA.
cthrc1	0.431	-2.32	collagen triple helix repeat containing 1 (CTHRC1), mRNA.
ccdc45	0.431	-2.32	coiled-coil domain containing 45 (CCDC45), mRNA.
c1orf163	0.431	-2.321	chromosome 1 open reading frame 163 (C1orf163), mRNA.
fbxo2	0.431	-2.321	F-box protein 2 (FBXO2), mRNA.
cabc1	0.431	-2.322	chaperone, ABC1 activity of bc1 complex like (S. pombe) (CABC1), mRNA.
hspc196	0.431	-2.323	transmembrane protein 138 (TMEM138), mRNA.
hypb	0.431	-2.323	huntingtin interacting protein B (HYPB), transcript variant 2, mRNA.
c17orf65	0.43	-2.323	chromosome 17 open reading frame 65 (C17orf65), mRNA.
chchd8	0.43	-2.323	coiled-coil-helix-coiled-coil-helix domain containing 8 (CHCHD8), mRNA.
foxq1	0.43	-2.323	forkhead box Q1 (FOXQ1), mRNA.
atp5o	0.43	-2.323	ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (oligomycin s ensitivity conferring protein) (ATP5O), nuclear gene encoding mitochondrial protein, mRNA.
elovl6	0.43	-2.323	ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) (ELOVL6), mRNA.
clip2	0.43	-2.324	CAP-GLY domain containing linker protein 2 (CLIP2), transcript variant 2, mRNA.
syf2	0.43	-2.324	SYF2 homolog, RNA splicing factor (S. cerevisiae) (SYF2), transcript variant 1, mRNA.
pank2	0.43	-2.324	pantothenate kinase 2 (Hallervorden-Spatz syndrome) (PANK2), transcript variant 5, mRNA.
stk36	0.43	-2.324	serine/threonine kinase 36, fused homolog (Drosophila) (STK36), mRNA.
sh3bgrl	0.43	-2.325	SH3 domain binding glutamic acid-rich protein like (SH3BGRL), mRNA.
commd7	0.43	-2.325	COMM domain containing 7 (COMMD7), mRNA.
drev1	0.43	-2.325	DORA reverse strand protein 1 (DREV1), mRNA.
neu1	0.43	-2.326	sialidase 1 (lysosomal sialidase) (NEU1), mRNA.
erp29	0.43	-2.326	endoplasmic reticulum protein 29 (ERP29), transcript variant 2, mRNA.
afg3l2	0.43	-2.326	AFG3 ATPase family gene 3-like 2 (yeast) (AFG3L2), nuclear gene encoding mitochondrial protein, mRNA.
giyd1	0.43	-2.327	GIY-YIG domain containing 1 (GIYD1), transcript variant 1, mRNA.
irf2bp2	0.43	-2.327	interferon regulatory factor 2 binding protein 2 (IRF2BP2), mRNA.
rnu14	0.43	-2.327	small nucleolar RNA, C/D box 14A (SNORD14A) on chromosome 11.
sod2	0.43	-2.327	superoxide dismutase 2, mitochondrial (SOD2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
slc4a5	0.43	-2.327	solute carrier family 4, sodium bicarbonate cotransporter, member 5 (SLC4A5), transcript variant d, mRNA.
c5orf5	0.43	-2.328	chromosome 5 open reading frame 5 (C5orf5), mRNA.
kiaa1267	0.43	-2.328	KIAA1267 (KIAA1267), mRNA.
mrps35	0.43	-2.328	mitochondrial ribosomal protein S35 (MRPS35), nuclear gene encoding mitochondrial protein, mRNA.
coro1b	0.43	-2.328	coronin, actin binding protein, 1B (CORO1B), transcript variant 1, mRNA.
gmppa	0.429	-2.328	GDP-mannose pyrophosphorylase A (GMPPA), transcript variant 2, mRNA.
fnbp1	0.429	-2.329	formin binding protein 1 (FNBP1), mRNA.
znf18	0.429	-2.329	zinc finger protein 18 (ZNF18), mRNA.

dkfzp762e13	0.429	-2.329	hypothetical protein DKFZp762E1312 (DKFZp762E1312), mRNA.
s100a11	0.429	-2.329	S100 calcium binding protein A11 (S100A11), mRNA.
nfil3	0.429	-2.33	nuclear factor, interleukin 3 regulated (NFIL3), mRNA.
nfkb2	0.429	-2.33	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100) (NFKB2), transcript variant 1, mRNA.
mfn2	0.429	-2.33	mitofusin 2 (MFN2), nuclear gene encoding mitochondrial protein, mRNA.
tcta	0.429	-2.33	T-cell leukemia translocation altered gene (TCTA), mRNA.
arf1	0.429	-2.331	ADP-ribosylation factor 1 (ARF1), transcript variant 3, mRNA.
tlcd1	0.429	-2.331	TLC domain containing 1 (TLCD1), mRNA.
c20orf116	0.429	-2.331	chromosome 20 open reading frame 116 (C20orf116), mRNA.
hsd17b4	0.429	-2.331	hydroxysteroid (17-beta) dehydrogenase 4 (HSD17B4), mRNA.
atpaf1	0.429	-2.331	ATP synthase mitochondrial F1 complex assembly factor 1 (ATPAF1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
flj21127	0.429	-2.331	tectonic (FLJ21127), mRNA.
ei24	0.429	-2.332	etoposide induced 2.4 mRNA (EI24), transcript variant 1, mRNA.
comtd1	0.429	-2.332	catechol-O-methyltransferase domain containing 1 (COMTD1), mRNA.
mcm2	0.429	-2.332	MCM2 minichromosome maintenance deficient 2, mitotin (S. cerevisiae) (MCM2), mRNA.
pop7	0.429	-2.333	processing of precursor 7, ribonuclease P subunit (S. cerevisiae) (POP7), mRNA.
c14orf78	0.429	-2.333	PREDICTED: chromosome 14 open reading frame 78, transcript variant 1 (C14orf78), mRNA.
mmp24	0.429	-2.333	matrix metallopeptidase 24 (membrane-inserted) (MMP24), mRNA.
tspan33	0.429	-2.333	tetraspanin 33 (TSPAN33), mRNA.
mgc72104	0.429	-2.333	similar to FRG1 protein (FSHD region gene 1 protein) (MGC72104), mRNA.
mrfap1l1	0.429	-2.334	Morf4 family associated protein 1-like 1 (MRFAP1L1), transcript variant 1, mRNA.
rab22a	0.428	-2.334	RAB22A, member RAS oncogene family (RAB22A), mRNA.
loc653879	0.428	-2.334	PREDICTED: similar to Complement C3 precursor (LOC653879), mRNA.
bet1	0.428	-2.334	BET1 homolog (S. cerevisiae) (BET1), mRNA.
osbpl10	0.428	-2.335	oxysterol binding protein-like 10 (OSBPL10), mRNA.
tiprl	0.428	-2.335	TIP41, TOR signalling pathway regulator-like (S. cerevisiae) (TIPRL), transcript variant 1, mRNA.
hs.143018	0.428	-2.335	BX105338 Soares_pregnant_uterus_NbHPU cDNA clone IMAGp998C114347, mRNA sequence
pes1	0.428	-2.335	pescadillo homolog 1, containing BRCT domain (zebrafish) (PES1), mRNA.
apex2	0.428	-2.335	APEX nuclease (apurinic/apyrimidinic endonuclease) 2 (APEX2), nuclear gene encoding mitochondrial protein, mRNA.
dennd4c	0.428	-2.335	DENN/MADD domain containing 4C (DENND4C), mRNA.
polr3k	0.428	-2.336	polymerase (RNA) III (DNA directed) polypeptide K, 12.3 kDa (POLR3K), mRNA.
gmfb	0.428	-2.336	glia maturation factor, beta (GMFB), mRNA.
ftsj3	0.428	-2.336	FtsJ homolog 3 (E. coli) (FTSJ3), mRNA.
fars2	0.428	-2.336	phenylalanine-tRNA synthetase 2 (mitochondrial) (FARS2), nuclear gene encoding mitochondrial protein, mRNA.
ass	0.428	-2.336	argininosuccinate synthetase (ASS), transcript variant 2, mRNA.
wdr62	0.428	-2.337	WD repeat domain 62 (WDR62), mRNA.
oxsm	0.428	-2.337	3-oxoacyl-ACP synthase, mitochondrial (OXSM), mRNA.
kcnn4	0.428	-2.337	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4 (KCNN4), mRNA.
kidins220	0.428	-2.337	kinase D-interacting substrate of 220 kDa (KIDINS220), mRNA.

mtx3	0.428	-2.338	metaxin 3 (MTX3), mRNA.
mrps31	0.428	-2.338	mitochondrial ribosomal protein S31 (MRPS31), nuclear gene encoding mitochondrial protein, mRNA.
mgc3121	0.428	-2.338	proline rich 14 (PRR14), mRNA.
spata18	0.428	-2.338	spermatogenesis associated 18 homolog (rat) (SPATA18), mRNA.
pcsk1n	0.428	-2.338	proprotein convertase subtilisin/kexin type 1 inhibitor (PCSK1N), mRNA.
timm8b	0.428	-2.338	translocase of inner mitochondrial membrane 8 homolog B (yeast) (TIMM8B), mRNA.
ugcgl2	0.428	-2.338	UDP-glucose ceramide glucosyltransferase-like 2 (UGCGL2), mRNA.
serpinb1	0.428	-2.338	serpin peptidase inhibitor, clade B (ovalbumin), member 1 (SERPINB1), mRNA.
kiaa0391	0.428	-2.338	KIAA0391 (KIAA0391), mRNA.
h1fx	0.428	-2.339	H1 histone family, member X (H1FX), mRNA.
neto2	0.428	-2.339	neuropilin (NRP) and tolloid (TLL)-like 2 (NETO2), mRNA.
plp2	0.427	-2.339	proteolipid protein 2 (colonic epithelium-enriched) (PLP2), mRNA.
kiaa1468	0.427	-2.34	KIAA1468 (KIAA1468), mRNA.
rnf31	0.427	-2.34	ring finger protein 31 (RNF31), mRNA.
mpv17	0.427	-2.34	MpV17 transgene, murine homolog, glomerulosclerosis (MPV17), mRNA.
ric8a	0.427	-2.34	resistance to inhibitors of cholinesterase 8 homolog A (C. elegans) (RIC8A), mRNA.
rrbp1	0.427	-2.341	ribosome binding protein 1 homolog 180kDa (dog) (RRBP1), transcript variant 1, mRNA.
polr2b	0.427	-2.342	polymerase (RNA) II (DNA directed) polypeptide B, 140kDa (POLR2B), mRNA.
hint2	0.427	-2.342	histidine triad nucleotide binding protein 2 (HINT2), mRNA.
mapk12	0.427	-2.342	mitogen-activated protein kinase 12 (MAPK12), mRNA.
pold4	0.427	-2.342	polymerase (DNA-directed), delta 4 (POLD4), mRNA.
spag7	0.427	-2.343	sperm associated antigen 7 (SPAG7), mRNA.
cand1	0.427	-2.343	cullin-associated and neddylation-dissociated 1 (CAND1), mRNA.
ganab	0.427	-2.343	glucosidase, alpha; neutral AB (GANAB), transcript variant 2, mRNA.
angel2	0.427	-2.343	angel homolog 2 (Drosophila) (ANGEL2), mRNA.
tfpi	0.427	-2.344	tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) (TFPI), transcript variant 1, mRNA.
rpl31	0.427	-2.344	ribosomal protein L31 (RPL31), mRNA.
mki67	0.426	-2.345	antigen identified by monoclonal antibody Ki-67 (MKI67), mRNA.
rbx1	0.426	-2.345	ring-box 1 (RBX1), mRNA.
rragc	0.426	-2.345	Ras-related GTP binding C (RRAGC), mRNA.
trmt1	0.426	-2.345	TRM1 tRNA methyltransferase 1 homolog (S. cerevisiae) (TRMT1), mRNA.
exosc8	0.426	-2.346	exosome component 8 (EXOSC8), mRNA.
5-Mar	0.426	-2.346	membrane-associated ring finger (C3HC4) 5 (MARCH5), mRNA.
akap13	0.426	-2.346	A kinase (PRKA) anchor protein 13 (AKAP13), transcript variant 2, mRNA.
tm9sf1	0.426	-2.347	transmembrane 9 superfamily member 1 (TM9SF1), transcript variant 2, mRNA.
nudt16l1	0.426	-2.347	nudix (nucleoside diphosphate linked moiety X)-type motif 16-like 1 (NUDT16L1), mRNA.
loc113386	0.426	-2.347	similar to envelope protein (LOC113386), mRNA.
rps20	0.426	-2.347	ribosomal protein S20 (RPS20), mRNA.
myo1e	0.426	-2.347	myosin IE (MYO1E), mRNA.
jmjd1c	0.426	-2.348	jumonji domain containing 1C (JMJD1C), transcript variant 1, mRNA.
scc-112	0.426	-2.349	PDS5, regulator of cohesion maintenance, homolog A (S. cerevisiae) (PDS5A), mRNA.
rps18	0.426	-2.349	ribosomal protein S18 (RPS18), mRNA.

plod3	0.426	-2.349	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3 (PLOD3), mRNA.
avpi1	0.426	-2.349	arginine vasopressin-induced 1 (AVPI1), mRNA.
usp14	0.426	-2.349	ubiquitin specific peptidase 14 (tRNA-guanine transglycosylase) (USP14), transcript variant 2, mRNA.
u2af1	0.426	-2.349	U2 small nuclear RNA auxiliary factor 1 (U2AF1), transcript variant b, mRNA.
cdk5rap2	0.426	-2.35	CDK5 regulatory subunit associated protein 2 (CDK5RAP2), transcript variant 1, mRNA.
ift20	0.426	-2.35	intraflagellar transport 20 homolog (Chlamydomonas) (IFT20), mRNA.
glb1	0.426	-2.35	galactosidase, beta 1 (GLB1), transcript variant 179423, mRNA.
pea15	0.426	-2.35	phosphoprotein enriched in astrocytes 15 (PEA15), mRNA.
smarcd3	0.426	-2.35	SWI/SNF related, matrix associated, actin dependent regulator of chromatin , subfamily d, member 3 (SMARCD3), transcript variant 2, mRNA.
mnd1	0.425	-2.35	meiotic nuclear divisions 1 homolog (S. cerevisiae) (MND1), mRNA.
id3	0.425	-2.351	inhibitor of DNA binding 3, dominant negative helix-loop-helix protein (ID3), mRNA.
cdc25b	0.425	-2.351	cell division cycle 25B (CDC25B), transcript variant 4, mRNA.
smo	0.425	-2.351	smoothened homolog (Drosophila) (SMO), mRNA.
tmem150	0.425	-2.351	transmembrane protein 150 (TMEM150), transcript variant 1, mRNA.
cuta	0.425	-2.352	cutA divalent cation tolerance homolog (E. coli) (CUTA), transcript variant 2, mRNA.
iqwd1	0.425	-2.353	IQ motif and WD repeats 1 (IQWD1), transcript variant 2, mRNA.
kiaa1840	0.425	-2.353	KIAA1840 (KIAA1840), mRNA.
faf1	0.425	-2.353	Fas (TNFRSF6) associated factor 1 (FAF1), mRNA.
copg2	0.425	-2.354	coatomer protein complex, subunit gamma 2 (COPG2), mRNA.
pla2g4c	0.425	-2.354	phospholipase A2, group IVC (cytosolic, calcium-independent) (PLA2G4C), mRNA.
lap3	0.425	-2.354	leucine aminopeptidase 3 (LAP3), mRNA.
hat1	0.425	-2.354	histone acetyltransferase 1 (HAT1), transcript variant 1, mRNA.
b3gnt6	0.425	-2.355	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 6 (B3GNT6), mRNA.
loc653219	0.425	-2.355	PREDICTED: similar to G antigen, family D, 2 isoform 1a, transcript variant 2 (LOC653219), mRNA.
acat1	0.425	-2.355	acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase) (ACAT1) , nuclear gene encoding mitochondrial protein, mRNA.
m-rip	0.424	-2.356	myosin phosphatase-Rho interacting protein (M-RIP), transcript variant 2, mRNA.
nhs	0.424	-2.356	PREDICTED: Nance-Horan syndrome (congenital cataracts and dental anomalies), transcript variant 3 (NHS), mRNA.
cdc2l5	0.424	-2.357	cell division cycle 2-like 5 (cholinesterase-related cell division controller) (CDC2L5), transcript variant 1, mRNA.
sdfr1	0.424	-2.357	stromal cell derived factor receptor 1 (SDFR1), transcript variant beta, mRNA.
pigs	0.424	-2.359	phosphatidylinositol glycan anchor biosynthesis, class S (PIGS), mRNA.
efna1	0.424	-2.359	ephrin-A1 (EFNA1), transcript variant 1, mRNA.
igf2bp2	0.424	-2.36	insulin-like growth factor 2 mRNA binding protein 2 (IGF2BP2), transcript variant 1, mRNA.
lipa	0.424	-2.36	lipase A, lysosomal acid, cholesterol esterase (Wolman disease) (LIPA), mRNA.
hoxb7	0.424	-2.361	homeo box B7 (HOXB7), mRNA.
loc644850	0.424	-2.361	PREDICTED: similar to phosducin-like 3 (LOC644850), mRNA.
cyb5d2	0.423	-2.362	cytochrome b5 domain containing 2 (CYB5D2), mRNA.
phkg2	0.423	-2.362	phosphorylase kinase, gamma 2 (testis) (PHKG2), mRNA.
timm8a	0.423	-2.362	translocase of inner mitochondrial membrane 8 homolog A (yeast) (TIMM8A), nuclear gene encoding mitochondrial protein, mRNA.
c12orf5	0.423	-2.363	chromosome 12 open reading frame 5 (C12orf5), mRNA.

cox6c	0.423	-2.363	cytochrome c oxidase subunit VIc (COX6C), mRNA.
pmm1	0.423	-2.363	phosphomannomutase 1 (PMM1), mRNA.
dkfzp564b14	0.423	-2.364	PREDICTED: DKFZP564B147 protein, transcript variant 10 (DKFZP564B147), misc RNA.
c6orf173	0.423	-2.364	chromosome 6 open reading frame 173 (C6orf173), mRNA.
loc643516	0.423	-2.364	PREDICTED: similar to 40S ribosomal protein S26 (LOC643516), mRNA.
isg20l2	0.423	-2.364	interferon stimulated exonuclease gene 20kDa-like 2 (ISG20L2), mRNA.
sco2	0.423	-2.364	SCO cytochrome oxidase deficient homolog 2 (yeast) (SCO2), nuclear gene encoding mitochondrial protein, mRNA
nap1l1	0.423	-2.364	nucleosome assembly protein 1-like 1 (NAP1L1), transcript variant 1, mRNA.
ythdf1	0.423	-2.364	YTH domain family, member 1 (YTHDF1), mRNA.
cdca5	0.423	-2.364	cell division cycle associated 5 (CDCA5), mRNA.
rnf167	0.423	-2.365	ring finger protein 167 (RNF167), mRNA.
surf1	0.423	-2.365	surfeit 1 (SURF1), nuclear gene encoding mitochondrial protein, mRNA.
casp6	0.423	-2.366	caspase 6, apoptosis-related cysteine peptidase (CASP6), transcript variant beta, mRNA.
hla-dma	0.423	-2.366	major histocompatibility complex, class II, DM alpha (HLA-DMA), mRNA.
n-pac	0.423	-2.366	cytokine-like nuclear factor n-pac (N-PAC), mRNA.
txlna	0.423	-2.366	taxilin alpha (TXLNA), mRNA.
scpep1	0.423	-2.366	serine carboxypeptidase 1 (SCPEP1), mRNA.
ptov1	0.423	-2.366	prostate tumor overexpressed gene 1 (PTOV1), mRNA.
tmem91	0.423	-2.367	transmembrane protein 91 (TMEM91), mRNA.
ascc3l1	0.422	-2.367	activating signal cointegrator 1 complex subunit 3-like 1 (ASCC3L1), mRNA.
msto1	0.422	-2.368	misato homolog 1 (Drosophila) (MSTO1), mRNA.
c12orf29	0.422	-2.369	chromosome 12 open reading frame 29 (C12orf29), mRNA.
bzw2	0.422	-2.369	basic leucine zipper and W2 domains 2 (BZW2), mRNA.
crip2	0.422	-2.369	cysteine-rich protein 2 (CRIP2), mRNA.
fbxo30	0.422	-2.369	F-box protein 30 (FBXO30), mRNA.
bcap31	0.422	-2.37	B-cell receptor-associated protein 31 (BCAP31), mRNA.
p8	0.422	-2.37	p8 protein (candidate of metastasis 1) (P8), mRNA.
gatad2b	0.422	-2.37	GATA zinc finger domain containing 2B (GATAD2B), mRNA.
nalp2	0.422	-2.37	NLR family, pyrin domain containing 2 (NLRP2), mRNA.
znf480	0.422	-2.371	zinc finger protein 480 (ZNF480), mRNA.
xylb	0.422	-2.371	xylulokinase homolog (H. influenzae) (XYLB), mRNA.
sub1	0.422	-2.371	SUB1 homolog (S. cerevisiae) (SUB1), mRNA.
trak2	0.422	-2.372	trafficking protein, kinesin binding 2 (TRAK2), mRNA.
mgc16597	0.422	-2.372	PREDICTED: similar to RIKEN cDNA 3110023B02, transcript variant 4 (MGC16597), mRNA.
abtb2	0.422	-2.372	ankyrin repeat and BTB (POZ) domain containing 2 (ABTB2), mRNA.
tgoln2	0.422	-2.372	trans-golgi network protein 2 (TGOLN2), mRNA.
pygb	0.422	-2.372	phosphorylase, glycogen; brain (PYGB), mRNA.
rad50	0.421	-2.373	RAD50 homolog (S. cerevisiae) (RAD50), transcript variant 1, mRNA.
ddx55	0.421	-2.373	DEAD (Asp-Glu-Ala-Asp) box polypeptide 55 (DDX55), mRNA.
c14orf94	0.421	-2.373	chromosome 14 open reading frame 94 (C14orf94), mRNA.
loc653778	0.421	-2.374	PREDICTED: similar to solute carrier family 25, member 37 (LOC653778), mRNA.
gne	0.421	-2.375	glucosamine (UDP-N-acetyl)-2-epimerase/N-acetylmannosamine kinase (GNE), mRNA.

arfgap1	0.421	-2.375	ADP-ribosylation factor GTPase activating protein 1 (ARFGAP1), transcript variant 2, mRNA.
gbe1	0.421	-2.375	glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV) (GBE1), mRNA.
eif2ak2	0.421	-2.375	eukaryotic translation initiation factor 2-alpha kinase 2 (EIF2AK2), mRNA.
yeats4	0.421	-2.375	YEATS domain containing 4 (YEATS4), mRNA.
fam89b	0.421	-2.375	family with sequence similarity 89, member B (FAM89B), mRNA.
chd1	0.421	-2.375	chromodomain helicase DNA binding protein 1 (CHD1), mRNA.
edem1	0.421	-2.375	ER degradation enhancer, mannosidase alpha-like 1 (EDEM1), mRNA.
etf1	0.421	-2.375	eukaryotic translation termination factor 1 (ETF1), mRNA.
gna11	0.421	-2.376	guanine nucleotide binding protein (G protein), alpha 11 (Gq class) (GNA11), mRNA.
tial1	0.421	-2.377	TIA1 cytotoxic granule-associated RNA binding protein-like 1 (TIAL1), transcript variant 1, mRNA. XM_945560 XM_945564 XM_945566 XM_945569 XM_945570 XM_945572 XM_945573
fto	0.421	-2.377	fat mass and obesity associated (FTO), mRNA.
ctsb	0.421	-2.377	cathepsin B (CTSB), transcript variant 1, mRNA.
pigh	0.421	-2.377	phosphatidylinositol glycan, class H (PIGH), mRNA.
park7	0.421	-2.377	Parkinson disease (autosomal recessive, early onset) 7 (PARK7), mRNA.
mett10d	0.421	-2.377	methyltransferase 10 domain containing (METT10D), mRNA.
nrbp1	0.421	-2.378	nuclear receptor binding protein 1 (NRBP1), mRNA.
znf462	0.421	-2.378	zinc finger protein 462 (ZNF462), mRNA.
ndufa7	0.421	-2.378	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7, 14.5kDa (NDUFA7), mRNA.
etnk1	0.421	-2.378	ethanolamine kinase 1 (ETNK1), mRNA.
aldh7a1	0.42	-2.378	PREDICTED: aldehyde dehydrogenase 7 family, member A1 (ALDH7A1), mRNA.
mlf2	0.42	-2.379	myeloid leukemia factor 2 (MLF2), mRNA.
polr3gl	0.42	-2.379	polymerase (RNA) III (DNA directed) polypeptide G (32kD)-like (POLR3GL), mRNA.
herc2	0.42	-2.38	hect domain and RLD 2 (HERC2), mRNA.
manbal	0.42	-2.38	mannosidase, beta A, lysosomal-like (MANBAL), transcript variant 2, mRNA.
yipf3	0.42	-2.381	Yip1 domain family, member 3 (YIPF3), mRNA.
ngfrap1	0.42	-2.381	nerve growth factor receptor (TNFRSF16) associated protein 1 (NGFRAP1), transcript variant 1, mRNA.
birc5	0.42	-2.382	baculoviral IAP repeat-containing 5 (survivin) (BIRC5), transcript variant 3, mRNA.
ift74	0.42	-2.382	intraflagellar transport 74 homolog (Chlamydomonas) (IFT74), mRNA.
yes1	0.42	-2.383	v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1 (YES1), mRNA.
znf271	0.42	-2.383	zinc finger protein 271 (ZNF271), mRNA.
zdhhc6	0.42	-2.383	zinc finger, DHHC-type containing 6 (ZDHHC6), mRNA.
foxd1	0.419	-2.384	forkhead box D1 (FOXD1), mRNA.
th1l	0.419	-2.384	TH1-like (Drosophila) (TH1L), transcript variant 2, mRNA.
c13orf7	0.419	-2.384	chromosome 13 open reading frame 7 (C13orf7), mRNA.
gpx1	0.419	-2.384	glutathione peroxidase 1 (GPX1), transcript variant 2, mRNA.
slc35b2	0.419	-2.385	solute carrier family 35, member B2 (SLC35B2), mRNA.
bat2	0.419	-2.385	HLA-B associated transcript 2 (BAT2), transcript variant 1, mRNA.
c20orf191	0.419	-2.386	chromosome 20 open reading frame 191 (C20orf191), mRNA.
cuedc2	0.419	-2.386	CUE domain containing 2 (CUEDC2), mRNA.
mlr2	0.419	-2.386	ligand-dependent corepressor (MLR2), mRNA.

ptms	0.419	-2.387	parathymosin (PTMS), mRNA.
chst12	0.419	-2.387	carbohydrate (chondroitin 4) sulfotransferase 12 (CHST12), mRNA.
gcsh	0.419	-2.387	glycine cleavage system protein H (aminomethyl carrier) (GCSH), mRNA.
loc648638	0.419	-2.388	PREDICTED: similar to Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 (Rotamase Pin4) (PPIase Pin4) (Parvulin 14) (Par14) (Peptidyl-prolyl cis/trans isomerase EPVH) (hPar14) (LOC648638), mRNA.
snhg5	0.419	-2.389	small nucleolar RNA host gene (non-protein coding) 5 (SNHG5) on chromosome 6.
loc285636	0.419	-2.389	hypothetical protein LOC285636 (LOC285636), mRNA.
cybasc3	0.418	-2.391	cytochrome b, ascorbate dependent 3 (CYBASC3), mRNA.
dkfzp686k16	0.418	-2.391	similar to BMP2 inducible kinase (DKFZp686K16132), mRNA.
znf22	0.418	-2.392	zinc finger protein 22 (KOX 15) (ZNF22), mRNA.
gstk1	0.418	-2.392	glutathione S-transferase kappa 1 (GSTK1), mRNA.
lamc2	0.418	-2.392	laminin, gamma 2 (LAMC2), transcript variant 1, mRNA.
hs.440088	0.418	-2.393	BX393727 NEUROBLASTOMA COT 25-NORMALIZED Homo sapiens cDNA clone CS0DC001YP02 5-PRIME, mRNA sequence
znf511	0.418	-2.393	zinc finger protein 511 (ZNF511), mRNA.
c1rl	0.418	-2.393	complement component 1, r subcomponent-like (C1RL), mRNA.
hs.193767	0.418	-2.393	cDNA FLJ26188 fis, clone ADG04821
eif2s1	0.418	-2.394	eukaryotic translation initiation factor 2, subunit 1 alpha, 35kDa (EIF2S1), mRNA.
leprel1	0.418	-2.394	leprecan-like 1 (LEPREL1), mRNA.
hcap-h2	0.418	-2.394	non-SMC condensin II complex, subunit H2 (NCAPH2), transcript variant 1, mRNA.
flj14466	0.418	-2.394	transmembrane protein 142A (TMEM142A), mRNA.
apeh	0.418	-2.395	N-acylaminoacyl-peptide hydrolase (APEH), mRNA.
hs.583806	0.418	-2.395	AGENCOURT_7908292 NIH_MGC_82 cDNA clone IMAGE:6102595 5, mRNA sequence
ubp1	0.418	-2.395	upstream binding protein 1 (LBP-1a) (UBP1), mRNA.
plcd3	0.417	-2.396	phospholipase C, delta 3 (PLCD3), mRNA.
nudcd3	0.417	-2.396	NudC domain containing 3 (NUDCD3), mRNA.
loc402694	0.417	-2.396	PREDICTED: similar to ribosomal protein L15 (LOC402694), mRNA.
vhl	0.417	-2.397	von Hippel-Lindau tumor suppressor (VHL), transcript variant 1, mRNA.
isca1	0.417	-2.397	iron-sulfur cluster assembly 1 homolog (S. cerevisiae) (ISCA1), mRNA.
loc388621	0.417	-2.397	PREDICTED: similar to ribosomal protein L21 isoform 1 (LOC388621), mRNA.
lzic	0.417	-2.397	leucine zipper and CTNNBIP1 domain containing (LZIC), mRNA.
ap2s1	0.417	-2.398	adaptor-related protein complex 2, sigma 1 subunit (AP2S1), transcript variant AP17, mRNA.
prkce	0.417	-2.399	protein kinase C, epsilon (PRKCE), mRNA.
rai17	0.417	-2.4	zinc finger, MIZ-type containing 1 (ZMIZ1), mRNA.
c5orf15	0.417	-2.4	chromosome 5 open reading frame 15 (C5orf15), mRNA.
ddx50	0.417	-2.4	DEAD (Asp-Glu-Ala-Asp) box polypeptide 50 (DDX50), mRNA.
c6orf160	0.417	-2.401	PREDICTED: chromosome 6 open reading frame 160, transcript variant 4 (C6orf160), mRNA.
c8orf52	0.417	-2.401	integrator complex subunit 8 (INTS8), mRNA.
mrps27	0.416	-2.401	mitochondrial ribosomal protein S27 (MRPS27), nuclear gene encoding mitochondrial protein, mRNA.
cnot8	0.416	-2.401	CCR4-NOT transcription complex, subunit 8 (CNOT8), mRNA.
flj21749	0.416	-2.402	hypothetical protein FLJ21749 (FLJ21749), mRNA.
tbl1x	0.416	-2.403	transducin (beta)-like 1X-linked (TBL1X), mRNA.

c9orf74	0.416	-2.403	ubiquitin related modifier 1 homolog (S. cerevisiae) (URM1), mRNA.
ubph	0.416	-2.403	similar to ubiquitin binding protein (UBPH), mRNA.
fahd2b	0.416	-2.403	fumarylacetoacetate hydrolase domain containing 2B (FAHD2B), mRNA.
zcd1	0.416	-2.403	zinc finger, CDGSH-type domain 1 (ZCD1), mRNA.
pgm2l1	0.416	-2.403	phosphoglucomutase 2-like 1 (PGM2L1), mRNA.
trappc1	0.416	-2.404	trafficking protein particle complex 1 (TRAPPC1), mRNA.
top3a	0.416	-2.404	topoisomerase (DNA) III alpha (TOP3A), mRNA.
fem1c	0.416	-2.404	fem-1 homolog c (C.elegans) (FEM1C), mRNA.
loc644969	0.416	-2.405	PREDICTED: similar to Ubiquinol-cytochrome c reductase complex 14 kDa protein (Complex III subunit VI) (QP-C) (LOC644969), mRNA.
ktn1	0.416	-2.405	kinectin 1 (kinesin receptor) (KTN1), mRNA.
thrap4	0.416	-2.405	thyroid hormone receptor associated protein 4 (THRAP4), mRNA.
plk2	0.416	-2.405	polo-like kinase 2 (Drosophila) (PLK2), mRNA.
csda	0.416	-2.405	cold shock domain protein A (CSDA), mRNA.
atp6v1g1	0.416	-2.405	ATPase, H+ transporting, lysosomal 13kDa, V1 subunit G isoform 1 (ATP6V1G1), mRNA.
smc4	0.416	-2.406	structural maintenance of chromosomes 4 (SMC4), transcript variant 2, mRNA.
susd1	0.416	-2.406	sushi domain containing 1 (SUSD1), mRNA.
dhrs4l2	0.416	-2.406	dehydrogenase/reductase (SDR family) member 4 like 2 (DHRS4L2), mRNA.
tor3a	0.416	-2.407	torsin family 3, member A (TOR3A), mRNA.
e2f2	0.415	-2.407	E2F transcription factor 2 (E2F2), mRNA.
hs.513971	0.415	-2.407	cDNA FLJ34018 fis, clone FCBBF2002801
abcf1	0.415	-2.407	ATP-binding cassette, sub-family F (GCN20), member 1 (ABCF1), transcript variant 2, mRNA.
slc39a14	0.415	-2.407	solute carrier family 39 (zinc transporter), member 14 (SLC39A14), mRNA.
gss	0.415	-2.408	glutathione synthetase (GSS), mRNA.
taf6l	0.415	-2.408	TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65kDa (TAF6L), mRNA.
loc286334	0.415	-2.409	PREDICTED: hypothetical protein LOC286334 (LOC286334), mRNA.
rnf20	0.415	-2.409	ring finger protein 20 (RNF20), mRNA.
als2	0.415	-2.409	amyotrophic lateral sclerosis 2 (juvenile) (ALS2), mRNA.
loc653171	0.415	-2.409	PREDICTED: similar to MAPK-interacting and spindle-stabilizing protein (LOC653171), mRNA.
gtf3c5	0.415	-2.41	general transcription factor IIIC, polypeptide 5, 63kDa (GTF3C5), mRNA.
c9orf156	0.415	-2.41	chromosome 9 open reading frame 156 (C9orf156), mRNA.
nsmce1	0.415	-2.41	non-SMC element 1 homolog (S. cerevisiae) (NSMCE1), mRNA.
pcnxl3	0.415	-2.41	PREDICTED: pecanex-like 3 (Drosophila), transcript variant 4 (PCNXL3), mRNA.
htra2	0.415	-2.411	HtrA serine peptidase 2 (HTRA2), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
pdk4	0.415	-2.411	pyruvate dehydrogenase kinase, isozyme 4 (PDK4), mRNA.
ncbp1	0.415	-2.411	nuclear cap binding protein subunit 1, 80kDa (NCBP1), mRNA.
nosip	0.415	-2.411	nitric oxide synthase interacting protein (NOSIP), mRNA.
unc45a	0.415	-2.412	unc-45 homolog A (C. elegans) (UNC45A), transcript variant 3, mRNA.
srgap1	0.415	-2.412	SLIT-ROBO Rho GTPase activating protein 1 (SRGAP1), mRNA.
rragb	0.415	-2.412	Ras-related GTP binding B (RRAGB), transcript variant RAGBs, mRNA.
wbp4	0.415	-2.412	WW domain binding protein 4 (formin binding protein 21) (WBP4), mRNA.
api5	0.415	-2.412	apoptosis inhibitor 5 (API5), mRNA.

rbaf600	0.414	-2.413	retinoblastoma-associated factor 600 (RBAF600), mRNA.
phyh	0.414	-2.413	phytanoyl-CoA hydroxylase (Refsum disease) (PHYH), mRNA.
kiaa2013	0.414	-2.414	KIAA2013 (KIAA2013), mRNA.
acot9	0.414	-2.414	acyl-CoA thioesterase 9 (ACOT9), mRNA.
flj34969	0.414	-2.415	PREDICTED: hypothetical protein FLJ34969 (FLJ34969), mRNA.
spen	0.414	-2.415	spen homolog, transcriptional regulator (Drosophila) (SPEN), mRNA.
mrpl17	0.414	-2.415	mitochondrial ribosomal protein L17 (MRPL17), nuclear gene encoding mitochondrial protein, mRNA.
cdk5rap1	0.414	-2.416	CDK5 regulatory subunit associated protein 1 (CDK5RAP1), transcript variant 1, mRNA.
baat	0.414	-2.416	bile acid Coenzyme A: amino acid N-acyltransferase (glycine N-choloyltransferase) (BAAT), mRNA.
c20orf14	0.414	-2.417	chromosome 20 open reading frame 14 (C20orf14), mRNA.
bfsp1	0.414	-2.418	beaded filament structural protein 1, filensin (BFSP1), mRNA.
atp6ap1	0.414	-2.418	ATPase, H+ transporting, lysosomal accessory protein 1 (ATP6AP1), mRNA.
vcl	0.414	-2.418	vinculin (VCL), transcript variant 1, mRNA.
abcc2	0.414	-2.418	ATP-binding cassette, sub-family C (CFTR/MRP), member 2 (ABCC2), mRNA.
loc652500	0.413	-2.419	PREDICTED: similar to butyrate-induced transcript 1 (LOC652500), mRNA.
znf266	0.413	-2.419	zinc finger protein 266 (ZNF266), mRNA.
dyrk1a	0.413	-2.419	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A (DYRK1A), transcript variant 2, mRNA.
rab11fip2	0.413	-2.419	RAB11 family interacting protein 2 (class I) (RAB11FIP2), mRNA.
mt2a	0.413	-2.419	metallothionein 2A (MT2A), mRNA.
mrpl41	0.413	-2.419	mitochondrial ribosomal protein L41 (MRPL41), nuclear gene encoding mitochondrial protein, mRNA.
pigg	0.413	-2.42	phosphatidylinositol glycan anchor biosynthesis, class G (PIGG), mRNA.
eif4b	0.413	-2.42	eukaryotic translation initiation factor 4B (EIF4B), mRNA.
c18orf37	0.413	-2.42	chromosome 18 open reading frame 37 (C18orf37), mRNA.
galntl4	0.413	-2.42	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 4 (GALNTL4), mRNA.
cri2	0.413	-2.42	EP300 interacting inhibitor of differentiation 2 (EID2), mRNA.
flj10213	0.413	-2.421	hypothetical protein FLJ10213 (FLJ10213), mRNA.
thbd	0.413	-2.421	thrombomodulin (THBD), mRNA.
c11orf48	0.413	-2.422	chromosome 11 open reading frame 48 (C11orf48), mRNA.
hexa	0.413	-2.422	hexosaminidase A (alpha polypeptide) (HEXA), mRNA.
tmem16f	0.413	-2.423	transmembrane protein 16F (TMEM16F), mRNA.
loc221143	0.413	-2.423	N-6 adenine-specific DNA methyltransferase 2 (putative) (N6AMT2), mRNA.
nagpa	0.413	-2.423	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase (NAGPA), mRNA.
hnrpa1l-2	0.413	-2.424	heterogeneous nuclear ribonucleoprotein A1 pseudogene (HNRPA1L-2) on chromosome 19.
igbp1	0.413	-2.424	immunoglobulin (CD79A) binding protein 1 (IGBP1), mRNA.
fcgrt	0.413	-2.424	Fc fragment of IgG, receptor, transporter, alpha (FCGRT), mRNA.
cib2	0.413	-2.424	calcium and integrin binding family member 2 (CIB2), mRNA.
mrps23	0.412	-2.425	mitochondrial ribosomal protein S23 (MRPS23), nuclear gene encoding mitochondrial protein, mRNA.
fhod1	0.412	-2.425	formin homology 2 domain containing 1 (FHOD1), mRNA.
serpina3	0.412	-2.426	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 3 (SERPINA3), mRNA.
dpp7	0.412	-2.426	dipeptidyl-peptidase 7 (DPP7), mRNA.
sp1	0.412	-2.426	Sp1 transcription factor (SP1), mRNA.
acox1	0.412	-2.426	acyl-Coenzyme A oxidase 1, palmitoyl (ACOX1), transcript variant 1, mRNA.

c20orf121	0.412	-2.427	chromosome 20 open reading frame 121 (C20orf121), mRNA.
rhbdd2	0.412	-2.427	rhomboid domain containing 2 (RHBDD2), mRNA.
nol11	0.412	-2.428	nucleolar protein 11 (NOL11), mRNA.
c5	0.412	-2.428	complement component 5 (C5), mRNA.
mrps2	0.412	-2.429	mitochondrial ribosomal protein S2 (MRPS2), nuclear gene encoding mitochondrial protein, mRNA.
plce1	0.412	-2.429	phospholipase C, epsilon 1 (PLCE1), mRNA.
c15orf39	0.412	-2.429	chromosome 15 open reading frame 39 (C15orf39), mRNA.
scn1b	0.412	-2.43	sodium channel, voltage-gated, type I, beta (SCN1B), transcript variant b, mRNA.
rpl13	0.411	-2.43	ribosomal protein L13 (RPL13), transcript variant 2, mRNA.
c8orf35	0.411	-2.431	chromosome 8 open reading frame 35 (C8orf35), mRNA.
loc643031	0.411	-2.432	PREDICTED: similar to NADH dehydrogenase subunit 5 (LOC643031), mRNA.
ergic3	0.411	-2.433	ERGIC and golgi 3 (ERGIC3), transcript variant 1, mRNA.
ny-sar-48	0.411	-2.433	sarcoma antigen NY-SAR-48 (NY-SAR-48), transcript variant 2, mRNA.
mrpl48	0.411	-2.433	mitochondrial ribosomal protein L48 (MRPL48), nuclear gene encoding mitochondrial protein, mRNA.
ccdc14	0.411	-2.434	coiled-coil domain containing 14 (CCDC14), mRNA.
stat2	0.411	-2.434	signal transducer and activator of transcription 2, 113kDa (STAT2), mRNA.
mrpl40	0.411	-2.434	mitochondrial ribosomal protein L40 (MRPL40), nuclear gene encoding mitochondrial protein, mRNA.
mtrf1l	0.411	-2.434	mitochondrial translational release factor 1-like (MTRF1L), mRNA.
znf529	0.411	-2.434	zinc finger protein 529 (ZNF529), mRNA.
wrb	0.411	-2.434	tryptophan rich basic protein (WRB), mRNA.
atp5l	0.411	-2.435	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit G (ATP5L) , nuclear gene encoding mitochondrial protein, mRNA.
flot1	0.411	-2.436	flotillin 1 (FLOT1), mRNA.
gpc1	0.41	-2.437	glypican 1 (GPC1), mRNA.
peg10	0.41	-2.437	paternally expressed 10 (PEG10), transcript variant 1, mRNA. XM_940378
hs.508682	0.41	-2.437	AV762101 MDS cDNA clone MDSEOA03 5, mRNA sequence
psmd6	0.41	-2.438	proteasome (prosome, macropain) 26S subunit, non-ATPase, 6 (PSMD6), mRNA.
ppm1h	0.41	-2.438	PREDICTED: protein phosphatase 1H (PP2C domain containing) (PPM1H), mRNA.
hs.57079	0.41	-2.438	cDNA FLJ13267 fis, clone OVARC1000964
btbd10	0.41	-2.438	BTB (POZ) domain containing 10 (BTBD10), mRNA.
vim	0.41	-2.439	vimentin (VIM), mRNA.
rassf2	0.41	-2.439	Ras association (RalGDS/AF-6) domain family 2 (RASSF2), transcript variant 2, mRNA.
abhd4	0.41	-2.439	abhydrolase domain containing 4 (ABHD4), mRNA.
nxf1	0.41	-2.44	nuclear RNA export factor 1 (NXF1), mRNA.
loc644191	0.41	-2.44	PREDICTED: similar to 40S ribosomal protein S26, transcript variant 2 (LOC644191), mRNA.
p2ry11	0.41	-2.44	purinergic receptor P2Y, G-protein coupled, 11 (P2RY11), mRNA.
purb	0.41	-2.441	purine-rich element binding protein B (PURB), mRNA.
hspc152	0.41	-2.441	hypothetical protein HSPC152 (HSPC152), mRNA.
tmem4	0.41	-2.441	transmembrane protein 4 (TMEM4), mRNA.
c1orf166	0.41	-2.441	chromosome 1 open reading frame 166 (C1orf166), mRNA.
ring1	0.41	-2.441	ring finger protein 1 (RING1), mRNA.
akap8l	0.409	-2.442	A kinase (PRKA) anchor protein 8-like (AKAP8L), mRNA.

fancg	0.409	-2.442	Fanconi anemia, complementation group G (FANCG), mRNA.
cdk5rap3	0.409	-2.442	CDK5 regulatory subunit associated protein 3 (CDK5RAP3), mRNA.
pttg3	0.409	-2.443	pituitary tumor-transforming 3 (PTTG3) on chromosome 8.
taf10	0.409	-2.443	TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 30kDa (TAF10), mRNA.
ube3c	0.409	-2.444	ubiquitin protein ligase E3C (UBE3C), mRNA.
ddx10	0.409	-2.444	DEAD (Asp-Glu-Ala-Asp) box polypeptide 10 (DDX10), mRNA.
pbef1	0.409	-2.444	pre-B-cell colony enhancing factor 1 (PBEF1), transcript variant 2, mRNA.
stk38	0.409	-2.444	serine/threonine kinase 38 (STK38), mRNA.
hnrpdl	0.409	-2.445	heterogeneous nuclear ribonucleoprotein D-like (HNRPDL), transcript variant 1, mRNA.
mical2	0.409	-2.445	microtubule associated monoxygenase, calponin and LIM domain containing 2 (MICAL2), mRNA.
cyb5a	0.409	-2.445	cytochrome b5 type A (microsomal) (CYB5A), transcript variant 2, mRNA.
wdr79	0.409	-2.445	WD repeat domain 79 (WDR79), mRNA.
loc652672	0.409	-2.445	PREDICTED: similar to damage-specific DNA binding protein 1 (LOC652672), mRNA.
tspan17	0.409	-2.446	tetraspanin 17 (TSPAN17), transcript variant 1, mRNA.
fgfr4	0.409	-2.446	fibroblast growth factor receptor 4 (FGFR4), transcript variant 2, mRNA.
sld5	0.409	-2.446	GINS complex subunit 4 (Sld5 homolog) (GINS4), mRNA.
c1orf122	0.409	-2.446	chromosome 1 open reading frame 122 (C1orf122), mRNA.
actr10	0.409	-2.446	actin-related protein 10 homolog (S. cerevisiae) (ACTR10), mRNA.
tspan5	0.409	-2.446	tetraspanin 5 (TSPAN5), mRNA.
ptplad2	0.409	-2.447	protein tyrosine phosphatase-like A domain containing 2 (PTPLAD2), mRNA.
manba	0.409	-2.447	mannosidase, beta A, lysosomal (MANBA), mRNA.
nfe2l1	0.408	-2.448	nuclear factor (erythroid-derived 2)-like 1 (NFE2L1), mRNA.
col16a1	0.408	-2.448	collagen, type XVI, alpha 1 (COL16A1), mRNA.
irf3	0.408	-2.448	interferon regulatory factor 3 (IRF3), mRNA.
kiaa1423	0.408	-2.449	PREDICTED: KIAA1423 (KIAA1423), mRNA.
mgc16169	0.408	-2.449	hypothetical protein MGC16169 (MGC16169), mRNA.
pabpc1	0.408	-2.449	poly(A) binding protein, cytoplasmic 1 (PABPC1), mRNA.
c21orf57	0.408	-2.45	chromosome 21 open reading frame 57 (C21orf57), transcript variant 2, mRNA.
nme3	0.408	-2.45	non-metastatic cells 3, protein expressed in (NME3), mRNA.
vdp	0.408	-2.45	vesicle docking protein p115 (VDP), mRNA.
eif3s4	0.408	-2.451	eukaryotic translation initiation factor 3, subunit 4 delta, 44kDa (EIF3S4), mRNA.
snft	0.408	-2.451	Jun dimerization protein p21SNFT (SNFT), mRNA.
c3orf9	0.408	-2.451	KTEL (Lys-Tyr-Glu-Leu) containing 1 (KTELC1), mRNA.
nme1-nme2	0.408	-2.451	NM23-LV (NME1-NME2), mRNA.
sh2d4a	0.408	-2.452	SH2 domain containing 4A (SH2D4A), mRNA.
gle1l	0.408	-2.452	GLE1 RNA export mediator-like (yeast) (GLE1L), transcript variant 2, mRNA.
loc440348	0.408	-2.453	similar to nuclear pore complex interacting protein (LOC440348), mRNA.
flj25222	0.408	-2.453	CXYorf1-related protein (FLJ25222), mRNA.
tgfbrap1	0.408	-2.454	transforming growth factor, beta receptor associated protein 1 (TGFBRAP1), mRNA.
nufip2	0.407	-2.454	nuclear fragile X mental retardation protein interacting protein 2 (NUFIP2), mRNA.
tarbp2	0.407	-2.454	TAR (HIV-1) RNA binding protein 2 (TARBP2), transcript variant 1, mRNA.
tnnt1	0.407	-2.455	troponin T type 1 (skeletal, slow) (TNNT1), mRNA.

tspyl5	0.407	-2.455	TSPY-like 5 (TSPYL5), mRNA.
bcl2l1	0.407	-2.455	BCL2-like 1 (BCL2L1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
cda	0.407	-2.456	cytidine deaminase (CDA), mRNA.
cant1	0.407	-2.456	calcium activated nucleotidase 1 (CANT1), mRNA.
flj20186	0.407	-2.456	hypothetical protein FLJ20186 (FLJ20186), transcript variant 1, mRNA.
c7orf24	0.407	-2.456	chromosome 7 open reading frame 24 (C7orf24), mRNA.
tm2d1	0.407	-2.456	TM2 domain containing 1 (TM2D1), mRNA.
plekhb2	0.407	-2.456	pleckstrin homology domain containing, family B (evectins) member 2 (PLEKHB2), transcript variant 2, mRNA.
akap8	0.407	-2.457	A kinase (PRKA) anchor protein 8 (AKAP8), mRNA.
pus7	0.407	-2.457	pseudouridylate synthase 7 homolog (S. cerevisiae) (PUS7), mRNA.
prps2	0.407	-2.457	phosphoribosyl pyrophosphate synthetase 2 (PRPS2), transcript variant 1, mRNA.
surf4	0.407	-2.458	surfeit 4 (SURF4), mRNA.
ehd2	0.407	-2.459	EH-domain containing 2 (EHD2), mRNA.
clip1	0.407	-2.459	CAP-GLY domain containing linker protein 1 (CLIP1), transcript variant 1, mRNA.
pdcd5	0.407	-2.46	programmed cell death 5 (PDCD5), mRNA.
atp6v0a1	0.406	-2.461	ATPase, H+ transporting, lysosomal V0 subunit a isoform 1 (ATP6V0A1), mRNA.
hdcma18p	0.406	-2.462	HDCMA18P protein (HDCMA18P), mRNA.
cxcl5	0.406	-2.462	chemokine (C-X-C motif) ligand 5 (CXCL5), mRNA.
gemin6	0.406	-2.462	gem (nuclear organelle) associated protein 6 (GEMIN6), mRNA.
rnd1	0.406	-2.463	Rho family GTPase 1 (RND1), mRNA.
ndufab1	0.406	-2.464	NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1, 8kDa (NDUFAB1), mRNA.
p2rx4	0.406	-2.464	purinergic receptor P2X, ligand-gated ion channel, 4 (P2RX4), transcript variant 3, mRNA.
znhit1	0.406	-2.465	zinc finger, HIT type 1 (ZNHIT1), mRNA.
ssr4	0.406	-2.465	signal sequence receptor, delta (translocon-associated protein delta) (SSR4), mRNA.
samd1	0.406	-2.465	sterile alpha motif domain containing 1 (SAMD1), mRNA.
setx	0.406	-2.465	senataxin (SETX), mRNA.
smox	0.406	-2.466	spermine oxidase (SMOX), transcript variant 4, mRNA.
ptpla	0.405	-2.466	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a (PTPLA), mRNA.
lppr2	0.405	-2.467	lipid phosphate phosphatase-related protein type 2 (LPPR2), mRNA.
anapc13	0.405	-2.467	anaphase promoting complex subunit 13 (ANAPC13), mRNA.
fam45a	0.405	-2.467	family with sequence similarity 45, member A (FAM45A), mRNA.
ccdc43	0.405	-2.467	coiled-coil domain containing 43 (CCDC43), mRNA.
copg	0.405	-2.467	coatomer protein complex, subunit gamma (COPG), mRNA.
ssna1	0.405	-2.467	Sjogren's syndrome nuclear autoantigen 1 (SSNA1), mRNA.
hs.184721	0.405	-2.467	EST366269 MAGE resequences, MAGC cDNA, mRNA sequence
acp2	0.405	-2.468	acid phosphatase 2, lysosomal (ACP2), mRNA.
ddx56	0.405	-2.468	DEAD (Asp-Glu-Ala-Asp) box polypeptide 56 (DDX56), mRNA.
loc147808	0.405	-2.468	zinc finger protein 784 (ZNF784), mRNA.
c14orf28	0.405	-2.469	chromosome 14 open reading frame 28 (C14orf28), mRNA.
suclg2	0.405	-2.469	succinate-CoA ligase, GDP-forming, beta subunit (SUCLG2), mRNA.
aamp	0.405	-2.469	angio-associated, migratory cell protein (AAMP), mRNA.
sh3pxd2b	0.405	-2.469	SH3 and PX domains 2B (SH3PXD2B), mRNA.

zbed1	0.405	-2.469	zinc finger, BED-type containing 1 (ZBED1), mRNA.
ewsr1	0.405	-2.47	Ewing sarcoma breakpoint region 1 (EWSR1), transcript variant EWS, mRNA.
cdk7	0.405	-2.47	cyclin-dependent kinase 7 (MO15 homolog, Xenopus laevis, cdk-activating kinase) (CDK7), mRNA.
sf4	0.405	-2.47	splicing factor 4 (SF4), transcript variant c, mRNA.
atpif1	0.405	-2.471	ATPase inhibitory factor 1 (ATPIF1), nuclear gene encoding mitochondrial protein, transcript variant 3, mRNA.
ap1s1	0.405	-2.471	adaptor-related protein complex 1, sigma 1 subunit (AP1S1), transcript variant 1, mRNA.
scarb1	0.405	-2.472	scavenger receptor class B, member 1 (SCARB1), mRNA.
zdhhc24	0.404	-2.472	zinc finger, DHHC-type containing 24 (ZDHHC24), mRNA.
klp1	0.404	-2.472	N-acetyltransferase 14 (NAT14), mRNA.
cst3	0.404	-2.473	cystatin C (amyloid angiopathy and cerebral hemorrhage) (CST3), mRNA.
wdr13	0.404	-2.473	WD repeat domain 13 (WDR13), mRNA.
slc29a3	0.404	-2.474	solute carrier family 29 (nucleoside transporters), member 3 (SLC29A3), mRNA.
tmem54	0.404	-2.474	transmembrane protein 54 (TMEM54), mRNA.
loc643949	0.404	-2.474	PREDICTED: similar to 60S acidic ribosomal protein P2 (LOC643949), mRNA.
loc134147	0.404	-2.474	similar to mouse 2310016A09Rik gene (LOC134147), mRNA.
tmed9	0.404	-2.474	transmembrane emp24 protein transport domain containing 9 (TMED9), mRNA.
psap	0.404	-2.474	prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy) (PSAP), mRNA.
flj10081	0.404	-2.476	hypothetical protein FLJ10081 (FLJ10081), mRNA.
rbm6	0.404	-2.476	RNA binding motif protein 6 (RBM6), mRNA.
ipo9	0.404	-2.476	importin 9 (IPO9), mRNA.
znf532	0.404	-2.477	zinc finger protein 532 (ZNF532), mRNA.
ica1	0.404	-2.477	islet cell autoantigen 1, 69kDa (ICA1), transcript variant 2, mRNA.
zbtb4	0.404	-2.478	zinc finger and BTB domain containing 4 (ZBTB4), mRNA.
hs.121525	0.403	-2.479	BX101252 NCI_CGAP_Lu24 cDNA clone IMAGp998I115625, mRNA sequence
nxn	0.403	-2.479	nucleoredoxin (NXN), mRNA.
odc1	0.403	-2.479	ornithine decarboxylase 1 (ODC1), mRNA.
mfap1	0.403	-2.479	microfibrillar-associated protein 1 (MFAP1), mRNA.
znf223	0.403	-2.48	zinc finger protein 223 (ZNF223), mRNA.
mcm5	0.403	-2.48	MCM5 minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae) (MCM5), mRNA.
ntan1	0.403	-2.481	PREDICTED: N-terminal asparagine amidase, transcript variant 3 (NTAN1), mRNA.
pcoln3	0.403	-2.481	procollagen (type III) N-endopeptidase (PCOLN3), mRNA.
map3k11	0.403	-2.481	mitogen-activated protein kinase kinase kinase 11 (MAP3K11), mRNA.
mc1r	0.403	-2.481	melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor) (MC1R), mRNA.
hs.461819	0.403	-2.482	cDNA FLJ41112 fis, clone BRACE1000239
herc5	0.403	-2.482	hect domain and RLD 5 (HERC5), mRNA.
adck1	0.403	-2.483	aarF domain containing kinase 1 (ADCK1), mRNA.
rbm10	0.403	-2.483	RNA binding motif protein 10 (RBM10), transcript variant 1, mRNA.
mrpl1	0.403	-2.483	mitochondrial ribosomal protein L1 (MRPL1), nuclear gene encoding mitochondrial protein, mRNA.
ercc5	0.403	-2.484	excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome)) (ERCC5), mRNA.
sumo2	0.403	-2.484	SMT3 suppressor of mif two 3 homolog 2 (S. cerevisiae) (SUMO2), transcript variant 2, mRNA.
15-Sep	0.403	-2.484	15 kDa selenoprotein (SEP15), transcript variant 1, mRNA.

za20d2	0.402	-2.485	zinc finger, AN1-type domain 5 (ZFAND5), mRNA.
pigk	0.402	-2.485	phosphatidylinositol glycan, class K (PIGK), mRNA.
parp12	0.402	-2.486	poly (ADP-ribose) polymerase family, member 12 (PARP12), mRNA.
loc440731	0.402	-2.486	PREDICTED: hypothetical LOC440731, transcript variant 2 (LOC440731), mRNA.
kiaa0427	0.402	-2.486	KIAA0427 (KIAA0427), mRNA.
drg2	0.402	-2.486	developmentally regulated GTP binding protein 2 (DRG2), mRNA.
cuedc1	0.402	-2.487	CUE domain containing 1 (CUEDC1), mRNA.
ephx2	0.402	-2.488	epoxide hydrolase 2, cytoplasmic (EPHX2), mRNA.
c12orf31	0.402	-2.488	chromosome 12 open reading frame 31 (C12orf31), mRNA.
fbn2	0.402	-2.489	fibrillin 2 (congenital contractural arachnodactyly) (FBN2), mRNA.
fkbp9l	0.402	-2.489	FK506 binding protein 9-like (FKBP9L), mRNA.
ctdsp1	0.402	-2.489	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 1 (CTDSP1), transcript variant 2, mRNA.
ly6e	0.402	-2.489	lymphocyte antigen 6 complex, locus E (LY6E), mRNA.
ndufv2	0.402	-2.49	NADH dehydrogenase (ubiquinone) flavoprotein 2, 24kDa (NDUFV2), mRNA.
hmfn0839	0.402	-2.49	lung cancer metastasis-associated protein (MAG1), mRNA.
cdc26	0.401	-2.491	cell division cycle 26 homolog (S. cerevisiae) (CDC26), mRNA.
tcf3	0.401	-2.493	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47) (TCF3), mRNA.
ublcp1	0.401	-2.493	ubiquitin-like domain containing CTD phosphatase 1 (UBLCP1), mRNA.
aarsl	0.401	-2.493	alanyl-tRNA synthetase like (AARSL), mRNA.
exo1	0.401	-2.493	exonuclease 1 (EXO1), transcript variant 2, mRNA.
ctnnbl1	0.401	-2.493	catenin, beta like 1 (CTNNBL1), mRNA.
spp1	0.401	-2.493	secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1) (SPP1), transcript variant 1, mRNA.
rg9mtd1	0.401	-2.494	PREDICTED: RNA (guanine-9-) methyltransferase domain containing 1 (Rg9mtd1), mRNA.
lhfpl2	0.401	-2.494	lipoma HMGIC fusion partner-like 2 (LHFPL2), mRNA.
man2b1	0.401	-2.496	mannosidase, alpha, class 2B, member 1 (MAN2B1), mRNA.
atad3a	0.401	-2.496	ATPase family, AAA domain containing 3A (ATAD3A), mRNA.
pdha1	0.401	-2.497	pyruvate dehydrogenase (lipoamide) alpha 1 (PDHA1), mRNA.
alkbh6	0.4	-2.497	alkB, alkylation repair homolog 6 (E. coli) (ALKBH6), transcript variant 1, mRNA.
cnnm1	0.4	-2.498	cyclin M1 (CNNM1), mRNA.
nr0b1	0.4	-2.499	nuclear receptor subfamily 0, group B, member 1 (NR0B1), mRNA.
grwd1	0.4	-2.499	glutamate-rich WD repeat containing 1 (GRWD1), mRNA.
mcm3	0.4	-2.499	MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) (MCM3), mRNA.
fez2	0.4	-2.5	fasciculation and elongation protein zeta 2 (zygin II) (FEZ2), mRNA.
grina	0.4	-2.5	glutamate receptor, ionotropic, N-methyl D-asparate-associated protein 1 (glutamate binding) (GRINA), transcript variant 2, mRNA.
slc35c1	0.4	-2.5	solute carrier family 35, member C1 (SLC35C1), mRNA.
prpf19	0.4	-2.5	PRP19/PSO4 pre-mRNA processing factor 19 homolog (S. cerevisiae) (PRPF19), mRNA.
dok4	0.4	-2.501	docking protein 4 (DOK4), mRNA.
pttg2	0.4	-2.502	pituitary tumor-transforming 2 (PTTG2), mRNA.
pcbp1	0.4	-2.502	poly(rC) binding protein 1 (PCBP1), mRNA.

c20orf4	0.4	-2.502	chromosome 20 open reading frame 4 (C20orf4), mRNA.
arpc1a	0.4	-2.503	actin related protein 2/3 complex, subunit 1A, 41kDa (ARPC1A), mRNA.
hnrpu	0.399	-2.503	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) (HNRPU), transcript variant 1, mRNA.
pop5	0.399	-2.503	processing of precursor 5, ribonuclease P/MRP subunit (S. cerevisiae) (POP5), transcript variant 2, mRNA.
kifc2	0.399	-2.504	kinesin family member C2 (KIFC2), mRNA.
pnn	0.399	-2.504	pinin, desmosome associated protein (PNN), mRNA.
flj10154	0.399	-2.504	hypothetical protein FLJ10154 (FLJ10154), mRNA.
rpl7	0.399	-2.504	ribosomal protein L7 (RPL7), mRNA.
loc649682	0.399	-2.505	PREDICTED: similar to ribosomal protein L31 (LOC649682), mRNA.
pgrmc2	0.399	-2.505	progesterone receptor membrane component 2 (PGRMC2), mRNA.
c5orf21	0.399	-2.505	chromosome 5 open reading frame 21 (C5orf21), mRNA.
vasp	0.399	-2.506	vasodilator-stimulated phosphoprotein (VASP), mRNA.
crop	0.399	-2.506	cisplatin resistance-associated overexpressed protein (CROP), transcript variant 2, mRNA.
mettl7b	0.399	-2.506	methyltransferase like 7B (METTL7B), mRNA.
chmp4a	0.399	-2.507	chromatin modifying protein 4A (CHMP4A), mRNA.
loc57149	0.399	-2.508	hypothetical protein A-211C6.1 (LOC57149), mRNA.
selm	0.399	-2.509	selenoprotein M (SELM), mRNA.
znf317	0.399	-2.509	zinc finger protein 317 (ZNF317), mRNA.
hs.363526	0.398	-2.509	cDNA clone IMAGE:4837650
hist1h4h	0.398	-2.51	histone cluster 1, H4h (HIST1H4H), mRNA.
bcl2l2	0.398	-2.511	BCL2-like 2 (BCL2L2), mRNA.
hs.575038	0.398	-2.511	cDNA: FLJ21027 fis, clone CAE07110
ipp	0.398	-2.511	intracisternal A particle-promoted polypeptide (IPP), mRNA.
snf1lk	0.398	-2.511	SNF1-like kinase (SNF1LK), mRNA.
gpc6	0.398	-2.511	glypican 6 (GPC6), mRNA.
plekha1	0.398	-2.512	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 (PLEKHA1), transcript variant 2, mRNA.
c1orf57	0.398	-2.513	chromosome 1 open reading frame 57 (C1orf57), mRNA.
loc644972	0.398	-2.513	PREDICTED: similar to ribosomal protein S3a (LOC644972), mRNA.
rhog	0.398	-2.513	ras homolog gene family, member G (rho G) (RHOG), mRNA.
st7	0.398	-2.513	suppression of tumorigenicity 7 (ST7), transcript variant a, mRNA.
znf281	0.398	-2.513	zinc finger protein 281 (ZNF281), mRNA.
thoc2	0.398	-2.513	THO complex 2 (THOC2), mRNA.
gtf3c1	0.398	-2.513	general transcription factor IIIC, polypeptide 1, alpha 220kDa (GTF3C1), mRNA.
loc134997	0.398	-2.514	peptidylprolyl isomerase A processed pseudogene (LOC134997) on chromosome 6.
bxdc2	0.398	-2.514	brix domain containing 2 (BXDC2), mRNA.
mterf	0.398	-2.515	mitochondrial transcription termination factor (MTERF), nuclear gene encoding mitochondrial protein, mRNA.
gpr137b	0.398	-2.515	G protein-coupled receptor 137B (GPR137B), mRNA.
perp	0.398	-2.515	PERP, TP53 apoptosis effector (PERP), mRNA.
myst3	0.398	-2.515	MYST histone acetyltransferase (monocytic leukemia) 3 (MYST3), mRNA.
nifun	0.398	-2.516	iron-sulfur cluster scaffold homolog (E. coli) (ISCU), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
hs.31532	0.397	-2.516	cDNA FLJ26784 fis, clone PRS04220

bard1	0.397	-2.517	BRCA1 associated RING domain 1 (BARD1), mRNA.
map2k2	0.397	-2.517	mitogen-activated protein kinase kinase 2 (MAP2K2), mRNA.
elf1	0.397	-2.517	E74-like factor 1 (ets domain transcription factor) (ELF1), mRNA.
smad3	0.397	-2.517	SMAD family member 3 (SMAD3), mRNA.
cables1	0.397	-2.517	Cdk5 and Abl enzyme substrate 1 (CABLES1), transcript variant 1, mRNA.
tpd52l2	0.397	-2.517	tumor protein D52-like 2 (TPD52L2), transcript variant 3, mRNA.
mbtps1	0.397	-2.517	membrane-bound transcription factor peptidase, site 1 (MBTPS1), mRNA.
loc285900	0.397	-2.517	PREDICTED: similar to 60S ribosomal protein L6 (TAX-responsive enhancer element binding protein 107) (TAXREB107) (Neoplasm-related protein C140), transcript variant 3 (LOC285900), mRNA.
setd1a	0.397	-2.518	SET domain containing 1A (SETD1A), mRNA.
s100a3	0.397	-2.519	S100 calcium binding protein A3 (S100A3), mRNA.
nol3	0.397	-2.519	nucleolar protein 3 (apoptosis repressor with CARD domain) (NOL3), mRNA.
dxs9879e	0.397	-2.519	DNA segment on chromosome X (unique) 9879 expressed sequence (DXS9879E), mRNA.
myo5a	0.397	-2.519	myosin VA (heavy polypeptide 12, myoxin) (MYO5A), mRNA.
ltb4dh	0.397	-2.52	leukotriene B4 12-hydroxydehydrogenase (LTB4DH), mRNA.
frap1	0.397	-2.52	FK506 binding protein 12-rapamycin associated protein 1 (FRAP1), mRNA.
lmna	0.397	-2.52	lamin A/C (LMNA), transcript variant 2, mRNA.
c20orf94	0.397	-2.52	chromosome 20 open reading frame 94 (C20orf94), mRNA.
gnb2l1	0.397	-2.521	guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1 (GNB2L1), mRNA.
ube2c	0.397	-2.521	ubiquitin-conjugating enzyme E2C (UBE2C), transcript variant 5, mRNA.
pdgfrl	0.397	-2.521	platelet-derived growth factor receptor-like (PDGFRL), mRNA.
ppp1r3c	0.396	-2.523	protein phosphatase 1, regulatory (inhibitor) subunit 3C (PPP1R3C), mRNA.
psmd11	0.396	-2.523	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 (PSMD11), mRNA.
pscd1	0.396	-2.524	pleckstrin homology, Sec7 and coiled-coil domains 1(cytohesin 1) (PSCD1), transcript variant 2, mRNA.
rexo2	0.396	-2.524	REX2, RNA exonuclease 2 homolog (S. cerevisiae) (REXO2), mRNA.
hs.347034	0.396	-2.524	cDNA FLJ27231 fis, clone SYN06240
anln	0.396	-2.524	anillin, actin binding protein (ANLN), mRNA.
cttn	0.396	-2.524	cortactin (CTTN), transcript variant 1, mRNA.
slc25a10	0.396	-2.524	solute carrier family 25 (mitochondrial carrier; dicarboxylate transporter), member 10 (SLC25A10), mRNA.
edd1	0.396	-2.524	E3 ubiquitin protein ligase, HECT domain containing, 1 (EDD1), mRNA.
c10orf58	0.396	-2.525	chromosome 10 open reading frame 58 (C10orf58), mRNA.
phip	0.396	-2.525	pleckstrin homology domain interacting protein (PHIP), mRNA.
pmvk	0.396	-2.525	phosphomevalonate kinase (PMVK), mRNA.
ceecam1	0.396	-2.526	cerebral endothelial cell adhesion molecule 1 (CEECAM1), mRNA.
b4galt4	0.396	-2.526	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4 (B4GALT4), transcript variant 2, mRNA.
xrcc3	0.396	-2.526	X-ray repair complementing defective repair in Chinese hamster cells 3 (XRCC3), mRNA.
cdk2ap2	0.396	-2.527	CDK2-associated protein 2 (CDK2AP2), mRNA.
pdia4	0.396	-2.527	protein disulfide isomerase family A, member 4 (PDIA4), mRNA.
sfrs6	0.396	-2.527	splicing factor, arginine/serine-rich 6 (SFRS6), mRNA.
aga	0.396	-2.527	aspartylglucosaminidase (AGA), mRNA.
bhlhb9	0.396	-2.528	basic helix-loop-helix domain containing, class B, 9 (BHLHB9), mRNA.
hs.538962	0.396	-2.528	mRNA; cDNA DKFZp564C152 (from clone DKFZp564C152)

tiparp	0.396	-2.528	TCDD-inducible poly(ADP-ribose) polymerase (TIPARP), mRNA.
gorasp1	0.395	-2.529	golgi reassembly stacking protein 1, 65kDa (GORASP1), mRNA.
ptpn1	0.395	-2.529	protein tyrosine phosphatase, non-receptor type 1 (PTPN1), mRNA.
c6orf136	0.395	-2.529	chromosome 6 open reading frame 136 (C6orf136), mRNA.
loc348262	0.395	-2.529	hypothetical protein LOC348262 (LOC348262), mRNA.
trpc4ap	0.395	-2.529	transient receptor potential cation channel, subfamily C, member 4 associated protein (TRPC4AP), transcript variant 1, mRNA.
ppia	0.395	-2.529	peptidylprolyl isomerase A (cyclophilin A) (PPIA), transcript variant 2, mRNA.
ap2a2	0.395	-2.529	adaptor-related protein complex 2, alpha 2 subunit (AP2A2), mRNA.
atp6v1c1	0.395	-2.53	ATPase, H+ transporting, lysosomal 42kDa, V1 subunit C1 (ATP6V1C1), transcript variant 1, mRNA.
ctsf	0.395	-2.531	cathepsin F (CTSF), mRNA.
cotl1	0.395	-2.532	coactosin-like 1 (Dictyostelium) (COTL1), mRNA.
flj20489	0.395	-2.533	hypothetical protein FLJ20489 (FLJ20489), mRNA.
topors	0.395	-2.533	topoisomerase I binding, arginine/serine-rich (TOPORS), mRNA.
c14orf100	0.395	-2.534	chromosome 14 open reading frame 100 (C14orf100), mRNA.
kiaa0528	0.395	-2.534	KIAA0528 (KIAA0528), mRNA.
degs1	0.395	-2.534	degenerative spermatocyte homolog 1, lipid desaturase (Drosophila) (DEGS1), transcript variant 1, mRNA.
lsm4	0.395	-2.535	LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae) (LSM4), mRNA.
pitrm1	0.394	-2.536	pitrilysin metallopeptidase 1 (PITRM1), mRNA.
loc644615	0.394	-2.536	PREDICTED: similar to Ras-related protein Rab-13 (LOC644615), mRNA.
yap1	0.394	-2.536	Yes-associated protein 1, 65kDa (YAP1), mRNA.
loc93081	0.394	-2.536	chromosome 13 open reading frame 27 (C13orf27), mRNA.
hs.512651	0.394	-2.537	PREDICTED: hypothetical protein DKFZp761G2113 (DKFZp761G2113), mRNA
herc1	0.394	-2.537	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 (HERC1), mRNA.
c1orf59	0.394	-2.538	chromosome 1 open reading frame 59 (C1orf59), mRNA.
wwc1	0.394	-2.538	WW, C2 and coiled-coil domain containing 1 (WWC1), mRNA.
trip11	0.394	-2.538	thyroid hormone receptor interactor 11 (TRIP11), mRNA.
nfkbia	0.394	-2.539	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha (NFKBIA), mRNA.
mapbpip	0.394	-2.54	mitogen-activated protein-binding protein-interacting protein (MAPBPIP), mRNA.
ube2q2	0.394	-2.54	ubiquitin-conjugating enzyme E2Q (putative) 2 (UBE2Q2), mRNA.
stox2	0.394	-2.54	storkhead box 2 (STOX2), mRNA.
ephx1	0.394	-2.541	epoxide hydrolase 1, microsomal (xenobiotic) (EPHX1), mRNA.
fam107b	0.393	-2.542	family with sequence similarity 107, member B (FAM107B), mRNA.
cald1	0.393	-2.542	caldesmon 1 (CALD1), transcript variant 1, mRNA.
ccdc72	0.393	-2.543	coiled-coil domain containing 72 (CCDC72), mRNA.
ndufa4	0.393	-2.544	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4, 9kDa (NDUFA4), nuclear gene encoding mitochondrial protein, mRNA.
tssc4	0.393	-2.545	tumor suppressing subtransferable candidate 4 (TSSC4), mRNA.
atp1b2	0.393	-2.545	ATPase, Na+/K+ transporting, beta 2 polypeptide (ATP1B2), mRNA.
sh3md2	0.393	-2.545	SH3 multiple domains 2 (SH3MD2), mRNA.
tparl	0.393	-2.545	transmembrane protein 165 (TMEM165), mRNA.
mgc4618	0.393	-2.546	transmembrane protein 175 (TMEM175), mRNA.
elp3	0.393	-2.546	elongation protein 3 homolog (S. cerevisiae) (ELP3), mRNA.
hs.531457	0.393	-2.547	cDNA FLJ37595 fis, clone BRCOC2007864

ppih	0.392	-2.548	peptidylprolyl isomerase H (cyclophilin H) (PPIH), mRNA.
tom1	0.392	-2.549	target of myb1 (chicken) (TOM1), mRNA.
rpl28	0.392	-2.549	ribosomal protein L28 (RPL28), mRNA.
hras	0.392	-2.549	v-Ha-ras Harvey rat sarcoma viral oncogene homolog (HRAS), transcript variant 2, mRNA.
pja2	0.392	-2.55	praja 2, RING-H2 motif containing (PJA2), mRNA.
numa1	0.392	-2.551	nuclear mitotic apparatus protein 1 (NUMA1), mRNA.
commd9	0.392	-2.551	COMM domain containing 9 (COMMD9), mRNA.
cdca4	0.392	-2.551	cell division cycle associated 4 (CDCA4), transcript variant 13, mRNA.
pofut2	0.392	-2.552	protein O-fucosyltransferase 2 (POFUT2), transcript variant 3, mRNA.
arse	0.392	-2.552	arylsulfatase E (chondrodysplasia punctata 1) (ARSE), mRNA.
serp1	0.392	-2.552	stress-associated endoplasmic reticulum protein 1 (SERP1), mRNA.
ixl	0.392	-2.552	mediator complex subunit 29 (MED29), mRNA.
impdh2	0.392	-2.553	IMP (inosine monophosphate) dehydrogenase 2 (IMPDH2), mRNA.
mgc17839	0.392	-2.554	transmembrane protein 136 (TMEM136), mRNA.
c16orf56	0.392	-2.554	chromosome 16 open reading frame 56 (C16orf56), mRNA.
kif21a	0.392	-2.554	kinesin family member 21A (KIF21A), mRNA.
zdhhc11	0.391	-2.555	PREDICTED: zinc finger, DHHC-type containing 11 (ZDHHC11), mRNA.
phca	0.391	-2.555	phytoceramidase, alkaline (PHCA), mRNA.
ndufaf1	0.391	-2.557	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, assembly factor 1 (NDUFAF1), mRNA.
hnrpul1	0.391	-2.557	heterogeneous nuclear ribonucleoprotein U-like 1 (HNRPUL1), transcript variant 4, mRNA.
col7a1	0.391	-2.557	collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive) (COL7A1), mRNA.
pik3r1	0.391	-2.558	phosphoinositide-3-kinase, regulatory subunit 1 (p85 alpha) (PIK3R1), transcript variant 2, mRNA.
mtr	0.391	-2.559	5-methyltetrahydrofolate-homocysteine methyltransferase (MTR), mRNA.
drap1	0.391	-2.56	DR1-associated protein 1 (negative cofactor 2 alpha) (DRAP1), mRNA.
trip6	0.391	-2.56	thyroid hormone receptor interactor 6 (TRIP6), mRNA.
actn4	0.391	-2.56	actinin, alpha 4 (ACTN4), mRNA.
loc643438	0.391	-2.561	PREDICTED: similar to Huntingtin interacting protein K (LOC643438), mRNA.
c17orf32	0.39	-2.561	chromosome 17 open reading frame 32 (C17orf32), mRNA.
c12orf48	0.39	-2.561	chromosome 12 open reading frame 48 (C12orf48), mRNA.
trmt12	0.39	-2.561	tRNA methyltransferase 12 homolog (S. cerevisiae) (TRMT12), mRNA.
rab5c	0.39	-2.561	RAB5C, member RAS oncogene family (RAB5C), transcript variant 2, mRNA.
dpp9	0.39	-2.562	dipeptidyl-peptidase 9 (DPP9), mRNA.
kpna3	0.39	-2.562	karyopherin alpha 3 (importin alpha 4) (KPNA3), mRNA.
c14orf106	0.39	-2.562	chromosome 14 open reading frame 106 (C14orf106), mRNA.
scd	0.39	-2.562	stearoyl-CoA desaturase (delta-9-desaturase) (SCD), mRNA.
mgc16028	0.39	-2.563	chromosome 14 open reading frame 179 (C14orf179), mRNA.
asf1b	0.39	-2.563	ASF1 anti-silencing function 1 homolog B (S. cerevisiae) (ASF1B), mRNA.
rheb	0.39	-2.563	Ras homolog enriched in brain (RHEB), mRNA.
ralbp1	0.39	-2.564	ralA binding protein 1 (RALBP1), mRNA.
loc649821	0.39	-2.564	PREDICTED: similar to 60S ribosomal protein L14 (CAG-ISL 7), transcript variant 1 (LOC649821), mRNA.
c7orf27	0.39	-2.564	chromosome 7 open reading frame 27 (C7orf27), mRNA.
snapc5	0.39	-2.565	small nuclear RNA activating complex, polypeptide 5, 19kDa (SNAPC5), mRNA.

hspc142	0.39	-2.565	chromosome 19 open reading frame 62 (C19orf62), transcript variant 2, mRNA.
hcfc1	0.39	-2.565	host cell factor C1 (VP16-accessory protein) (HCFC1), mRNA.
tceal8	0.39	-2.565	transcription elongation factor A (SII)-like 8 (TCEAL8), transcript variant 1, mRNA.
ndufs3	0.39	-2.566	NADH dehydrogenase (ubiquinone) Fe-S protein 3, 30kDa (NADH-coenzyme Q reductase) (NDUFS3), mRNA.
atp6v1f	0.39	-2.566	ATPase, H+ transporting, lysosomal 14kDa, V1 subunit F (ATP6V1F), mRNA.
stx4a	0.39	-2.566	syntaxin 4 (STX4), mRNA.
psma4	0.389	-2.568	proteasome (prosome, macropain) subunit, alpha type, 4 (PSMA4), mRNA.
loc651423	0.389	-2.568	PREDICTED: similar to mitogen-activated protein kinase kinase 3 isoform A (LOC651423), mRNA.
loc389674	0.389	-2.568	PREDICTED: similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein) (hnRNP core protein A1) (HDP-1) (Topoisomerase-inhibitor suppressed) (LOC389674), mRNA.
ccdc127	0.389	-2.569	coiled-coil domain containing 127 (CCDC127), mRNA.
ctsd	0.389	-2.569	cathepsin D (CTSD), mRNA.
lsp1	0.389	-2.569	lymphocyte-specific protein 1 (LSP1), transcript variant 3, mRNA.
clic3	0.389	-2.569	chloride intracellular channel 3 (CLIC3), mRNA.
ckb	0.389	-2.569	creatine kinase, brain (CKB), mRNA.
eif2b5	0.389	-2.57	eukaryotic translation initiation factor 2B, subunit 5 epsilon, 82kDa (EIF2B5), mRNA.
procr	0.389	-2.57	protein C receptor, endothelial (EPCR) (PROCR), mRNA.
rpp40	0.389	-2.57	ribonuclease P 40kDa subunit (RPP40), mRNA.
acot4	0.389	-2.571	acyl-CoA thioesterase 4 (ACOT4), mRNA.
rbm4b	0.389	-2.571	RNA binding motif protein 4B (RBM4B), mRNA.
loc649518	0.389	-2.572	PREDICTED: similar to 40S ribosomal protein S26 (LOC649518), mRNA.
rab36	0.389	-2.572	RAB36, member RAS oncogene family (RAB36), mRNA.
actr3	0.389	-2.572	ARP3 actin-related protein 3 homolog (yeast) (ACTR3), mRNA.
mgc5242	0.389	-2.573	chromosome 7 open reading frame 49 (C7orf49), mRNA.
tram2	0.389	-2.573	translocation associated membrane protein 2 (TRAM2), mRNA.
foxred2	0.389	-2.573	FAD-dependent oxidoreductase domain containing 2 (FOXRED2), mRNA.
tcea2	0.389	-2.573	transcription elongation factor A (SII), 2 (TCEA2), transcript variant 2, mRNA.
zdhhc7	0.389	-2.573	zinc finger, DHHC-type containing 7 (ZDHHC7), mRNA.
ppa2	0.389	-2.574	inorganic pyrophosphatase 2 (PPA2), transcript variant 1, mRNA.
ddah2	0.388	-2.575	dimethylarginine dimethylaminohydrolase 2 (DDAH2), mRNA.
glce	0.388	-2.575	glucuronic acid epimerase (GLCE), mRNA.
loc88523	0.388	-2.576	CG016 (LOC88523), mRNA.
exosc4	0.388	-2.576	exosome component 4 (EXOSC4), mRNA.
snrpa	0.388	-2.577	small nuclear ribonucleoprotein polypeptide A (SNRPA), mRNA.
rab40b	0.388	-2.577	RAB40B, member RAS oncogene family (RAB40B), mRNA.
ndrg1	0.388	-2.577	N-myc downstream regulated gene 1 (NDRG1), mRNA.
kiaa0830	0.388	-2.577	PREDICTED: KIAA0830 protein, transcript variant 4 (KIAA0830), mRNA.
mybbp1a	0.388	-2.578	MYB binding protein (P160) 1a (MYBBP1A), mRNA.
amy1c	0.388	-2.579	amylase, alpha 1C (salivary) (AMY1C), mRNA.
srp19	0.388	-2.579	signal recognition particle 19kDa (SRP19), mRNA.
hs.482497	0.388	-2.579	cDNA FLJ42778 fis, clone BRAWH3005146
fuca1	0.388	-2.579	fucosidase, alpha-L- 1, tissue (FUCA1), mRNA.

axl	0.388	-2.58	AXL receptor tyrosine kinase (AXL), transcript variant 1, mRNA.
prkab1	0.388	-2.58	protein kinase, AMP-activated, beta 1 non-catalytic subunit (PRKAB1), mRNA.
cripak	0.388	-2.581	cysteine-rich PAK1 inhibitor (CRIPAK), mRNA.
c18orf56	0.388	-2.581	chromosome 18 open reading frame 56 (C18orf56), mRNA.
rbm12b	0.387	-2.581	RNA binding motif protein 12B (RBM12B), mRNA.
nrd1	0.387	-2.581	nardilysin (N-arginine dibasic convertase) (NRD1), mRNA.
plekhg2	0.387	-2.582	pleckstrin homology domain containing, family G (with RhoGef domain) member 2 (PLEKHG2), mRNA.
c14orf112	0.387	-2.582	chromosome 14 open reading frame 112 (C14orf112), mRNA.
tada3l	0.387	-2.582	transcriptional adaptor 3 (NGG1 homolog, yeast)-like (TADA3L), transcript variant 2, mRNA.
cerk	0.387	-2.583	ceramide kinase (CERK), transcript variant 1, mRNA.
tsen34	0.387	-2.583	tRNA splicing endonuclease 34 homolog (S. cerevisiae) (TSEN34), mRNA.
cxorf6	0.387	-2.583	chromosome X open reading frame 6 (CXorf6), mRNA.
mcts1	0.387	-2.584	malignant T cell amplified sequence 1 (MCTS1), mRNA.
kiaa1904	0.387	-2.584	KIAA1904 protein (KIAA1904), mRNA.
ptplb	0.387	-2.584	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b (PTPLB), mRNA.
chmp1b	0.387	-2.585	chromatin modifying protein 1B (CHMP1B), mRNA.
neil2	0.387	-2.586	nei like 2 (E. coli) (NEIL2), mRNA.
pcca	0.387	-2.586	propionyl Coenzyme A carboxylase, alpha polypeptide (PCCA), mRNA.
hscb	0.387	-2.586	HscB iron-sulfur cluster co-chaperone homolog (E. coli) (HSCB), mRNA.
psmb5	0.387	-2.586	proteasome (prosome, macropain) subunit, beta type, 5 (PSMB5), mRNA.
cebpz	0.387	-2.587	CCAAT/enhancer binding protein zeta (CEBPZ), mRNA.
scoc	0.387	-2.587	short coiled-coil protein (SCOC), mRNA.
tpm4	0.386	-2.588	tropomyosin 4 (TPM4), mRNA.
rhoq	0.386	-2.589	ras homolog gene family, member Q (RHOQ), mRNA.
kiaa0319l	0.386	-2.589	KIAA0319-like (KIAA0319L), transcript variant 2, mRNA.
als2cr13	0.386	-2.59	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 13 (ALS2CR13), mRNA.
mgat1	0.386	-2.59	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (MGAT1), mRNA.
ahr	0.386	-2.592	aryl hydrocarbon receptor (AHR), mRNA.
hs.190748	0.386	-2.592	PREDICTED: hypothetical LOC388214 (LOC388214), mRNA
nudt22	0.386	-2.592	nudix (nucleoside diphosphate linked moiety X)-type motif 22 (NUDT22), mRNA.
hax1	0.386	-2.592	HCLS1 associated protein X-1 (HAX1), transcript variant 1, mRNA.
sharpin	0.386	-2.593	SHANK-associated RH domain interactor (SHARPIN), mRNA.
ppcs	0.386	-2.593	phosphopantothenoylcysteine synthetase (PPCS), mRNA.
tmem5	0.386	-2.593	transmembrane protein 5 (TMEM5), mRNA.
ubl3	0.386	-2.594	ubiquitin-like 3 (UBL3), mRNA.
trappc6a	0.385	-2.594	trafficking protein particle complex 6A (TRAPPC6A), mRNA.
cenpj	0.385	-2.594	centromere protein J (CENPJ), mRNA.
farsla	0.385	-2.595	phenylalanine-tRNA synthetase-like, alpha subunit (FARSLA), mRNA.
snip1	0.385	-2.596	Smad nuclear interacting protein 1 (SNIP1), mRNA.
tceb2	0.385	-2.597	transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B) (TCEB2), transcript variant 1, mRNA.
nudt1	0.385	-2.597	nudix (nucleoside diphosphate linked moiety X)-type motif 1 (NUDT1), transcript variant 2A, mRNA.
gca	0.385	-2.597	grancalcin, EF-hand calcium binding protein (GCA), mRNA.

leprel2	0.385	-2.597	leprecan-like 2 (LEPREL2), mRNA.
sepw1	0.385	-2.598	selenoprotein W, 1 (SEPW1), mRNA.
cad	0.385	-2.598	carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase (CAD), mRNA.
top1mt	0.385	-2.598	topoisomerase (DNA) I, mitochondrial (TOP1MT), nuclear gene encoding mitochondrial protein, mRNA.
sdsl	0.385	-2.598	serine dehydratase-like (SDSL), mRNA.
itpa	0.385	-2.6	inosine triphosphatase (nucleoside triphosphate pyrophosphatase) (ITPA), transcript variant 2, mRNA.
tesc	0.385	-2.6	tescalcin (TESC), mRNA.
hs.25892	0.385	-2.6	cDNA FLJ38860 fis, clone MESAN2011977
prpf8	0.385	-2.6	PRP8 pre-mRNA processing factor 8 homolog (yeast) (PRPF8), mRNA.
rer1	0.385	-2.601	RER1 retention in endoplasmic reticulum 1 homolog (S. cerevisiae) (RER1), mRNA.
c14orf169	0.384	-2.601	chromosome 14 open reading frame 169 (C14orf169), mRNA.
impad1	0.384	-2.602	inositol monophosphatase domain containing 1 (IMPAD1), mRNA.
mgc10433	0.384	-2.602	hypothetical protein MGC10433 (MGC10433), mRNA.
sumf2	0.384	-2.603	sulfatase modifying factor 2 (SUMF2), transcript variant 4, mRNA.
aph1b	0.384	-2.604	anterior pharynx defective 1 homolog B (C. elegans) (APH1B), mRNA.
moxd1	0.384	-2.604	monooxygenase, DBH-like 1 (MOXD1), transcript variant 1, mRNA.
hs.444999	0.384	-2.604	BX114974 NCI_CGAP_Kid3 cDNA clone IMAGp998C023886, mRNA sequence
ing2	0.384	-2.605	inhibitor of growth family, member 2 (ING2), mRNA.
ddb2	0.384	-2.605	damage-specific DNA binding protein 2, 48kDa (DDB2), mRNA.
brca1	0.384	-2.605	breast cancer 1, early onset (BRCA1), transcript variant BRCA1b, mRNA.
nsun5	0.384	-2.605	NOL1/NOP2/Sun domain family, member 5 (NSUN5), transcript variant 2, mRNA.
npepl1	0.384	-2.606	aminopeptidase-like 1 (NPEPL1), mRNA.
ubap2l	0.384	-2.606	ubiquitin associated protein 2-like (UBAP2L), mRNA.
st13	0.384	-2.606	suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein) (ST13), mRNA.
wbp1	0.384	-2.606	WW domain binding protein 1 (WBP1), mRNA.
snrpc	0.384	-2.607	small nuclear ribonucleoprotein polypeptide C (SNRPC), mRNA.
hs.444329	0.384	-2.607	mRNA; cDNA DKFZp686H20120 (from clone DKFZp686H20120)
nudt2	0.383	-2.608	nudix (nucleoside diphosphate linked moiety X)-type motif 2 (NUDT2), transcript variant 3, mRNA.
loc387753	0.383	-2.608	PREDICTED: similar to 60S ribosomal protein L21 (LOC387753), mRNA.
hs.193784	0.383	-2.608	mRNA; cDNA DKFZp586K1922 (from clone DKFZp586K1922)
tceal1	0.383	-2.609	transcription elongation factor A (SII)-like 1 (TCEAL1), transcript variant 3, mRNA.
gdi1	0.383	-2.609	GDP dissociation inhibitor 1 (GDI1), mRNA.
tars	0.383	-2.609	threonyl-tRNA synthetase (TARS), mRNA.
trim65	0.383	-2.61	tripartite motif-containing 65 (TRIM65), mRNA.
loc440704	0.383	-2.611	PREDICTED: hypothetical gene supported by BC042042 (LOC440704), mRNA.
loc220433	0.383	-2.611	PREDICTED: similar to 40S ribosomal protein S4, X isoform (LOC220433), mRNA.
anxa8l1	0.383	-2.611	annexin A8 (ANXA8), mRNA. XM_931361 XM_931369 XM_931374 XM_931375 XM_931378 XM_931383 XM_931388 XM_931391 XM_931399 XM_931404 XM_931411
tmem113	0.383	-2.611	transmembrane protein 113 (TMEM113), mRNA.
kiaa0446	0.383	-2.612	solute carrier family 25, member 44 (SLC25A44), mRNA.
blcap	0.383	-2.613	bladder cancer associated protein (BLCAP), mRNA.
hs.201854	0.383	-2.613	Homo sapiens, clone IMAGE:4429392, mRNA, partial cds

midn	0.383	-2.613	midnolin (MIDN), mRNA.
txndc9	0.383	-2.613	thioredoxin domain containing 9 (TXNDC9), mRNA.
jub	0.383	-2.613	jub, ajuba homolog (Xenopus laevis) (JUB), transcript variant 1, mRNA.
plod1	0.383	-2.614	procollagen-lysine 1, 2-oxoglutarate 5-dioxygenase 1 (PLOD1), mRNA.
flj12118	0.382	-2.615	cysteinyl-tRNA synthetase 2, mitochondrial (putative) (CARS2), mRNA.
dkfzp686l181	0.382	-2.615	lin-54 homolog (C. elegans) (LIN54), mRNA.
c2orf30	0.382	-2.617	chromosome 2 open reading frame 30 (C2orf30), mRNA.
lrrc14	0.382	-2.618	leucine rich repeat containing 14 (LRRC14), mRNA.
crkrs	0.382	-2.618	Cdc2-related kinase, arginine/serine-rich (CRKRS), mRNA.
lcor	0.382	-2.618	ligand dependent nuclear receptor corepressor (LCOR), mRNA.
arhgdia	0.382	-2.618	Rho GDP dissociation inhibitor (GDI) alpha (ARHGDIA), mRNA.
slc16a4	0.382	-2.619	solute carrier family 16, member 4 (monocarboxylic acid transporter 5) (SLC16A4), mRNA.
gtpbp4	0.382	-2.619	GTP binding protein 4 (GTPBP4), mRNA.
lta4h	0.382	-2.619	leukotriene A4 hydrolase (LTA4H), mRNA.
rbm23	0.382	-2.619	RNA binding motif protein 23 (RBM23), transcript variant 3, mRNA.
pisd	0.382	-2.619	phosphatidylserine decarboxylase (PISD), mRNA.
trim25	0.382	-2.621	tripartite motif-containing 25 (TRIM25), mRNA.
gsto1	0.382	-2.621	glutathione S-transferase omega 1 (GSTO1), mRNA.
pdss2	0.382	-2.621	prenyl (decaprenyl) diphosphate synthase, subunit 2 (PDSS2), mRNA.
loc56902	0.381	-2.621	partner of NOB1 homolog (S. cerevisiae) (PNO1), mRNA.
tcirg1	0.381	-2.623	T-cell, immune regulator 1, ATPase, H+ transporting, lysosomal V0 subunit A3 (TCIRG1), transcript variant 1, mRNA.
pdpr	0.381	-2.623	pyruvate dehydrogenase phosphatase regulatory subunit (PDPR), mRNA.
grn	0.381	-2.626	granulin (GRN), transcript variant 2, mRNA.
c8orf38	0.381	-2.627	chromosome 8 open reading frame 38 (C8orf38), mRNA.
rpp25	0.381	-2.627	ribonuclease P 25kDa subunit (RPP25), mRNA.
hoxc8	0.381	-2.627	homeobox C8 (HOXC8), mRNA.
ciapin1	0.381	-2.627	cytokine induced apoptosis inhibitor 1 (CIAPIN1), mRNA.
eif4g3	0.381	-2.627	eukaryotic translation initiation factor 4 gamma, 3 (EIF4G3), mRNA.
sec23b	0.381	-2.627	Sec23 homolog B (S. cerevisiae) (SEC23B), transcript variant 2, mRNA.
por	0.381	-2.627	P450 (cytochrome) oxidoreductase (POR), mRNA.
tspan4	0.381	-2.628	tetraspanin 4 (TSPAN4), transcript variant 2, mRNA.
tm4sf1	0.381	-2.628	transmembrane 4 L six family member 1 (TM4SF1), mRNA.
ddx17	0.381	-2.628	DEAD (Asp-Glu-Ala-Asp) box polypeptide 17 (DDX17), transcript variant 1, mRNA.
btbd1	0.381	-2.628	BTB (POZ) domain containing 1 (BTBD1), transcript variant 1, mRNA.
nudt14	0.381	-2.628	nudix (nucleoside diphosphate linked moiety X)-type motif 14 (NUDT14), mRNA.
tubb6	0.38	-2.628	PREDICTED: tubulin, beta 6 (TUBB6), mRNA.
znf526	0.38	-2.629	zinc finger protein 526 (ZNF526), mRNA.
pfdn6	0.38	-2.63	prefoldin subunit 6 (PFDN6), mRNA.
mrps26	0.38	-2.63	mitochondrial ribosomal protein S26 (MRPS26), nuclear gene encoding mitochondrial protein, mRNA.
nr1h3	0.38	-2.631	nuclear receptor subfamily 1, group H, member 3 (NR1H3), mRNA.
camsap1l1	0.38	-2.631	calmodulin regulated spectrin-associated protein 1-like 1 (CAMSAP1L1), mRNA.
aspm	0.38	-2.631	asp (abnormal spindle)-like, microcephaly associated (Drosophila) (ASPM), mRNA.

ptd015	0.38	-2.632	chromosome 11 open reading frame 67 (C11orf67), mRNA.
mlstd2	0.38	-2.632	male sterility domain containing 2 (MLSTD2), mRNA.
hcfc1r1	0.38	-2.632	host cell factor C1 regulator 1 (XPO1 dependent) (HCFC1R1), transcript variant 2, mRNA.
rnu35b	0.38	-2.632	small nucleolar RNA, C/D box 35B (SNORD35B) on chromosome 19.
hs.551128	0.38	-2.633	MSTP131 (MST131) mRNA, complete cds
lrrc47	0.38	-2.633	leucine rich repeat containing 47 (LRRC47), mRNA.
sfrs10	0.38	-2.633	splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila) (SFRS10), mRNA.
c18orf21	0.38	-2.633	chromosome 18 open reading frame 21 (C18orf21), mRNA.
wdr72	0.38	-2.634	WD repeat domain 72 (WDR72), mRNA.
yy1	0.379	-2.635	YY1 transcription factor (YY1), mRNA.
rab8b	0.379	-2.635	RAB8B, member RAS oncogene family (RAB8B), mRNA.
smarcc1	0.379	-2.635	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1 (SMARCC1), mRNA.
kiaa0999	0.379	-2.636	KIAA0999 protein (KIAA0999), mRNA.
ilvbl	0.379	-2.636	ilvB (bacterial acetolactate synthase)-like (ILVBL), mRNA.
sephs2	0.379	-2.637	selenophosphate synthetase 2 (SEPHS2), mRNA.
kiaa0664	0.379	-2.637	KIAA0664 (KIAA0664), mRNA.
loc402055	0.379	-2.638	similar to SRR1-like protein (LOC402055), mRNA.
samd9	0.379	-2.638	sterile alpha motif domain containing 9 (SAMD9), mRNA.
ddit4l	0.379	-2.64	DNA-damage-inducible transcript 4-like (DDIT4L), mRNA.
pip5k2c	0.379	-2.641	phosphatidylinositol-4-phosphate 5-kinase, type II, gamma (PIP5K2C), mRNA.
hsdl2	0.379	-2.641	hydroxysteroid dehydrogenase like 2 (HSDL2), mRNA.
loc440160	0.379	-2.641	PREDICTED: hypothetical LOC440160 (LOC440160), mRNA.
ldoc1	0.379	-2.641	leucine zipper, down-regulated in cancer 1 (LDOC1), mRNA.
immp2l	0.378	-2.642	IMP2 inner mitochondrial membrane peptidase-like (S. cerevisiae) (IMMP2L), mRNA.
c11orf51	0.378	-2.642	chromosome 11 open reading frame 51 (C11orf51), mRNA.
pole	0.378	-2.643	polymerase (DNA directed), epsilon (POLE), mRNA.
sf3b4	0.378	-2.643	splicing factor 3b, subunit 4, 49kDa (SF3B4), mRNA.
loc391370	0.378	-2.644	PREDICTED: similar to ribosomal protein S12 (LOC391370), mRNA.
flj11259	0.378	-2.645	hypothetical protein FLJ11259 (FLJ11259), mRNA.
drb1	0.378	-2.646	developmentally regulated RNA-binding protein 1 (DRB1), mRNA.
myl5	0.378	-2.646	myosin, light chain 5, regulatory (MYL5), mRNA.
dnase1l1	0.378	-2.646	deoxyribonuclease I-like 1 (DNASE1L1), transcript variant 2, mRNA.
c20orf108	0.378	-2.646	chromosome 20 open reading frame 108 (C20orf108), mRNA.
mgc52000	0.378	-2.647	CXYorf1-related protein (MGC52000), mRNA.
sec8l1	0.378	-2.648	SEC8-like 1 (S. cerevisiae) (SEC8L1), mRNA.
c5orf14	0.378	-2.648	chromosome 5 open reading frame 14 (C5orf14), mRNA.
loc648000	0.378	-2.648	PREDICTED: similar to 60S ribosomal protein L7, transcript variant 1 (LOC648000), mRNA.
bclaf1	0.377	-2.65	BCL2-associated transcription factor 1 (BCLAF1), transcript variant 1, mRNA.
ttc17	0.377	-2.65	tetratricopeptide repeat domain 17 (TTC17), mRNA.
loc653377	0.377	-2.652	PREDICTED: similar to family with sequence similarity 36, member A (LOC653377), mRNA.
emr1	0.377	-2.652	egf-like module containing, mucin-like, hormone receptor-like 1 (EMR1), mRNA.
pigf	0.377	-2.652	phosphatidylinositol glycan, class F (PIGF), transcript variant 2, mRNA.

dscr2	0.377	-2.652	Down syndrome critical region gene 2 (DSCR2), transcript variant 1, mRNA.
ttc14	0.377	-2.653	tetratricopeptide repeat domain 14 (TTC14), mRNA.
dnpep	0.377	-2.653	aspartyl aminopeptidase (DNPEP), mRNA.
hnrpd	0.377	-2.653	heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kDa) (HNRPD), transcript variant 2, mRNA.
aadacl1	0.377	-2.653	arylacetamide deacetylase-like 1 (AADACL1), mRNA.
anxa2p1	0.377	-2.654	annexin A2 pseudogene 1 (ANXA2P1) on chromosome 4.
cmpk	0.377	-2.654	cytidylate kinase (CMPK), mRNA.
npc2	0.377	-2.654	Niemann-Pick disease, type C2 (NPC2), mRNA.
dctd	0.377	-2.655	dCMP deaminase (DCTD), transcript variant 2, mRNA.
c9orf30	0.377	-2.655	chromosome 9 open reading frame 30 (C9orf30), mRNA.
snupn	0.377	-2.655	snurportin 1 (SNUPN), transcript variant 2, mRNA.
nola2	0.377	-2.655	nucleolar protein family A, member 2 (H/ACA small nucleolar RNPs) (NOLA2), transcript variant 2, mRNA.
dgkd	0.377	-2.655	diacylglycerol kinase, delta 130kDa (DGKD), transcript variant 1, mRNA.
ccdc22	0.377	-2.655	coiled-coil domain containing 22 (CCDC22), mRNA.
mrpl19	0.377	-2.656	mitochondrial ribosomal protein L19 (MRPL19), nuclear gene encoding mitochondrial protein, mRNA.
tmub2	0.377	-2.656	transmembrane and ubiquitin-like domain containing 2 (TMUB2), transcript variant 1, mRNA.
cdv3	0.376	-2.656	CDV3 homolog (mouse) (CDV3), mRNA. XM_945284 XM_945286 XM_945287
klf9	0.376	-2.658	Kruppel-like factor 9 (KLF9), mRNA.
pnkd	0.376	-2.658	paroxysmal nonkinesiogenic dyskinesia (PNKD), transcript variant 1, mRNA.
xpo5	0.376	-2.658	exportin 5 (XPO5), mRNA.
akap12	0.376	-2.659	A kinase (PRKA) anchor protein (gravin) 12 (AKAP12), transcript variant 1, mRNA.
rpa2	0.376	-2.659	replication protein A2, 32kDa (RPA2), mRNA.
thrap5	0.376	-2.659	mediator complex subunit 16 (MED16), mRNA.
lman2l	0.376	-2.659	lectin, mannose-binding 2-like (LMAN2L), mRNA.
rnf130	0.376	-2.659	ring finger protein 130 (RNF130), mRNA.
hs.554410	0.376	-2.66	cDNA FLJ43451 fis, clone OCBBF2033413
c5orf13	0.376	-2.66	chromosome 5 open reading frame 13 (C5orf13), mRNA.
ctsz	0.376	-2.66	cathepsin Z (CTSZ), mRNA.
xpo6	0.376	-2.661	exportin 6 (XPO6), mRNA.
loc205251	0.376	-2.662	hypothetical protein LOC205251 (LOC205251), mRNA.
mvp	0.375	-2.663	major vault protein (MVP), transcript variant 2, mRNA.
tp53i3	0.375	-2.664	tumor protein p53 inducible protein 3 (TP53I3), transcript variant 1, mRNA.
sms	0.375	-2.665	spermine synthase (SMS), mRNA.
serpinb7	0.375	-2.665	serpin peptidase inhibitor, clade B (ovalbumin), member 7 (SERPINB7), mRNA.
mgst2	0.375	-2.665	microsomal glutathione S-transferase 2 (MGST2), mRNA.
heg1	0.375	-2.666	PREDICTED: HEG homolog 1 (zebrafish) (HEG1), mRNA.
fam79a	0.375	-2.666	family with sequence similarity 79, member A (FAM79A), mRNA.
trim28	0.375	-2.667	tripartite motif-containing 28 (TRIM28), mRNA.
agtrap	0.375	-2.667	angiotensin II receptor-associated protein (AGTRAP), transcript variant 2, mRNA.
creld2	0.375	-2.668	cysteine-rich with EGF-like domains 2 (CRELD2), mRNA.
arl8b	0.375	-2.668	ADP-ribosylation factor-like 8B (ARL8B), mRNA.
c6orf166	0.375	-2.668	chromosome 6 open reading frame 166 (C6orf166), mRNA.

tp53i13	0.375	-2.668	tumor protein p53 inducible protein 13 (TP53I13), mRNA.
dhrs7b	0.374	-2.671	dehydrogenase/reductase (SDR family) member 7B (DHRS7B), mRNA.
srpx	0.374	-2.671	sushi-repeat-containing protein, X-linked (SRPX), mRNA.
aco2	0.374	-2.671	aconitase 2, mitochondrial (ACO2), nuclear gene encoding mitochondrial protein, mRNA.
pnma1	0.374	-2.671	paraneoplastic antigen MA1 (PNMA1), mRNA.
hmmr	0.374	-2.671	hyaluronan-mediated motility receptor (RHAMM) (HMMR), transcript variant 2, mRNA.
m6prbp1	0.374	-2.671	mannose-6-phosphate receptor binding protein 1 (M6PRBP1), mRNA.
loc401233	0.374	-2.672	similar to HIV TAT specific factor 1; cofactor required for Tat activation of HIV-1 transcription (LOC401233), mRNA.
ppp4r1	0.374	-2.673	protein phosphatase 4, regulatory subunit 1 (PPP4R1), mRNA.
msn	0.374	-2.674	moesin (MSN), mRNA.
ilf3	0.374	-2.674	interleukin enhancer binding factor 3, 90kDa (ILF3), transcript variant 2, mRNA.
kiaa0746	0.374	-2.675	KIAA0746 protein (KIAA0746), mRNA.
utp14a	0.374	-2.676	UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast) (UTP14A), mRNA.
mtx2	0.374	-2.676	metaxin 2 (MTX2), transcript variant 2, mRNA.
dmtf1	0.374	-2.677	cyclin D binding myb-like transcription factor 1 (DMTF1), mRNA.
mad1l1	0.374	-2.677	MAD1 mitotic arrest deficient-like 1 (yeast) (MAD1L1), transcript variant 2, mRNA.
c10orf11	0.374	-2.677	chromosome 10 open reading frame 11 (C10orf11), mRNA.
coch	0.373	-2.678	coagulation factor C homolog, cochlin (Limulus polyphemus) (COCH), mRNA.
slc16a3	0.373	-2.678	solute carrier family 16, member 3 (monocarboxylic acid transporter 4) (SLC16A3), transcript variant 2, mRNA.
kif11	0.373	-2.678	kinesin family member 11 (KIF11), mRNA.
cps1	0.373	-2.679	carbamoyl-phosphate synthetase 1, mitochondrial (CPS1), mRNA.
recql4	0.373	-2.679	RecQ protein-like 4 (RECQL4), mRNA.
gsto2	0.373	-2.68	glutathione S-transferase omega 2 (GSTO2), mRNA.
col12a1	0.373	-2.68	collagen, type XII, alpha 1 (COL12A1), transcript variant long, mRNA.
mcfd2	0.373	-2.681	multiple coagulation factor deficiency 2 (MCFD2), mRNA.
sdhc	0.373	-2.681	succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa (SDHC), nuclear gene encoding mitochondrial protein, mRNA.
flj14154	0.373	-2.682	hypothetical protein FLJ14154 (FLJ14154), mRNA.
stk35	0.373	-2.682	serine/threonine kinase 35 (STK35), mRNA.
rcn1	0.373	-2.682	reticulocalbin 1, EF-hand calcium binding domain (RCN1), mRNA.
znf518	0.373	-2.682	zinc finger protein 518 (ZNF518), mRNA.
sdpr	0.373	-2.682	serum deprivation response (phosphatidylserine binding protein) (SDPR), mRNA.
kbtbd2	0.373	-2.683	kelch repeat and BTB (POZ) domain containing 2 (KBTBD2), mRNA.
itga5	0.373	-2.683	integrin, alpha 5 (fibronectin receptor, alpha polypeptide) (ITGA5), mRNA.
chpt1	0.373	-2.683	choline phosphotransferase 1 (CHPT1), mRNA.
bfar	0.373	-2.684	bifunctional apoptosis regulator (BFAR), mRNA.
loc440093	0.373	-2.684	similar to H3 histone, family 3B (LOC440093), mRNA.
kiaa0310	0.372	-2.685	KIAA0310 (KIAA0310), mRNA.
ormdl1	0.372	-2.685	ORM1-like 1 (S. cerevisiae) (ORMDL1), mRNA.
hs.125395	0.372	-2.685	BX110351 NCI_CGAP_Kid5 cDNA clone IMAGp998C243778, mRNA sequence
tfpt	0.372	-2.686	TCF3 (E2A) fusion partner (in childhood Leukemia) (TFPT), mRNA.
hsp90aa1	0.372	-2.687	heat shock protein 90kDa alpha (cytosolic), class A member 1 (HSP90AA1), transcript variant 2, mRNA.

mrps28	0.372	-2.687	mitochondrial ribosomal protein S28 (MRPS28), nuclear gene encoding mitochondrial protein, mRNA.
ahcyl1	0.372	-2.687	S-adenosylhomocysteine hydrolase-like 1 (AHCYL1), mRNA.
chchd3	0.372	-2.688	coiled-coil-helix-coiled-coil-helix domain containing 3 (CHCHD3), mRNA.
sec31l1	0.372	-2.689	SEC31 homolog A (S. cerevisiae) (SEC31A), transcript variant 1, mRNA.
usp34	0.372	-2.69	ubiquitin specific peptidase 34 (USP34), mRNA.
cdc25a	0.372	-2.69	cell division cycle 25A (CDC25A), transcript variant 1, mRNA.
b3galt6	0.372	-2.691	UDP-Gal:betaGal beta 1,3-galactosyltransferase polypeptide 6 (B3GALT6), mRNA.
fam120a	0.371	-2.692	family with sequence similarity 120A (FAM120A), mRNA.
psmd2	0.371	-2.693	proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 (PSMD2), mRNA.
nlgn2	0.371	-2.693	neuroligin 2 (NLGN2), mRNA.
itgb5	0.371	-2.693	PREDICTED: integrin, beta 5, transcript variant 3 (ITGB5), mRNA.
frag1	0.371	-2.693	FGF receptor activating protein 1 (FRAG1), mRNA.
ugdh	0.371	-2.693	UDP-glucose dehydrogenase (UGDH), mRNA.
nat9	0.371	-2.693	N-acetyltransferase 9 (NAT9), mRNA.
zscan5	0.371	-2.694	zinc finger and SCAN domain containing 5 (ZSCAN5), mRNA.
serinc1	0.371	-2.694	serine incorporator 1 (SERINC1), mRNA.
usp11	0.371	-2.695	ubiquitin specific peptidase 11 (USP11), mRNA.
loc643287	0.371	-2.695	PREDICTED: similar to prothymosin, alpha (gene sequence 28), transcript variant 1 (LOC643287), mRNA.
c19orf2	0.371	-2.695	chromosome 19 open reading frame 2 (C19orf2), transcript variant 2, mRNA.
fam111a	0.371	-2.696	family with sequence similarity 111, member A (FAM111A), transcript variant 1, mRNA.
asb8	0.371	-2.696	ankyrin repeat and SOCS box-containing 8 (ASB8), mRNA.
phb2	0.371	-2.697	prohibitin 2 (PHB2), mRNA.
prmt5	0.371	-2.697	protein arginine methyltransferase 5 (PRMT5), mRNA.
nubpl	0.371	-2.697	nucleotide binding protein-like (NUBPL), mRNA.
denr	0.371	-2.698	density-regulated protein (DENR), mRNA.
palld	0.371	-2.698	palladin, cytoskeletal associated protein (PALLD), mRNA.
trim26	0.37	-2.7	tripartite motif-containing 26 (TRIM26), mRNA.
c19orf10	0.37	-2.7	chromosome 19 open reading frame 10 (C19orf10), mRNA.
nfia	0.37	-2.7	nuclear factor I/A (NFIA), mRNA.
znf219	0.37	-2.701	zinc finger protein 219 (ZNF219), mRNA.
mfsd1	0.37	-2.702	major facilitator superfamily domain containing 1 (MFSD1), mRNA.
lgp1	0.37	-2.702	homolog of mouse LGP1 (LGP1), mRNA.
vamp3	0.37	-2.702	vesicle-associated membrane protein 3 (cellubrevin) (VAMP3), mRNA.
mesp1	0.37	-2.702	mesoderm posterior 1 homolog (mouse) (MESP1), mRNA.
nrcam	0.37	-2.703	neuronal cell adhesion molecule (NRCAM), mRNA.
ltbr	0.37	-2.703	lymphotoxin beta receptor (TNFR superfamily, member 3) (LTBR), mRNA.
psmd14	0.37	-2.703	proteasome (prosome, macropain) 26S subunit, non-ATPase, 14 (PSMD14), mRNA.
fads1	0.37	-2.704	fatty acid desaturase 1 (FADS1), mRNA.
ubqln2	0.37	-2.705	ubiquilin 2 (UBQLN2), mRNA.
kiaa1600	0.37	-2.705	KIAA1600 (KIAA1600), mRNA.
c10orf57	0.37	-2.705	chromosome 10 open reading frame 57 (C10orf57), mRNA.
plekha9	0.37	-2.706	pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 9 (PLEKHA9), mRNA.

paf1	0.37	-2.706	Paf1, RNA polymerase II associated factor, homolog (S. cerevisiae) (PAF1), mRNA.
sh3bp4	0.37	-2.706	SH3-domain binding protein 4 (SH3BP4), mRNA.
polr2a	0.369	-2.707	polymerase (RNA) II (DNA directed) polypeptide A, 220kDa (POLR2A), mRNA.
wipi1	0.369	-2.707	WD repeat domain, phosphoinositide interacting 1 (WIPI1), mRNA.
plek2	0.369	-2.707	pleckstrin 2 (PLEK2), mRNA.
rps6ka5	0.369	-2.707	ribosomal protein S6 kinase, 90kDa, polypeptide 5 (RPS6KA5), transcript variant 2, mRNA.
mtcp1	0.369	-2.707	mature T-cell proliferation 1 (MTCP1), nuclear gene encoding mitochondrial protein, transcript variant B1, mRNA.
calm3	0.369	-2.707	calmodulin 3 (phosphorylase kinase, delta) (CALM3), mRNA.
slc5a6	0.369	-2.709	solute carrier family 5 (sodium-dependent vitamin transporter), member 6 (SLC5A6), mRNA.
rfc4	0.369	-2.71	replication factor C (activator 1) 4, 37kDa (RFC4), transcript variant 1, mRNA.
thoc1	0.369	-2.71	THO complex 1 (THOC1), mRNA.
lrrc41	0.369	-2.71	leucine rich repeat containing 41 (LRRC41), mRNA.
gpkow	0.369	-2.711	G patch domain and KOW motifs (GPKOW), mRNA.
fscn1	0.369	-2.711	fascin homolog 1, actin-bundling protein (Strongylocentrotus purpuratus) (FSCN1), mRNA.
loc650029	0.369	-2.712	PREDICTED: similar to RNA-binding protein 4 (RNA-binding motif protein 4) (Lark homolog) (Hlark) (RNA-binding motif protein 4a), transcript variant 2 (LOC650029), mRNA.
yipf4	0.369	-2.712	Yip1 domain family, member 4 (YIPF4), mRNA.
6-Mar	0.369	-2.713	membrane-associated ring finger (C3HC4) 6 (MARCH6), mRNA.
cyb5r1	0.369	-2.713	cytochrome b5 reductase 1 (CYB5R1), mRNA.
slc7a6	0.368	-2.715	solute carrier family 7 (cationic amino acid transporter, y+ system), member 6 (SLC7A6), transcript variant 2, mRNA.
aldh3a2	0.368	-2.715	aldehyde dehydrogenase 3 family, member A2 (ALDH3A2), transcript variant 1, mRNA.
mt1f	0.368	-2.716	metallothionein 1F (MT1F), mRNA.
nagk	0.368	-2.717	N-acetylglucosamine kinase (NAGK), mRNA.
asb1	0.368	-2.718	ankyrin repeat and SOCS box-containing 1 (ASB1), mRNA.
gm2a	0.368	-2.718	GM2 ganglioside activator (GM2A), mRNA.
slc35c2	0.368	-2.718	solute carrier family 35, member C2 (SLC35C2), transcript variant 1, mRNA.
tmem106c	0.368	-2.719	transmembrane protein 106C (TMEM106C), mRNA.
adnp	0.368	-2.721	activity-dependent neuroprotector homeobox (ADNP), transcript variant 1, mRNA.
mpdz	0.367	-2.721	multiple PDZ domain protein (MPDZ), mRNA.
cnfn	0.367	-2.722	cornifelin (CNFN), mRNA.
gyg2	0.367	-2.722	glycogenin 2 (GYG2), mRNA.
tfb2m	0.367	-2.723	transcription factor B2, mitochondrial (TFB2M), mRNA.
nkx3-1	0.367	-2.723	NK3 homeobox 1 (NKX3-1), mRNA.
yipf1	0.367	-2.724	Yip1 domain family, member 1 (YIPF1), mRNA.
pcgf4	0.367	-2.724	BMI1 polycomb ring finger oncogene (BMI1), mRNA.
dnm1	0.367	-2.725	dynamin 1 (DNM1), transcript variant 1, mRNA.
cog4	0.367	-2.726	component of oligomeric golgi complex 4 (COG4), mRNA.
cbx5	0.367	-2.726	chromobox homolog 5 (HP1 alpha homolog, Drosophila) (CBX5), mRNA.
mrps17	0.367	-2.726	mitochondrial ribosomal protein S17 (MRPS17), nuclear gene encoding mitochondrial protein, mRNA.
sec61b	0.367	-2.727	Sec61 beta subunit (SEC61B), mRNA.
preb	0.367	-2.728	prolactin regulatory element binding (PREB), mRNA.
rps28	0.367	-2.728	ribosomal protein S28 (RPS28), mRNA.

mxd4	0.367	-2.728	MAX dimerization protein 4 (MXD4), mRNA.
kiaa0690	0.367	-2.729	ribosomal RNA processing 12 homolog (S. cerevisiae) (RRP12), mRNA.
eif4ebp1	0.366	-2.729	eukaryotic translation initiation factor 4E binding protein 1 (EIF4EBP1), mRNA.
g1p3	0.366	-2.729	interferon, alpha-inducible protein (clone IFI-6-16) (G1P3), transcript variant 1, mRNA.
baiap2	0.366	-2.732	BAI1-associated protein 2 (BAIAP2), transcript variant 2, mRNA.
adamts9	0.366	-2.732	ADAM metallopeptidase with thrombospondin type 1 motif, 9 (ADAMTS9), mRNA.
loc652226	0.366	-2.733	PREDICTED: hypothetical protein LOC652226 (LOC652226), mRNA.
litaf	0.366	-2.735	lipopolysaccharide-induced TNF factor (LITAF), mRNA.
alg5	0.365	-2.737	asparagine-linked glycosylation 5 homolog (S. cerevisiae, dolichyl-phosphate beta-glucosyltransferase) (ALG5), mRNA.
nts	0.365	-2.738	neurotensin (NTS), mRNA.
ptrh2	0.365	-2.739	peptidyl-tRNA hydrolase 2 (PTRH2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
itpk1	0.365	-2.739	inositol 1,3,4-triphosphate 5/6 kinase (ITPK1), mRNA.
c12orf57	0.365	-2.74	chromosome 12 open reading frame 57 (C12orf57), mRNA.
hscarg	0.365	-2.742	NmrA-like family domain containing 1 (NMRAL1), mRNA.
gddr	0.365	-2.742	gastrokine 2 (GKN2), mRNA.
abce1	0.365	-2.743	ATP-binding cassette, sub-family E (OABP), member 1 (ABCE1), mRNA.
pcf11	0.365	-2.743	PCF11, cleavage and polyadenylation factor subunit, homolog (S. cerevisiae) (PCF11), mRNA.
hs.193557	0.365	-2.743	cDNA FLJ32401 fis, clone SKMUS2000339
rad21	0.364	-2.744	RAD21 homolog (S. pombe) (RAD21), mRNA.
slc30a1	0.364	-2.744	solute carrier family 30 (zinc transporter), member 1 (SLC30A1), mRNA.
elovl1	0.364	-2.744	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1 (ELOVL1), mRNA.
mrpl15	0.364	-2.746	mitochondrial ribosomal protein L15 (MRPL15), nuclear gene encoding mitochondrial protein, mRNA.
eml4	0.364	-2.746	echinoderm microtubule associated protein like 4 (EML4), mRNA.
ehd4	0.364	-2.746	EH-domain containing 4 (EHD4), mRNA.
uqcrc1	0.364	-2.747	ubiquinol-cytochrome c reductase core protein I (UQCRC1), mRNA.
mte	0.364	-2.747	metallothionein E (MTE), mRNA.
psmb1	0.364	-2.747	proteasome (prosome, macropain) subunit, beta type, 1 (PSMB1), mRNA.
frmd6	0.364	-2.747	FERM domain containing 6 (FRMD6), mRNA.
gpr175	0.364	-2.748	G protein-coupled receptor 175 (GPR175), mRNA.
ets2	0.364	-2.749	v-ets erythroblastosis virus E26 oncogene homolog 2 (avian) (ETS2), mRNA.
eif4g1	0.364	-2.749	eukaryotic translation initiation factor 4 gamma, 1 (EIF4G1), transcript variant 5, mRNA.
fam92a1	0.364	-2.75	PREDICTED: family with sequence similarity 92, member A1, transcript variant 3 (FAM92A1), mRNA.
nme4	0.364	-2.75	non-metastatic cells 4, protein expressed in (NME4), mRNA.
ipo13	0.364	-2.751	importin 13 (IPO13), mRNA.
ccng1	0.364	-2.751	cyclin G1 (CCNG1), transcript variant 2, mRNA.
c9orf3	0.363	-2.751	chromosome 9 open reading frame 3 (C9orf3), mRNA.
vti1b	0.363	-2.752	vesicle transport through interaction with t-SNAREs homolog 1B (yeast) (VTI1B), mRNA.
parn	0.363	-2.752	poly(A)-specific ribonuclease (deadenylation nuclease) (PARN), mRNA.
edg3	0.363	-2.752	endothelial differentiation, sphingolipid G-protein-coupled receptor, 3 (EDG3), mRNA.
ccdc55	0.363	-2.753	coiled-coil domain containing 55 (CCDC55), transcript variant 1, mRNA.
hspb1	0.363	-2.753	heat shock 27kDa protein 1 (HSPB1), mRNA.
flj25801	0.363	-2.753	hypothetical protein FLJ25801 (FLJ25801), mRNA.

eif3s12	0.363	-2.754	eukaryotic translation initiation factor 3, subunit 12 (EIF3S12), mRNA.
rtn4	0.363	-2.755	reticulon 4 (RTN4), transcript variant 3, mRNA.
dnttip1	0.363	-2.755	deoxynucleotidyltransferase, terminal, interacting protein 1 (DNTTIP1), mRNA.
dock10	0.363	-2.755	dedicator of cytokinesis 10 (DOCK10), mRNA.
zbtb9	0.363	-2.756	zinc finger and BTB domain containing 9 (ZBTB9), mRNA.
neo1	0.363	-2.757	neogenin homolog 1 (chicken) (NEO1), mRNA.
sil1	0.363	-2.757	SIL1 homolog, endoplasmic reticulum chaperone (S. cerevisiae) (SIL1), mRNA.
loc219854	0.363	-2.757	PREDICTED: hypothetical protein LOC219854, transcript variant 11 (LOC219854), mRNA.
cep27	0.363	-2.758	centrosomal protein 27kDa (CEP27), mRNA.
fastk	0.362	-2.76	Fas-activated serine/threonine kinase (FASTK), transcript variant 1, mRNA.
nucks1	0.362	-2.76	nuclear casein kinase and cyclin-dependent kinase substrate 1 (NUCKS1), mRNA.
acacb	0.362	-2.76	acetyl-Coenzyme A carboxylase beta (ACACB), mRNA.
npas2	0.362	-2.761	neuronal PAS domain protein 2 (NPAS2), mRNA.
eral1	0.362	-2.761	Era G-protein-like 1 (E. coli) (ERAL1), mRNA.
fam101b	0.362	-2.761	family with sequence similarity 101, member B (FAM101B), mRNA.
pop4	0.362	-2.762	processing of precursor 4, ribonuclease P/MRP subunit (S. cerevisiae) (POP4), mRNA.
ugcg	0.362	-2.763	UDP-glucose ceramide glucosyltransferase (UGCG), mRNA.
prpf31	0.362	-2.763	PRP31 pre-mRNA processing factor 31 homolog (S. cerevisiae) (PRPF31), mRNA.
trip12	0.362	-2.763	thyroid hormone receptor interactor 12 (TRIP12), mRNA.
sdhd	0.362	-2.765	succinate dehydrogenase complex, subunit D, integral membrane protein (SDHD), nuclear gene encoding mitochondrial protein, mRNA.
sedlp	0.362	-2.766	spondyloepiphyseal dysplasia, late, pseudogene (SEDLP) on chromosome 19.
secisbp2	0.361	-2.767	SECIS binding protein 2 (SECISBP2), mRNA.
loc642236	0.361	-2.767	PREDICTED: similar to FRG1 protein (FSHD region gene 1 protein), transcript variant 8 (LOC642236), mRNA.
srxn1	0.361	-2.768	sulfiredoxin 1 homolog (S. cerevisiae) (SRXN1), mRNA.
hs.368255	0.361	-2.768	PREDICTED: KIAA0368 (KIAA0368), mRNA
mrpl2	0.361	-2.769	mitochondrial ribosomal protein L2 (MRPL2), nuclear gene encoding mitochondrial protein, mRNA.
mlh1	0.361	-2.77	mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli) (MLH1), mRNA.
khdrbs1	0.361	-2.77	KH domain containing, RNA binding, signal transduction associated 1 (KHDRBS1), mRNA.
mgc23909	0.361	-2.77	hypothetical protein MGC23909 (MGC23909), mRNA.
fgb	0.361	-2.771	fibrinogen beta chain (FGB), mRNA.
hs.555252	0.361	-2.771	DA371742 BRTHA2 cDNA clone BRTHA2001741 5, mRNA sequence
eif3s6ip	0.361	-2.772	eukaryotic translation initiation factor 3, subunit E interacting protein (EIF3EIP), mRNA.
sdf4	0.361	-2.772	stromal cell derived factor 4 (SDF4), mRNA.
c4orf34	0.361	-2.773	chromosome 4 open reading frame 34 (C4orf34), mRNA.
tpm2	0.361	-2.774	tropomyosin 2 (beta) (TPM2), transcript variant 1, mRNA.
srrm2	0.36	-2.775	serine/arginine repetitive matrix 2 (SRRM2), mRNA.
c10orf32	0.36	-2.775	chromosome 10 open reading frame 32 (C10orf32), mRNA.
lsm10	0.36	-2.775	LSM10, U7 small nuclear RNA associated (LSM10), mRNA.
csf2ra	0.36	-2.776	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) (CSF2RA), transcript variant 6, mRNA.
mtmr4	0.36	-2.777	myotubularin related protein 4 (MTMR4), mRNA.
nt5c2	0.36	-2.778	5'-nucleotidase, cytosolic II (NT5C2), mRNA.

loc440927	0.36	-2.778	PREDICTED: similar to 60S acidic ribosomal protein P1, transcript variant 4 (LOC440927), mRNA.
c14orf82	0.36	-2.778	PREDICTED: chromosome 14 open reading frame 82 (C14orf82), mRNA.
flj22662	0.36	-2.778	hypothetical protein FLJ22662 (FLJ22662), mRNA.
aplp1	0.36	-2.778	amyloid beta (A4) precursor-like protein 1 (APLP1), transcript variant 2, mRNA.
mgc24381	0.36	-2.779	chromosome 16 open reading frame 42 (C16orf42), mRNA.
bambi	0.36	-2.781	BMP and activin membrane-bound inhibitor homolog (Xenopus laevis) (BAMBI), mRNA.
gba	0.36	-2.781	glucosidase, beta; acid (includes glucosylceramidase) (GBA), transcript variant 3, mRNA.
smbp	0.36	-2.781	transmembrane 9 superfamily member 3 (TM9SF3), mRNA.
setmar	0.36	-2.781	SET domain and mariner transposase fusion gene (SETMAR), mRNA.
nab1	0.359	-2.782	NGFI-A binding protein 1 (EGR1 binding protein 1) (NAB1), mRNA.
cd68	0.359	-2.783	CD68 antigen (CD68), mRNA.
zzef1	0.359	-2.783	zinc finger, ZZ-type with EF-hand domain 1 (ZZEF1), mRNA.
jarid2	0.359	-2.783	jumonji, AT rich interactive domain 2 (JARID2), mRNA.
tubb2a	0.359	-2.784	tubulin, beta 2A (TUBB2A), mRNA.
igfbp7	0.359	-2.784	insulin-like growth factor binding protein 7 (IGFBP7), mRNA.
cdc34	0.359	-2.785	cell division cycle 34 homolog (S. cerevisiae) (CDC34), mRNA.
cul4b	0.359	-2.785	cullin 4B (CUL4B), mRNA.
slc35a5	0.359	-2.785	solute carrier family 35, member A5 (SLC35A5), mRNA.
magoh	0.359	-2.786	mago-nashi homolog, proliferation-associated (Drosophila) (MAGOH), mRNA.
ctsh	0.359	-2.786	cathepsin H (CTSH), transcript variant 1, mRNA.
loc647954	0.359	-2.788	PREDICTED: similar to Keratin, type II cytoskeletal 8 (Cytokeratin-8) (CK-8) (Keraton-8) (K8) (LOC647954), mRNA.
cdc16	0.359	-2.788	cell division cycle 16 homolog (S. cerevisiae) (CDC16), transcript variant 2, mRNA.
tcp1	0.359	-2.789	t-complex 1 (TCP1), transcript variant 1, mRNA.
obfc2a	0.359	-2.789	oligonucleotide/oligosaccharide-binding fold containing 2A (OBFC2A), transcript variant 2, mRNA.
tra16	0.359	-2.789	TR4 orphan receptor associated protein TRA16 (TRA16), mRNA.
ctsl1	0.358	-2.79	cathepsin L1 (CTSL1), transcript variant 2, mRNA.
flvcr	0.358	-2.79	feline leukemia virus subgroup C cellular receptor 1 (FLVCR1), mRNA.
rnf103	0.358	-2.791	ring finger protein 103 (RNF103), mRNA.
hs.14555	0.358	-2.791	cDNA: FLJ21513 fis, clone COL05778
polr2c	0.358	-2.792	polymerase (RNA) II (DNA directed) polypeptide C, 33kDa (POLR2C), transcript variant gamma, mRNA.
rps6ka4	0.358	-2.792	ribosomal protein S6 kinase, 90kDa, polypeptide 4 (RPS6KA4), transcript variant 1, mRNA.
mrps12	0.358	-2.792	mitochondrial ribosomal protein S12 (MRPS12), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
tnip1	0.358	-2.794	TNFAIP3 interacting protein 1 (TNIP1), mRNA.
taf15	0.358	-2.794	TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 68kDa (TAF15), transcript variant 2, mRNA.
gyg1	0.358	-2.794	glycogenin 1 (GYG1), mRNA.
sumf1	0.358	-2.794	sulfatase modifying factor 1 (SUMF1), mRNA.
rbm12	0.358	-2.795	RNA binding motif protein 12 (RBM12), transcript variant 1, mRNA.
pink1	0.358	-2.796	PTEN induced putative kinase 1 (PINK1), mRNA.
aco1	0.357	-2.797	aconitase 1, soluble (ACO1), mRNA.
fads2	0.357	-2.797	fatty acid desaturase 2 (FADS2), mRNA.
flj11200	0.357	-2.798	chromosome 4 open reading frame 20 (C4orf20), mRNA.
apbb1ip	0.357	-2.798	amyloid beta (A4) precursor protein-binding, family B, member 1 interacting protein (APBB1IP), mRNA.

klf2	0.357	-2.798	Kruppel-like factor 2 (lung) (KLF2), mRNA.
dpp8	0.357	-2.799	dipeptidyl-peptidase 8 (DPP8), transcript variant 3, mRNA.
g10	0.357	-2.799	BUD31 homolog (S. cerevisiae) (BUD31), mRNA.
nt5e	0.357	-2.801	5'-nucleotidase, ecto (CD73) (NT5E), mRNA.
nrip1	0.357	-2.802	nuclear receptor interacting protein 1 (NRIP1), mRNA.
ncbp2	0.357	-2.802	nuclear cap binding protein subunit 2, 20kDa (NCBP2), mRNA.
il32	0.357	-2.802	interleukin 32 (IL32), transcript variant 3, mRNA.
prpf3	0.357	-2.802	PRP3 pre-mRNA processing factor 3 homolog (S. cerevisiae) (PRPF3), mRNA.
entpd4	0.357	-2.802	ectonucleoside triphosphate diphosphohydrolase 4 (ENTPD4), mRNA.
kcnf1	0.357	-2.803	potassium voltage-gated channel, subfamily F, member 1 (KCNF1), mRNA.
ptprf	0.357	-2.804	protein tyrosine phosphatase, receptor type, F (PTPRF), transcript variant 2, mRNA.
lxn	0.357	-2.804	latexin (LXN), mRNA.
usp7	0.356	-2.805	ubiquitin specific peptidase 7 (herpes virus-associated) (USP7), mRNA.
ard1a	0.356	-2.805	ARD1 homolog A, N-acetyltransferase (S. cerevisiae) (ARD1A), mRNA.
loc653780	0.356	-2.807	PREDICTED: similar to ubiquitin-conjugating enzyme E2C isoform 2 (LOC653780), mRNA.
rdh5	0.356	-2.808	retinol dehydrogenase 5 (11-cis/9-cis) (RDH5), mRNA.
mrpl27	0.356	-2.808	mitochondrial ribosomal protein L27 (MRPL27), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
mlc1sa	0.356	-2.809	myosin light chain 1 slow a (MLC1SA), mRNA.
synj2bp	0.356	-2.81	synaptojanin 2 binding protein (SYNJ2BP), mRNA.
hspd1	0.356	-2.81	heat shock 60kDa protein 1 (chaperonin) (HSPD1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
rab34	0.356	-2.812	RAB34, member RAS oncogene family (RAB34), mRNA.
fam3a	0.355	-2.813	family with sequence similarity 3, member A (FAM3A), mRNA.
loc653888	0.355	-2.814	PREDICTED: similar to Actin-related protein 2/3 complex subunit 1B (ARP2/3 complex 41 kDa subunit) (p41-ARC) (LOC653888), mRNA.
xpr1	0.355	-2.814	xenotropic and polytropic retrovirus receptor (XPR1), mRNA.
c9orf119	0.355	-2.815	PREDICTED: chromosome 9 open reading frame 119 (C9orf119), mRNA.
timm23	0.355	-2.815	PREDICTED: translocase of inner mitochondrial membrane 23 homolog (yeast) (TIMM23), mRNA.
b2m	0.355	-2.816	beta-2-microglobulin (B2M), mRNA.
ltb4r	0.355	-2.816	leukotriene B4 receptor (LTB4R), mRNA.
lman2	0.355	-2.816	lectin, mannose-binding 2 (LMAN2), mRNA.
rdh11	0.355	-2.817	retinol dehydrogenase 11 (all-trans/9-cis/11-cis) (RDH11), mRNA.
loc648113	0.355	-2.817	PREDICTED: similar to M-phase phosphoprotein, mpp8, transcript variant 1 (LOC648113), mRNA.
dmwd	0.355	-2.818	dystrophia myotonica-containing WD repeat motif (DMWD), mRNA.
phldb2	0.355	-2.82	pleckstrin homology-like domain, family B, member 2 (PHLDB2), mRNA.
trove2	0.355	-2.821	TROVE domain family, member 2 (TROVE2), transcript variant 1, mRNA.
mgc15416	0.354	-2.822	chromosome 16 open reading frame 14 (C16orf14), mRNA.
timp2	0.354	-2.822	TIMP metallopeptidase inhibitor 2 (TIMP2), mRNA.
mgc4825	0.354	-2.823	apolipoprotein O (APOO), mRNA.
atp5h	0.354	-2.823	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d (ATP5H), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
lsg1	0.354	-2.823	large subunit GTPase 1 homolog (S. cerevisiae) (LSG1), mRNA.
grhpr	0.354	-2.825	glyoxylate reductase/hydroxypyruvate reductase (GRHPR), mRNA.

loc644422	0.354	-2.825	PREDICTED: similar to arginine/serine-rich splicing factor 6, transcript variant 1 (LOC644422), mRNA.
popdc3	0.354	-2.825	popeye domain containing 3 (POPDC3), mRNA.
hs.5724	0.354	-2.827	mRNA; cDNA DKFZp779O0231 (from clone DKFZp779O0231)
armcx6	0.354	-2.827	armadillo repeat containing, X-linked 6 (ARMCX6), transcript variant 1, mRNA.
cflar	0.354	-2.827	CASP8 and FADD-like apoptosis regulator (CFLAR), mRNA.
c6orf49	0.354	-2.827	chromosome 6 open reading frame 49 (C6orf49), mRNA.
tp53bp1	0.354	-2.828	tumor protein p53 binding protein 1 (TP53BP1), mRNA.
trit1	0.353	-2.829	tRNA isopentenyltransferase 1 (TRIT1), mRNA.
mgc15912	0.353	-2.829	hypothetical protein MGC15912 (MGC15912), mRNA.
rbm15	0.353	-2.83	RNA binding motif protein 15 (RBM15), mRNA.
elf3	0.353	-2.833	E74-like factor 3 (ets domain transcription factor, epithelial-specific) (ELF3), mRNA.
trim41	0.353	-2.833	tripartite motif-containing 41 (TRIM41), transcript variant 1, mRNA.
loc648852	0.353	-2.833	PREDICTED: hypothetical protein LOC648852 (LOC648852), mRNA.
fkbp10	0.353	-2.835	FK506 binding protein 10, 65 kDa (FKBP10), mRNA.
ccdc12	0.353	-2.835	coiled-coil domain containing 12 (CCDC12), mRNA.
c9orf19	0.353	-2.836	chromosome 9 open reading frame 19 (C9orf19), mRNA.
kti12	0.353	-2.836	KTI12 homolog, chromatin associated (S. cerevisiae) (KTI12), mRNA.
gnl3l	0.353	-2.837	guanine nucleotide binding protein-like 3 (nucleolar)-like (GNL3L), mRNA.
znf259	0.352	-2.837	zinc finger protein 259 (ZNF259), mRNA.
c1orf107	0.352	-2.838	chromosome 1 open reading frame 107 (C1orf107), mRNA.
nuak1	0.352	-2.838	NUAK family, SNF1-like kinase, 1 (NUAK1), mRNA.
glud1	0.352	-2.838	glutamate dehydrogenase 1 (GLUD1), mRNA.
qscn6	0.352	-2.839	quiescin Q6 sulfhydryl oxidase 1 (QSOX1), transcript variant 1, mRNA.
casc3	0.352	-2.839	cancer susceptibility candidate 3 (CASC3), mRNA.
mrpl46	0.352	-2.839	mitochondrial ribosomal protein L46 (MRPL46), nuclear gene encoding mitochondrial protein, mRNA.
loc152485	0.352	-2.839	hypothetical protein LOC152485 (LOC152485), mRNA.
ptpn11	0.352	-2.839	protein tyrosine phosphatase, non-receptor type 11 (Noonan syndrome 1) (PTPN11), mRNA.
abhd12	0.352	-2.84	abhydrolase domain containing 12 (ABHD12), transcript variant 1, mRNA.
cdh2	0.352	-2.841	cadherin 2, type 1, N-cadherin (neuronal) (CDH2), mRNA.
stim1	0.352	-2.841	stromal interaction molecule 1 (STIM1), mRNA.
raftlin	0.352	-2.841	raftlin, lipid raft linker 1 (RFTN1), mRNA.
fn3krp	0.352	-2.842	fructosamine-3-kinase-related protein (FN3KRP), mRNA.
tnrc5	0.352	-2.842	trinucleotide repeat containing 5 (TNRC5), transcript variant 1, mRNA.
flj20516	0.352	-2.843	TIMELESS interacting protein (TIPIN), mRNA.
skiv2l2	0.352	-2.843	superkiller viralicidic activity 2-like 2 (S. cerevisiae) (SKIV2L2), mRNA.
supt5h	0.352	-2.844	suppressor of Ty 5 homolog (S. cerevisiae) (SUPT5H), mRNA.
aven	0.352	-2.844	apoptosis, caspase activation inhibitor (AVEN), mRNA.
c8orf59	0.352	-2.845	PREDICTED: chromosome 8 open reading frame 59, transcript variant 2 (C8orf59), mRNA.
arf3	0.352	-2.845	ADP-ribosylation factor 3 (ARF3), mRNA.
c20orf22	0.351	-2.845	chromosome 20 open reading frame 22 (C20orf22), mRNA.
sfrs15	0.351	-2.847	splicing factor, arginine/serine-rich 15 (SFRS15), mRNA.
mbd4	0.351	-2.848	methyl-CpG binding domain protein 4 (MBD4), mRNA.

ichthyin	0.351	-2.849	PREDICTED: ichthyin protein (ICHTHYIN), mRNA.
isg20l1	0.351	-2.849	interferon stimulated exonuclease gene 20kDa-like 1 (ISG20L1), mRNA.
orc6l	0.351	-2.849	origin recognition complex, subunit 6 like (yeast) (ORC6L), mRNA.
trmt5	0.351	-2.849	TRM5 tRNA methyltransferase 5 homolog (S. cerevisiae) (TRMT5), mRNA.
fosb	0.351	-2.852	FBJ murine osteosarcoma viral oncogene homolog B (FOSB), mRNA.
sarm1	0.351	-2.853	sterile alpha and TIR motif containing 1 (SARM1), mRNA.
c9orf5	0.35	-2.854	chromosome 9 open reading frame 5 (C9orf5), mRNA.
tmem66	0.35	-2.856	transmembrane protein 66 (TMEM66), mRNA.
tceb1	0.35	-2.857	transcription elongation factor B (SIII), polypeptide 1 (15kDa, elongin C) (TCEB1), mRNA.
kcnj16	0.35	-2.857	potassium inwardly-rectifying channel, subfamily J, member 16 (KCNJ16), transcript variant 1, mRNA.
gbf1	0.35	-2.857	golgi-specific brefeldin A resistance factor 1 (GBF1), mRNA.
mrpl11	0.35	-2.858	mitochondrial ribosomal protein L11 (MRPL11), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
dkfzp686a01	0.35	-2.858	LIM and calponin homology domains 1 (LIMCH1), mRNA.
cdc2l2	0.35	-2.861	cell division cycle 2-like 2 (PITSLRE proteins) (CDC2L2), transcript variant 5, mRNA.
itgae	0.349	-2.861	integrin, alpha E (antigen CD103, human mucosal lymphocyte antigen 1; alpha polypeptide) (ITGAE), mRNA.
rbm9	0.349	-2.863	RNA binding motif protein 9 (RBM9), transcript variant 2, mRNA.
sfrp1	0.349	-2.863	secreted frizzled-related protein 1 (SFRP1), mRNA.
snrp70	0.349	-2.863	small nuclear ribonucleoprotein 70kDa polypeptide (RNP antigen) (SNRP70), transcript variant 2, mRNA.
hs.27048	0.349	-2.864	cDNA FLJ31750 fis, clone NT2RI2007406
loc644914	0.349	-2.864	PREDICTED: similar to H3 histone, family 3B (LOC644914), mRNA.
s100p	0.349	-2.865	S100 calcium binding protein P (S100P), mRNA.
gtf2h5	0.349	-2.866	general transcription factor IIH, polypeptide 5 (GTF2H5), mRNA.
tnfaip1	0.349	-2.866	tumor necrosis factor, alpha-induced protein 1 (endothelial) (TNFAIP1), mRNA.
pfn1	0.349	-2.867	profilin 1 (PFN1), mRNA.
gnb5	0.349	-2.868	guanine nucleotide binding protein (G protein), beta 5 (GNB5), transcript variant 2, mRNA.
nfic	0.349	-2.868	nuclear factor I/C (CCAAT-binding transcription factor) (NFIC), transcript variant 1, mRNA.
irf1	0.349	-2.868	interferon regulatory factor 1 (IRF1), mRNA.
supt6h	0.348	-2.87	suppressor of Ty 6 homolog (S. cerevisiae) (SUPT6H), mRNA.
ube2z	0.348	-2.872	ubiquitin-conjugating enzyme E2Z (putative) (UBE2Z), mRNA.
tmem44	0.348	-2.873	transmembrane protein 44 (TMEM44), transcript variant 1, mRNA.
aff4	0.348	-2.874	AF4/FMR2 family, member 4 (AFF4), mRNA.
c19orf37	0.348	-2.877	zinc finger protein 428 (ZNF428), mRNA.
blvra	0.347	-2.88	biliverdin reductase A (BLVRA), mRNA.
mfsd5	0.347	-2.88	major facilitator superfamily domain containing 5 (MFSD5), mRNA.
aldh1b1	0.347	-2.88	aldehyde dehydrogenase 1 family, member B1 (ALDH1B1), nuclear gene encoding mitochondrial protein, mRNA.
ube2e1	0.347	-2.881	ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast) (UBE2E1), transcript variant 1, mRNA.
tln2	0.347	-2.881	talin 2 (TLN2), mRNA.
mars	0.347	-2.881	methionyl-tRNA synthetase (MARS), mRNA.
exoc7	0.347	-2.883	exocyst complex component 7 (EXOC7), transcript variant 1, mRNA.
kiaa0323	0.347	-2.883	KIAA0323 (KIAA0323), mRNA.
ecgf1	0.347	-2.883	endothelial cell growth factor 1 (platelet-derived) (ECGF1), mRNA.
trim13	0.347	-2.884	tripartite motif-containing 13 (TRIM13), transcript variant 4, mRNA.

rab9a	0.347	-2.885	RAB9A, member RAS oncogene family (RAB9A), mRNA.
msl3l1	0.347	-2.885	male-specific lethal 3-like 1 (Drosophila) (MSL3L1), transcript variant 4, mRNA.
cox6a1	0.347	-2.886	cytochrome c oxidase subunit VIa polypeptide 1 (COX6A1), nuclear gene encoding mitochondrial protein, mRNA.
lum	0.347	-2.886	lumican (LUM), mRNA.
actr5	0.346	-2.887	ARP5 actin-related protein 5 homolog (yeast) (ACTR5), mRNA.
rtn4ip1	0.346	-2.887	reticulon 4 interacting protein 1 (RTN4IP1), nuclear gene encoding mitochondrial protein, mRNA.
hs.535028	0.346	-2.89	cDNA: FLJ22720 fis, clone HSI14320
calm2	0.346	-2.89	calmodulin 2 (phosphorylase kinase, delta) (CALM2), mRNA.
btg1	0.346	-2.891	B-cell translocation gene 1, anti-proliferative (BTG1), mRNA.
tbc1d7	0.346	-2.894	TBC1 domain family, member 7 (TBC1D7), mRNA.
cxxc5	0.346	-2.894	CXXC finger 5 (CXXC5), mRNA.
chst3	0.345	-2.895	carbohydrate (chondroitin 6) sulfotransferase 3 (CHST3), mRNA.
uqcrc2	0.345	-2.895	ubiquinol-cytochrome c reductase core protein II (UQCRC2), mRNA.
rpl13a	0.345	-2.896	ribosomal protein L13a (RPL13A), mRNA.
mgc40405	0.345	-2.897	hypothetical protein MGC40405 (MGC40405), mRNA.
pomt2	0.345	-2.898	protein-O-mannosyltransferase 2 (POMT2), mRNA.
hcap-g	0.345	-2.898	non-SMC condensin I complex, subunit G (NCAPG), mRNA.
col6a1	0.345	-2.9	collagen, type VI, alpha 1 (COL6A1), mRNA.
elk1	0.345	-2.9	ELK1, member of ETS oncogene family (ELK1), mRNA.
nav1	0.345	-2.901	neuron navigator 1 (NAV1), mRNA.
hmgb3	0.345	-2.902	high-mobility group box 3 (HMGB3), mRNA.
nans	0.344	-2.903	N-acetylneuraminic acid synthase (sialic acid synthase) (NANS), mRNA.
smarce1	0.344	-2.903	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1 (SMARCE1), mRNA.
nqo2	0.344	-2.906	NAD(P)H dehydrogenase, quinone 2 (NQO2), mRNA.
dpp3	0.344	-2.908	dipeptidyl-peptidase 3 (DPP3), transcript variant 1, mRNA.
hdac6	0.344	-2.908	histone deacetylase 6 (HDAC6), mRNA.
ddost	0.344	-2.909	dolichyl-diphosphooligosaccharide-protein glycosyltransferase (DDOST), mRNA.
c16orf34	0.344	-2.91	hematological and neurological expressed 1-like (HN1L), mRNA.
cyr61	0.344	-2.911	cysteine-rich, angiogenic inducer, 61 (CYR61), mRNA.
thumpd1	0.344	-2.911	THUMP domain containing 1 (THUMPD1), mRNA.
rps15a	0.343	-2.912	ribosomal protein S15a (RPS15A), transcript variant 2, mRNA.
lass2	0.343	-2.914	LAG1 longevity assurance homolog 2 (S. cerevisiae) (LASS2), transcript variant 3, mRNA.
fam46a	0.343	-2.914	family with sequence similarity 46, member A (FAM46A), mRNA.
hdac2	0.343	-2.914	histone deacetylase 2 (HDAC2), mRNA.
hnrpa3	0.343	-2.914	heterogeneous nuclear ribonucleoprotein A3 (HNRPA3), mRNA.
tgfbi	0.343	-2.915	transforming growth factor, beta-induced, 68kDa (TGFBI), mRNA.
flj43339	0.343	-2.916	chromosome 15 open reading frame 52 (C15orf52), mRNA.
gtf2a2	0.343	-2.916	general transcription factor IIA, 2, 12kDa (GTF2A2), mRNA.
c3orf17	0.343	-2.916	chromosome 3 open reading frame 17 (C3orf17), transcript variant 2, mRNA.
fcar	0.343	-2.918	Fc fragment of IgA, receptor for (FCAR), transcript variant 10, mRNA.
pcolce2	0.343	-2.919	procollagen C-endopeptidase enhancer 2 (PCOLCE2), mRNA.
insig1	0.343	-2.919	insulin induced gene 1 (INSIG1), transcript variant 2, mRNA.

crmp1	0.343	-2.919	collapsin response mediator protein 1 (CRMP1), transcript variant 1, mRNA.
timp1	0.343	-2.919	TIMP metallopeptidase inhibitor 1 (TIMP1), mRNA.
kiaa1794	0.343	-2.919	KIAA1794 (KIAA1794), mRNA.
zfand2a	0.343	-2.92	zinc finger, AN1-type domain 2A (ZFAND2A), mRNA.
dtx3	0.342	-2.921	deltex 3 homolog (Drosophila) (DTX3), mRNA.
efnb2	0.342	-2.921	ephrin-B2 (EFNB2), mRNA.
loc654103	0.342	-2.922	PREDICTED: similar to solute carrier family 25, member 37 (LOC654103), mRNA.
pfdn5	0.342	-2.923	prefoldin subunit 5 (PFDN5), transcript variant 3, mRNA.
spfh1	0.342	-2.924	ER lipid raft associated 1 (ERLIN1), mRNA.
pelp1	0.342	-2.924	proline, glutamic acid and leucine rich protein 1 (PELP1), mRNA.
plec1	0.342	-2.924	plectin 1, intermediate filament binding protein 500kDa (PLEC1), transcript variant 1, mRNA.
slc30a9	0.342	-2.925	solute carrier family 30 (zinc transporter), member 9 (SLC30A9), mRNA.
hs.570308	0.342	-2.925	BX099724 Soares_fetal_liver_spleen_1NFLS_S1 cDNA clone IMAGp998F201004, mRNA sequence
loc642489	0.342	-2.926	PREDICTED: similar to FK506-binding protein 1A (LOC642489), mRNA.
btbd6	0.342	-2.926	BTB (POZ) domain containing 6 (BTBD6), mRNA.
sema3c	0.342	-2.926	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C (SEMA3C), mRNA.
pdcl3	0.342	-2.927	phosducin-like 3 (PDCL3), mRNA.
dhrs7	0.342	-2.928	dehydrogenase/reductase (SDR family) member 7 (DHRS7), mRNA.
elof1	0.341	-2.929	elongation factor 1 homolog (ELF1, S. cerevisiae) (ELOF1), mRNA.
cebpa	0.341	-2.929	CCAAT/enhancer binding protein (C/EBP), alpha (CEBPA), mRNA.
ssh1	0.341	-2.93	slingshot homolog 1 (Drosophila) (SSH1), mRNA.
eppb9	0.341	-2.93	B9 protein (EPPB9), mRNA.
loc493856	0.341	-2.931	zinc finger, CDGSH-type domain 2 (ZCD2), mRNA.
coil	0.341	-2.931	coilin (COIL), mRNA.
atp1b1	0.341	-2.932	ATPase, Na+/K+ transporting, beta 1 polypeptide (ATP1B1), transcript variant 1, mRNA.
gps2	0.341	-2.932	G protein pathway suppressor 2 (GPS2), mRNA.
cct6ap1	0.341	-2.934	chaperonin containing TCP1, subunit 6A (zeta 1) pseudogene 1 (CCT6AP1) on chromosome 7.
igfbp6	0.341	-2.934	insulin-like growth factor binding protein 6 (IGFBP6), mRNA.
fah	0.341	-2.935	fumarylacetoacetate hydrolase (fumarylacetoacetase) (FAH), mRNA.
lrrc54	0.341	-2.936	tsukushin (TSKU), mRNA.
rhbdd3	0.34	-2.937	rhomboid domain containing 3 (RHBDD3), mRNA.
pdgfc	0.34	-2.939	platelet derived growth factor C (PDGFC), mRNA.
hddc2	0.34	-2.94	HD domain containing 2 (HDDC2), mRNA.
alox5ap	0.34	-2.941	arachidonate 5-lipoxygenase-activating protein (ALOX5AP), mRNA.
hipk2	0.34	-2.942	homeodomain interacting protein kinase 2 (HIPK2), mRNA.
smek2	0.34	-2.944	SMEK homolog 2, suppressor of mek1 (Dictyostelium) (SMEK2), mRNA.
thumpd2	0.34	-2.945	THUMP domain containing 2 (THUMPD2), mRNA.
atbf1	0.339	-2.947	AT-binding transcription factor 1 (ATBF1), mRNA.
inpp5a	0.339	-2.948	PREDICTED: inositol polyphosphate-5-phosphatase, 40kDa (INPP5A), mRNA.
dpm1	0.339	-2.948	dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic subunit (DPM1), mRNA.
dus1l	0.339	-2.948	dihydrouridine synthase 1-like (S. cerevisiae) (DUS1L), mRNA.
loc151579	0.339	-2.949	PREDICTED: similar to basic leucine zipper and W2 domains 1 (LOC151579), mRNA.

loc650428	0.339	-2.949	PREDICTED: similar to keratin, hair, basic, 6 (LOC650428), mRNA.
ckap4	0.339	-2.95	cytoskeleton-associated protein 4 (CKAP4), mRNA.
loc728643	0.339	-2.951	heterogeneous nuclear ribonucleoprotein A1 pseudogene (LOC728643) on chromosome 10.
fut8	0.339	-2.952	fucosyltransferase 8 (alpha (1,6) fucosyltransferase) (FUT8), transcript variant 4, mRNA.
nup210	0.339	-2.953	nucleoporin 210kDa (NUP210), mRNA.
mettl1	0.339	-2.954	methyltransferase like 1 (METTL1), transcript variant 2, mRNA.
loc644931	0.338	-2.955	PREDICTED: hypothetical LOC644931 (LOC644931), mRNA.
ep400	0.338	-2.955	E1A binding protein p400 (EP400), mRNA.
c17orf40	0.338	-2.955	UTP6, small subunit (SSU) processome component, homolog (yeast) (UTP6), mRNA.
srp14p1	0.338	-2.959	signal recognition particle 14kDa (homologous Alu RNA binding protein) pseudogene 1 (SRP14P1) on chromosome 12.
rin2	0.338	-2.961	Ras and Rab interactor 2 (RIN2), mRNA.
loc493869	0.338	-2.961	similar to RIKEN cDNA 2310016C16 (LOC493869), mRNA.
c19orf48	0.338	-2.962	chromosome 19 open reading frame 48 (C19orf48), mRNA.
cd59	0.338	-2.963	CD59 molecule, complement regulatory protein (CD59), transcript variant 4, mRNA.
c6orf153	0.338	-2.963	chromosome 6 open reading frame 153 (C6orf153), mRNA.
fam113a	0.338	-2.963	family with sequence similarity 113, member A (FAM113A), mRNA.
dpm3	0.337	-2.963	dolichyl-phosphate mannosyltransferase polypeptide 3 (DPM3), transcript variant 2, mRNA.
c10orf38	0.337	-2.964	chromosome 10 open reading frame 38 (C10orf38), mRNA.
fam73a	0.337	-2.966	family with sequence similarity 73, member A (FAM73A), mRNA.
huwe1	0.337	-2.968	HECT, UBA and WWE domain containing 1 (HUWE1), mRNA.
mpp1	0.337	-2.968	membrane protein, palmitoylated 1, 55kDa (MPP1), mRNA.
igfbp4	0.337	-2.969	insulin-like growth factor binding protein 4 (IGFBP4), mRNA.
pts	0.337	-2.969	6-pyruvoyltetrahydropterin synthase (PTS), mRNA.
loc644250	0.337	-2.969	PREDICTED: hypothetical protein LOC644250 (LOC644250), mRNA.
ctgf	0.337	-2.969	connective tissue growth factor (CTGF), mRNA.
klhl9	0.337	-2.969	kelch-like 9 (Drosophila) (KLHL9), mRNA.
ndufb3	0.337	-2.97	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12kDa (NDUFB3), mRNA.
smarcad1	0.337	-2.97	SWI/SNF-related, matrix-associated actin-dependent regulator of chromatin , subfamily a, containing DEAD/H box 1 (SMARCAD1), mRNA.
wdr21a	0.336	-2.972	WD repeat domain 21A (WDR21A), transcript variant 1, mRNA.
ypel5	0.336	-2.973	yippee-like 5 (Drosophila) (YPEL5), mRNA.
aytl2	0.336	-2.973	acyltransferase like 2 (AYTL2), mRNA.
ahsa2	0.336	-2.974	AHA1, activator of heat shock 90kDa protein ATPase homolog 2 (yeast) (AHSA2), mRNA.
ddef2	0.336	-2.974	development and differentiation enhancing factor 2 (DDEF2), mRNA.
mdp-1	0.336	-2.975	magnesium-dependent phosphatase 1 (MDP-1), mRNA.
c10orf114	0.336	-2.975	chromosome 10 open reading frame 114 (C10orf114), mRNA.
adar	0.336	-2.979	adenosine deaminase, RNA-specific (ADAR), transcript variant 2, mRNA.
hs.542993	0.336	-2.98	xj89b12.x1 Soares_NFL_T_GBC_S1 cDNA clone IMAGE:2664383 3, mRNA sequence
gabarapl1	0.336	-2.981	GABA(A) receptor-associated protein like 1 (GABARAPL1), mRNA.
zc3h15	0.335	-2.985	zinc finger CCCH-type containing 15 (ZC3H15), mRNA.
rbck1	0.335	-2.985	RanBP-type and C3HC4-type zinc finger containing 1 (RBCK1), transcript variant 1, mRNA.
creg1	0.335	-2.985	cellular repressor of E1A-stimulated genes 1 (CREG1), mRNA.

dynlrb1	0.335	-2.986	dynein, light chain, roadblock-type 1 (DYNLRB1), mRNA.
wasf2	0.335	-2.986	WAS protein family, member 2 (WASF2), mRNA.
ahsa1	0.335	-2.986	AHA1, activator of heat shock 90kDa protein ATPase homolog 1 (yeast) (AHSA1), mRNA.
c20orf172	0.335	-2.987	DSN1, MIND kinetochore complex component, homolog (S. cerevisiae) (DSN1), mRNA.
znf593	0.335	-2.987	zinc finger protein 593 (ZNF593), mRNA.
ftsj1	0.335	-2.988	FtsJ homolog 1 (E. coli) (FTSJ1), transcript variant 1, mRNA.
gpr126	0.335	-2.989	G protein-coupled receptor 126 (GPR126), transcript variant a2, mRNA.
rp1-112k5.2	0.334	-2.99	TSR2, 20S rRNA accumulation, homolog (S. cerevisiae) (TSR2), mRNA.
brd8	0.334	-2.992	bromodomain containing 8 (BRD8), transcript variant 3, mRNA.
dffa	0.334	-2.993	DNA fragmentation factor, 45kDa, alpha polypeptide (DFFA), transcript variant 2, mRNA.
slc16a2	0.334	-2.994	solute carrier family 16, member 2 (monocarboxylic acid transporter 8) (SLC16A2), mRNA.
tarbp1	0.334	-2.995	Tar (HIV-1) RNA binding protein 1 (TARBP1), mRNA.
loc645138	0.334	-2.996	PREDICTED: similar to ribosomal protein S11 (LOC645138), mRNA.
rnf135	0.334	-2.996	ring finger protein 135 (RNF135), transcript variant 2, mRNA.
znhit2	0.334	-2.997	zinc finger, HIT type 2 (ZNHIT2), mRNA.
flj20701	0.334	-2.997	phosphotyrosine interaction domain containing 1 (PID1), mRNA.
tuft1	0.334	-2.998	tuftelin 1 (TUFT1), mRNA.
loc346887	0.334	-2.998	PREDICTED: similar to solute carrier family 16 (monocarboxylic acid transporters), member 14 (LOC346887), mRNA.
nudt5	0.334	-2.998	nudix (nucleoside diphosphate linked moiety X)-type motif 5 (NUDT5), mRNA.
nucb2	0.333	-2.999	nucleobindin 2 (NUCB2), mRNA.
hmg1l1	0.333	-2.999	high-mobility group (nonhistone chromosomal) protein 1-like 1 (HMG1L1), mRNA.
taf7	0.333	-2.999	TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 55kDa (TAF7), mRNA.
stub1	0.333	-2.999	STIP1 homology and U-box containing protein 1 (STUB1), mRNA.
fzd2	0.333	-3	frizzled homolog 2 (Drosophila) (FZD2), mRNA.
c17orf71	0.333	-3	chromosome 17 open reading frame 71 (C17orf71), mRNA.
cnot7	0.333	-3.001	CCR4-NOT transcription complex, subunit 7 (CNOT7), transcript variant 2, mRNA.
cnih4	0.333	-3.002	cornichon homolog 4 (Drosophila) (CNIH4), mRNA.
c20orf30	0.333	-3.003	chromosome 20 open reading frame 30 (C20orf30), transcript variant 2, mRNA.
mllt11	0.333	-3.003	myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 11 (MLLT11), mRNA.
rpl37a	0.333	-3.004	ribosomal protein L37a (RPL37A), mRNA.
eif2ak4	0.333	-3.004	eukaryotic translation initiation factor 2 alpha kinase 4 (EIF2AK4), mRNA.
loc653658	0.333	-3.004	PREDICTED: similar to ribosomal protein S23 (LOC653658), mRNA.
ywhag	0.333	-3.004	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma polypeptide (YWHAG), mRNA.
fn5	0.333	-3.005	chromosome 11 open reading frame 75 (C11orf75), mRNA.
bex1	0.332	-3.009	brain expressed, X-linked 1 (BEX1), mRNA.
kiaa0372	0.332	-3.01	KIAA0372 (KIAA0372), mRNA.
sec14l4	0.332	-3.013	SEC14-like 4 (S. cerevisiae) (SEC14L4), mRNA.
tspyl1	0.332	-3.013	TSPY-like 1 (TSPYL1), mRNA.
loc400948	0.332	-3.015	PREDICTED: similar to CG33774-PA (LOC400948), mRNA.
epas1	0.332	-3.015	endothelial PAS domain protein 1 (EPAS1), mRNA.
kiaa0152	0.332	-3.016	KIAA0152 (KIAA0152), mRNA.
chgn	0.331	-3.017	chondroitin beta1,4 N-acetylgalactosaminyltransferase (ChGn), mRNA.

bmpr2	0.331	-3.017	bone morphogenetic protein receptor, type II (serine/threonine kinase) (BMPR2), mRNA.
rabac1	0.331	-3.02	Rab acceptor 1 (prenylated) (RABAC1), mRNA.
mrpl34	0.331	-3.023	mitochondrial ribosomal protein L34 (MRPL34), nuclear gene encoding mitochondrial protein, mRNA.
zfr	0.331	-3.025	zinc finger RNA binding protein (ZFR), mRNA.
dhrs10	0.33	-3.027	hydroxysteroid (17-beta) dehydrogenase 14 (HSD17B14), mRNA.
mgmt	0.33	-3.027	O-6-methylguanine-DNA methyltransferase (MGMT), mRNA.
kiaa0831	0.33	-3.028	KIAA0831 (KIAA0831), mRNA.
mki67ip	0.33	-3.03	MKI67 (FHA domain) interacting nucleolar phosphoprotein (MKI67IP), mRNA.
golga3	0.33	-3.03	golgi autoantigen, golgin subfamily a, 3 (GOLGA3), mRNA.
c1orf176	0.33	-3.03	chromosome 1 open reading frame 176 (C1orf176), mRNA.
klhdc2	0.33	-3.034	kelch domain containing 2 (KLHDC2), mRNA.
ext1	0.33	-3.034	exostoses (multiple) 1 (EXT1), mRNA.
mrpl10	0.33	-3.034	mitochondrial ribosomal protein L10 (MRPL10), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
syt11	0.33	-3.034	synaptotagmin XI (SYT11), mRNA.
cdkn1a	0.33	-3.035	cyclin-dependent kinase inhibitor 1A (p21, Cip1) (CDKN1A), transcript variant 2, mRNA.
peo1	0.329	-3.037	progressive external ophthalmoplegia 1 (PEO1), mRNA.
ddah1	0.329	-3.038	dimethylarginine dimethylaminohydrolase 1 (DDAH1), mRNA.
topbp1	0.329	-3.038	topoisomerase (DNA) II binding protein 1 (TOPBP1), mRNA.
krthb1	0.329	-3.043	keratin 81 (KRT81), mRNA.
rps26l	0.328	-3.047	40S ribosomal protein S26-like (RPS26L) on chromosome 13.
loc440354	0.328	-3.047	PI-3-kinase-related kinase SMG-1 pseudogene (LOC440354) on chromosome 16.
bcdin3	0.328	-3.047	methylphosphate capping enzyme (MEPCE), mRNA.
nol5a	0.328	-3.048	nucleolar protein 5A (56kDa with KKE/D repeat) (NOL5A), mRNA.
axin2	0.328	-3.048	axin 2 (conductin, axil) (AXIN2), mRNA.
snapc1	0.328	-3.049	small nuclear RNA activating complex, polypeptide 1, 43kDa (SNAPC1), mRNA.
pdcd4	0.328	-3.05	programmed cell death 4 (neoplastic transformation inhibitor) (PDCD4), transcript variant 2, mRNA.
ddef1	0.328	-3.05	development and differentiation enhancing factor 1 (DDEF1), mRNA.
rps3a	0.328	-3.051	ribosomal protein S3A (RPS3A), mRNA.
arhgap10	0.328	-3.051	Rho GTPase activating protein 10 (ARHGAP10), mRNA.
vegfc	0.327	-3.054	vascular endothelial growth factor C (VEGFC), mRNA.
ldhb	0.327	-3.057	lactate dehydrogenase B (LDHB), mRNA.
mgc2408	0.327	-3.058	hypothetical protein MGC2408 (MGC2408), mRNA.
loc644934	0.327	-3.058	PREDICTED: similar to 40S ribosomal protein S26, transcript variant 2 (LOC644934), mRNA.
hs.397465	0.327	-3.059	mRNA; cDNA DKFZp686F1546 (from clone DKFZp686F1546)
stx8	0.327	-3.06	syntaxin 8 (STX8), mRNA.
creb3l2	0.327	-3.061	cAMP responsive element binding protein 3-like 2 (CREB3L2), mRNA.
flj13912	0.327	-3.061	GINS complex subunit 3 (Psf3 homolog) (GINS3), mRNA.
capzb	0.327	-3.062	capping protein (actin filament) muscle Z-line, beta (CAPZB), mRNA.
gas6	0.326	-3.066	growth arrest-specific 6 (GAS6), mRNA.
erh	0.326	-3.066	enhancer of rudimentary homolog (Drosophila) (ERH), mRNA.
loc647048	0.326	-3.066	PREDICTED: similar to hypothetical gene LOC283846 (LOC647048), mRNA.
slc4a2	0.326	-3.067	solute carrier family 4, anion exchanger, member 2 (erythrocyte membrane protein band 3-like 1) (SLC4A2), mRNA.

nox4	0.325	-3.073	NADPH oxidase 4 (NOX4), mRNA.
loc643509	0.325	-3.074	PREDICTED: similar to Dihydrofolate reductase, transcript variant 1 (LOC643509), mRNA.
laptm4a	0.325	-3.074	lysosomal-associated protein transmembrane 4 alpha (LAPTM4A), mRNA.
fbxo32	0.325	-3.075	F-box protein 32 (FBXO32), transcript variant 2, mRNA.
rrs1	0.325	-3.076	RRS1 ribosome biogenesis regulator homolog (S. cerevisiae) (RRS1), mRNA.
lrp11	0.325	-3.076	low density lipoprotein receptor-related protein 11 (LRP11), mRNA.
hs.202577	0.325	-3.077	cDNA FLJ34585 fis, clone KIDNE2008758
isoc2	0.325	-3.077	isochorismatase domain containing 2 (ISOC2), mRNA.
slc22a4	0.325	-3.079	solute carrier family 22 (organic cation transporter), member 4 (SLC22A4), mRNA.
mrpl13	0.325	-3.08	mitochondrial ribosomal protein L13 (MRPL13), nuclear gene encoding mitochondrial protein, mRNA.
jmjd1b	0.325	-3.08	jumonji domain containing 1B (JMJD1B), mRNA.
rhoa	0.325	-3.08	ras homolog gene family, member A (RHOA), mRNA.
socs2	0.325	-3.081	suppressor of cytokine signaling 2 (SOCS2), mRNA.
hnrpa0	0.324	-3.082	heterogeneous nuclear ribonucleoprotein A0 (HNRPA0), mRNA.
golph2	0.324	-3.083	golgi phosphoprotein 2 (GOLPH2), transcript variant 2, mRNA.
hist2h2ac	0.324	-3.084	histone cluster 2, H2ac (HIST2H2AC), mRNA.
kiaa0556	0.324	-3.088	KIAA0556 (KIAA0556), mRNA.
c17orf25	0.324	-3.091	glyoxalase domain containing 4 (GLOD4), mRNA.
rars	0.324	-3.091	arginyl-tRNA synthetase (RARS), mRNA.
gripap1	0.324	-3.091	GRIP1 associated protein 1 (GRIPAP1), transcript variant 2, mRNA.
c20orf177	0.323	-3.092	chromosome 20 open reading frame 177 (C20orf177), mRNA.
hdgfrp3	0.323	-3.094	hepatoma-derived growth factor, related protein 3 (HDGFRP3), mRNA.
hist1h2bd	0.323	-3.095	histone cluster 1, H2bd (HIST1H2BD), transcript variant 1, mRNA.
alg8	0.323	-3.095	asparagine-linked glycosylation 8 homolog (yeast, alpha-1,3-glucosyltransferase) (ALG8), transcript variant 2, mRNA.
c1orf43	0.323	-3.095	chromosome 1 open reading frame 43 (C1orf43), transcript variant 1, mRNA.
tspan10	0.323	-3.095	tetraspanin 10 (TSPAN10), mRNA.
hnrph3	0.323	-3.097	heterogeneous nuclear ribonucleoprotein H3 (2H9) (HNRPH3), transcript variant 2H9A, mRNA.
gcnt3	0.323	-3.099	glucosaminyl (N-acetyl) transferase 3, mucin type (GCNT3), mRNA.
gmpr2	0.323	-3.1	guanosine monophosphate reductase 2 (GMPR2), transcript variant 2, mRNA.
cdk5	0.323	-3.1	cyclin-dependent kinase 5 (CDK5), mRNA.
stx3a	0.323	-3.1	syntaxin 3 (STX3), mRNA.
pola1	0.323	-3.1	polymerase (DNA directed), alpha 1 (POLA1), mRNA.
hlx1	0.323	-3.1	H2.0-like homeobox (HLX), mRNA.
wdr73	0.323	-3.101	WD repeat domain 73 (WDR73), mRNA.
nexn	0.322	-3.101	nexilin (F actin binding protein) (NEXN), mRNA.
gaa	0.322	-3.104	glucosidase, alpha; acid (Pompe disease, glycogen storage disease type II) (GAA), mRNA.
leprot	0.322	-3.104	leptin receptor overlapping transcript (LEPROT), mRNA.
usp36	0.322	-3.105	ubiquitin specific peptidase 36 (USP36), mRNA.
rock2	0.322	-3.106	Rho-associated, coiled-coil containing protein kinase 2 (ROCK2), mRNA.
nup62	0.322	-3.107	nucleoporin 62kDa (NUP62), transcript variant 1, mRNA.
nadsyn1	0.322	-3.108	NAD synthetase 1 (NADSYN1), mRNA.
pomgnt1	0.321	-3.115	protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase (POMGNT1), mRNA.

arl6ip5	0.321	-3.115	ADP-ribosylation-like factor 6 interacting protein 5 (ARL6IP5), mRNA.
atf5	0.321	-3.116	activating transcription factor 5 (ATF5), mRNA.
znf263	0.321	-3.116	zinc finger protein 263 (ZNF263), mRNA.
pigy	0.32	-3.12	phosphatidylinositol glycan, class Y (PIGY), mRNA.
ddx28	0.32	-3.122	DEAD (Asp-Glu-Ala-Asp) box polypeptide 28 (DDX28), nuclear gene encoding mitochondrial protein, mRNA.
blmh	0.32	-3.122	bleomycin hydrolase (BLMH), mRNA.
nsun2	0.32	-3.123	NOL1/NOP2/Sun domain family, member 2 (NSUN2), mRNA.
c17orf62	0.32	-3.124	chromosome 17 open reading frame 62 (C17orf62), mRNA.
zic2	0.32	-3.125	Zic family member 2 (odd-paired homolog, Drosophila) (ZIC2), mRNA.
dld	0.32	-3.129	dihydrolipoamide dehydrogenase (E3 component of pyruvate dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid dehydrogenase complex) (DLD), mRNA.
cct5	0.32	-3.13	chaperonin containing TCP1, subunit 5 (epsilon) (CCT5), mRNA.
ddx27	0.319	-3.13	DEAD (Asp-Glu-Ala-Asp) box polypeptide 27 (DDX27), mRNA.
tmem2	0.319	-3.131	transmembrane protein 2 (TMEM2), mRNA.
bax	0.319	-3.132	BCL2-associated X protein (BAX), transcript variant beta, mRNA.
phtf1	0.319	-3.133	putative homeodomain transcription factor 1 (PHTF1), mRNA.
cebpd	0.319	-3.133	CCAAT/enhancer binding protein (C/EBP), delta (CEBPD), mRNA.
loc652356	0.319	-3.133	PREDICTED: similar to voltage-dependent anion channel 2 (LOC652356), mRNA.
chst7	0.319	-3.135	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7 (CHST7), mRNA.
mrpl16	0.319	-3.136	mitochondrial ribosomal protein L16 (MRPL16), nuclear gene encoding mitochondrial protein, mRNA.
bcl9l	0.319	-3.137	B-cell CLL/lymphoma 9-like (BCL9L), mRNA.
ina	0.319	-3.139	internexin neuronal intermediate filament protein, alpha (INA), mRNA.
srgn	0.318	-3.141	serglycin (SRGN), mRNA.
mfge8	0.318	-3.141	milk fat globule-EGF factor 8 protein (MFGE8), mRNA.
mknk2	0.318	-3.142	MAP kinase interacting serine/threonine kinase 2 (MKNK2), transcript variant 2, mRNA.
ndrg3	0.318	-3.142	NDRG family member 3 (NDRG3), transcript variant 2, mRNA.
pa2g4	0.318	-3.143	proliferation-associated 2G4, 38kDa (PA2G4), mRNA.
flj10781	0.318	-3.143	hypothetical protein FLJ10781 (FLJ10781), mRNA.
anxa8	0.318	-3.144	annexin A8-like 2 (ANXA8L2), mRNA.
fam20b	0.318	-3.144	family with sequence similarity 20, member B (FAM20B), mRNA.
mgc24039	0.318	-3.145	hypothetical protein MGC24039 (MGC24039), mRNA.
sac3d1	0.318	-3.147	SAC3 domain containing 1 (SAC3D1), mRNA.
gmnn	0.318	-3.147	geminin, DNA replication inhibitor (GMNN), mRNA.
serpine1	0.318	-3.148	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1 (SERPINE1), mRNA.
spr	0.318	-3.149	sepiapterin reductase (7,8-dihydrobiopterin:NADP+ oxidoreductase) (SPR), mRNA.
sod1	0.318	-3.15	superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1 (adult)) (SOD1), mRNA.
c7orf36	0.317	-3.15	chromosome 7 open reading frame 36 (C7orf36), mRNA.
dync1h1	0.317	-3.151	dynein, cytoplasmic 1, heavy chain 1 (DYNC1H1), mRNA.
tubgcp5	0.317	-3.154	tubulin, gamma complex associated protein 5 (TUBGCP5), mRNA.
sqrdl	0.317	-3.156	sulfide quinone reductase-like (yeast) (SQRDL), mRNA.
ankrd29	0.317	-3.156	ankyrin repeat domain 29 (ANKRD29), mRNA.
cd55	0.317	-3.157	CD55 molecule, decay accelerating factor for complement (Cromer blood group) (CD55), mRNA.

fhod3	0.316	-3.161	formin homology 2 domain containing 3 (FHOD3), mRNA.
mobkl2b	0.316	-3.162	MOB1, Mps One Binder kinase activator-like 2B (yeast) (MOBKL2B), mRNA.
atp5j	0.316	-3.163	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit F6 (ATP5J) , nuclear gene encoding mitochondrial protein, transcript variant 3, mRNA.
papd5	0.316	-3.163	PAP associated domain containing 5 (PAPD5), mRNA.
tpm1	0.316	-3.163	tropomyosin 1 (alpha) (TPM1), transcript variant 3, mRNA.
gapdh	0.316	-3.164	glyceraldehyde-3-phosphate dehydrogenase (GAPDH), mRNA.
loc388789	0.316	-3.167	PREDICTED: hypothetical gene supported by AF147354 (LOC388789), mRNA.
zbtb20	0.316	-3.169	zinc finger and BTB domain containing 20 (ZBTB20), mRNA.
ppp1r10	0.316	-3.169	protein phosphatase 1, regulatory (inhibitor) subunit 10 (PPP1R10), mRNA.
rage	0.315	-3.17	renal tumor antigen (RAGE), mRNA.
flj13149	0.315	-3.17	FAST kinase domains 5 (FASTKD5), mRNA.
c6orf62	0.315	-3.173	chromosome 6 open reading frame 62 (C6orf62), mRNA.
c6orf111	0.315	-3.175	chromosome 6 open reading frame 111 (C6orf111), mRNA.
fam81a	0.315	-3.175	family with sequence similarity 81, member A (FAM81A), mRNA.
ccdc24	0.315	-3.176	coiled-coil domain containing 24 (CCDC24), mRNA.
dkfzp762i137	0.315	-3.176	zinc finger protein 786 (ZNF786), mRNA.
rnuxa	0.315	-3.177	RNA U, small nuclear RNA export adaptor (phosphorylation regulated) (RNUXA), mRNA.
chd8	0.315	-3.177	chromodomain helicase DNA binding protein 8 (CHD8), mRNA.
mif4gd	0.315	-3.179	MIF4G domain containing (MIF4GD), mRNA.
cul1	0.315	-3.179	cullin 1 (CUL1), mRNA.
kiaa0907	0.314	-3.181	KIAA0907 (KIAA0907), mRNA.
lamb2	0.314	-3.182	laminin, beta 2 (laminin S) (LAMB2), mRNA.
kcng1	0.314	-3.183	potassium voltage-gated channel, subfamily G, member 1 (KCNG1), transcript variant 2, mRNA.
loc644560	0.314	-3.183	PREDICTED: region containing F-box protein 10; chromosome 9 open reading frame 105, transcript variant 1 (LOC644560), mRNA
lmod3	0.314	-3.183	leiomodin 3 (fetal) (LMOD3), mRNA.
id2	0.314	-3.184	inhibitor of DNA binding 2, dominant negative helix-loop-helix protein (ID2), mRNA.
c12orf62	0.314	-3.185	chromosome 12 open reading frame 62 (C12orf62), mRNA.
slitl2	0.314	-3.186	slit-like 2 (Drosophila) (SLITL2), mRNA.
ict1	0.314	-3.187	immature colon carcinoma transcript 1 (ICT1), mRNA.
psma3	0.314	-3.189	proteasome (prosome, macropain) subunit, alpha type, 3 (PSMA3), transcript variant 1, mRNA.
rpn2	0.313	-3.192	ribophorin II (RPN2), mRNA.
rbm25	0.313	-3.192	RNA binding motif protein 25 (RBM25), mRNA.
hnrph2	0.313	-3.192	heterogeneous nuclear ribonucleoprotein H2 (H') (HNRPH2), transcript variant 2, mRNA.
ptrf	0.313	-3.193	polymerase I and transcript release factor (PTRF), mRNA.
pparbp	0.313	-3.193	PPAR binding protein (PPARBP), mRNA.
golt1b	0.313	-3.193	golgi transport 1 homolog B (S. cerevisiae) (GOLT1B), mRNA.
uchl1	0.313	-3.194	ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase) (UCHL1), mRNA.
loc644760	0.313	-3.195	PREDICTED: hypothetical protein LOC644760 (LOC644760), mRNA.
irak1	0.313	-3.196	interleukin-1 receptor-associated kinase 1 (IRAK1), transcript variant 3, mRNA.
gpx4	0.313	-3.196	glutathione peroxidase 4 (phospholipid hydroperoxidase) (GPX4), transcript variant 2, mRNA.
loc653147	0.313	-3.198	PREDICTED: similar to 60S ribosomal protein L26-like 1 (LOC653147), mRNA.

dullard	0.313	-3.199	dullard homolog (Xenopus laevis) (DULLARD), mRNA.
pafah1b1	0.312	-3.205	platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit 45kDa (PAFAH1B1), mRNA.
sympk	0.312	-3.205	symplekin (SYMPK), mRNA.
rspo3	0.312	-3.206	R-spondin 3 homolog (Xenopus laevis) (RSPO3), mRNA.
tmem132a	0.312	-3.207	transmembrane protein 132A (TMEM132A), transcript variant 2, mRNA.
tnfrsf12a	0.312	-3.208	tumor necrosis factor receptor superfamily, member 12A (TNFRSF12A), mRNA.
arpc5	0.312	-3.208	actin related protein 2/3 complex, subunit 5, 16kDa (ARPC5), mRNA.
cdc91l1	0.312	-3.209	phosphatidylinositol glycan anchor biosynthesis, class U (PIGU), mRNA.
gpr37	0.311	-3.217	G protein-coupled receptor 37 (endothelin receptor type B-like) (GPR37), mRNA.
fcmd	0.311	-3.221	Fukuyama type congenital muscular dystrophy (fukutin) (FCMD), mRNA.
h2bfs	0.31	-3.222	H2B histone family, member S (H2BFS), mRNA.
cd63	0.31	-3.223	CD63 antigen (melanoma 1 antigen) (CD63), mRNA.
hn1	0.31	-3.224	hematological and neurological expressed 1 (HN1), transcript variant 1, mRNA.
klhdc5	0.31	-3.228	kelch domain containing 5 (KLHDC5), mRNA.
mtdh	0.31	-3.229	metadherin (MTDH), mRNA.
yars2	0.31	-3.23	tyrosyl-tRNA synthetase 2 (mitochondrial) (YARS2), mRNA.
glt8d1	0.31	-3.23	glycosyltransferase 8 domain containing 1 (GLT8D1), transcript variant 3, mRNA.
bcor	0.31	-3.231	BCL6 co-repressor (BCOR), transcript variant 2, mRNA.
hs.159264	0.309	-3.232	Human clone 23948 mRNA sequence
igfbp1	0.309	-3.233	insulin-like growth factor binding protein 1 (IGFBP1), mRNA.
thy1	0.309	-3.24	Thy-1 cell surface antigen (THY1), mRNA.
pigm	0.308	-3.242	phosphatidylinositol glycan, class M (PIGM), mRNA.
eif5a	0.308	-3.242	eukaryotic translation initiation factor 5A (EIF5A), mRNA.
dnajc7	0.308	-3.243	DnaJ (Hsp40) homolog, subfamily C, member 7 (DNAJC7), mRNA.
mgea5	0.308	-3.244	meningioma expressed antigen 5 (hyaluronidase) (MGEA5), mRNA.
irs2	0.308	-3.245	insulin receptor substrate 2 (IRS2), mRNA.
loc646197	0.308	-3.247	PREDICTED: similar to Heat shock protein HSP 90-beta (HSP 84) (Tumor-specific transplantation 84 kDa antigen) (TSTA) (LOC646197), mRNA.
mrpl24	0.308	-3.251	mitochondrial ribosomal protein L24 (MRPL24), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
zyx	0.308	-3.252	zyxin (ZYX), transcript variant 1, mRNA.
loc130074	0.307	-3.254	hypothetical protein LOC130074 (LOC130074), mRNA.
ankrd13	0.307	-3.259	ankyrin repeat domain 13 (ANKRD13), mRNA.
coro1c	0.307	-3.261	coronin, actin binding protein, 1C (CORO1C), mRNA.
bsdc1	0.307	-3.262	BSD domain containing 1 (BSDC1), mRNA.
peci	0.306	-3.264	peroxisomal D3,D2-enoyl-CoA isomerase (PECI), transcript variant 2, mRNA.
atpbd1b	0.306	-3.265	ATP binding domain 1 family, member B (ATPBD1B), mRNA.
ppapdc1b	0.306	-3.266	phosphatidic acid phosphatase type 2 domain containing 1B (PPAPDC1B), mRNA.
atp1b3	0.306	-3.266	ATPase, Na+/K+ transporting, beta 3 polypeptide (ATP1B3), mRNA. XM_945518
stxbp5	0.306	-3.269	syntaxin binding protein 5 (tomosyn) (STXBP5), mRNA.
c14orf142	0.306	-3.272	chromosome 14 open reading frame 142 (C14orf142), mRNA.
mgc20255	0.306	-3.272	coiled-coil domain containing 97 (CCDC97), mRNA.
pvr	0.306	-3.273	poliovirus receptor (PVR), mRNA.

nfkbib	0.305	-3.275	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta (NFKBIB), transcript variant 1, mRNA.
loc653086	0.305	-3.276	PREDICTED: similar to RAN-binding protein 2-like 1 isoform 2, transcript variant 6 (LOC653086), mRNA.
rn7sl1	0.305	-3.281	RNA, 7SL, cytoplasmic 1 (RN7SL1) on chromosome 14.
golgb1	0.305	-3.283	golgi autoantigen, golgin subfamily b, macrogolgin (with transmembrane signal), 1 (GOLGB1), mRNA.
oxtr	0.304	-3.284	oxytocin receptor (OXTR), mRNA.
ttc3	0.304	-3.287	tetratricopeptide repeat domain 3 (TTC3), transcript variant 1, mRNA.
urod	0.304	-3.288	uroporphyrinogen decarboxylase (UROD), mRNA.
loc649841	0.304	-3.289	PREDICTED: similar to protein immuno-reactive with anti-PTH polyclonal antibodies (LOC649841), mRNA.
blvrb	0.304	-3.29	biliverdin reductase B (flavin reductase (NADPH)) (BLVRB), mRNA.
tsta3	0.304	-3.291	tissue specific transplantation antigen P35B (TSTA3), mRNA.
tmed1	0.304	-3.294	transmembrane emp24 protein transport domain containing 1 (TMED1), mRNA.
fam14a	0.303	-3.295	family with sequence similarity 14, member A (FAM14A), mRNA.
mrpl45	0.303	-3.303	mitochondrial ribosomal protein L45 (MRPL45), nuclear gene encoding mitochondrial protein, mRNA.
tubb4q	0.303	-3.304	tubulin, beta polypeptide 4, member Q (TUBB4Q), mRNA.
naglu	0.303	-3.304	N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB) (NAGLU), mRNA.
fam36a	0.303	-3.304	family with sequence similarity 36, member A (FAM36A), mRNA.
hsd17b7	0.302	-3.306	hydroxysteroid (17-beta) dehydrogenase 7 (HSD17B7), mRNA.
nfib	0.302	-3.307	nuclear factor I/B (NFIB), mRNA.
ubap2	0.302	-3.313	ubiquitin associated protein 2 (UBAP2), mRNA.
morf4l1	0.302	-3.313	mortality factor 4 like 1 (MORF4L1), transcript variant 2, mRNA.
fth1	0.302	-3.314	ferritin, heavy polypeptide 1 (FTH1), mRNA.
acaa2	0.302	-3.314	acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) (ACAA2), nuclear gene encoding mitochondrial protein, mRNA.
sertad1	0.302	-3.315	SERTA domain containing 1 (SERTAD1), mRNA.
papss2	0.302	-3.316	3'-phosphoadenosine 5'-phosphosulfate synthase 2 (PAPSS2), transcript variant 1, mRNA.
adprhl2	0.302	-3.317	ADP-ribosylhydrolase like 2 (ADPRHL2), mRNA.
mgc14376	0.301	-3.317	hypothetical protein MGC14376 (MGC14376), transcript variant 2, mRNA.
dyrk2	0.301	-3.323	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 (DYRK2), transcript variant 2, mRNA.
leng4	0.301	-3.324	leukocyte receptor cluster (LRC) member 4 (LENG4), mRNA.
map7	0.301	-3.327	microtubule-associated protein 7 (MAP7), mRNA.
tomm22	0.301	-3.327	translocase of outer mitochondrial membrane 22 homolog (yeast) (TOMM22), nuclear gene encoding mitochondrial protein, mRNA.
npc1	0.3	-3.328	Niemann-Pick disease, type C1 (NPC1), mRNA.
ankfy1	0.3	-3.331	ankyrin repeat and FYVE domain containing 1 (ANKFY1), transcript variant 2, mRNA.
tacc3	0.3	-3.333	transforming, acidic coiled-coil containing protein 3 (TACC3), mRNA.
ptges	0.299	-3.341	prostaglandin E synthase (PTGES), mRNA.
hspe1	0.299	-3.343	heat shock 10kDa protein 1 (chaperonin 10) (HSPE1), mRNA.
apoc1	0.299	-3.345	apolipoprotein C-I (APOC1), mRNA.
nmi	0.299	-3.346	N-myc (and STAT) interactor (NMI), mRNA.
polr2j3	0.299	-3.347	RPB11b2alpha protein (POLR2J3), mRNA.
slc25a14	0.299	-3.347	solute carrier family 25 (mitochondrial carrier, brain), member 14 (SLC25A14), nuclear gene encoding mitochondrial protein, transcript variant short, mRNA.

hoxb5	0.298	-3.352	homeo box B5 (HOXB5), mRNA.
hadhb	0.298	-3.353	hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit (HADHB), nuclear gene encoding mitochondrial protein, mRNA.
tm9sf4	0.298	-3.354	transmembrane 9 superfamily protein member 4 (TM9SF4), mRNA.
higd2a	0.298	-3.355	HIG1 domain family, member 2A (HIGD2A), mRNA.
mgc2803	0.298	-3.356	chromosome 19 open reading frame 43 (C19orf43), mRNA.
syap1	0.297	-3.362	synapse associated protein 1, SAP47 homolog (Drosophila) (SYAP1), mRNA.
dapk3	0.297	-3.362	death-associated protein kinase 3 (DAPK3), mRNA.
hs.544637	0.297	-3.362	xr14b10.x1 NCI_CGAP_Lu28 cDNA clone IMAGE:2760091 3, mRNA sequence
ctsl2	0.297	-3.363	cathepsin L2 (CTSL2), mRNA.
gprc5c	0.297	-3.364	G protein-coupled receptor, family C, group 5, member C (GPRC5C), transcript variant 2, mRNA.
sec11l3	0.297	-3.364	SEC11 homolog C (S. cerevisiae) (SEC11C), mRNA.
flj40504	0.297	-3.366	hypothetical protein FLJ40504 (FLJ40504), mRNA.
nup214	0.297	-3.369	nucleoporin 214kDa (NUP214), mRNA.
prss23	0.297	-3.37	protease, serine, 23 (PRSS23), mRNA.
sc65	0.296	-3.373	synaptonemal complex protein SC65 (SC65), mRNA.
mfhas1	0.296	-3.373	malignant fibrous histiocytoma amplified sequence 1 (MFHAS1), mRNA.
ccdc58	0.296	-3.373	coiled-coil domain containing 58 (CCDC58), mRNA.
slc25a13	0.296	-3.374	solute carrier family 25, member 13 (citrin) (SLC25A13), mRNA.
phlda3	0.296	-3.374	pleckstrin homology-like domain, family A, member 3 (PHLDA3), mRNA.
c20orf31	0.296	-3.374	ER degradation enhancer, mannosidase alpha-like 2 (EDEM2), mRNA.
adck4	0.296	-3.376	aarF domain containing kinase 4 (ADCK4), mRNA.
hspa9b	0.296	-3.376	heat shock 70kDa protein 9B (mortalin-2) (HSPA9B), nuclear gene encoding mitochondrial protein, mRNA.
eno3	0.296	-3.38	enolase 3 (beta, muscle) (ENO3), transcript variant 1, mRNA.
npepps	0.296	-3.381	aminopeptidase puromycin sensitive (NPEPPS), mRNA.
fgf2	0.296	-3.381	fibroblast growth factor 2 (basic) (FGF2), mRNA.
rbm34	0.295	-3.393	RNA binding motif protein 34 (RBM34), mRNA.
tdg	0.294	-3.396	thymine-DNA glycosylase (TDG), mRNA.
prkca	0.294	-3.402	protein kinase C, alpha (PRKCA), mRNA.
h2afy	0.294	-3.402	H2A histone family, member Y (H2AFY), transcript variant 2, mRNA.
fthl3	0.294	-3.403	ferritin, heavy polypeptide-like 3 (FTHL3) on chromosome 2.
hs.553187	0.294	-3.405	cDNA FLJ45619 fis, clone BRTHA3027318
dusp3	0.294	-3.406	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) (DUSP3), mRNA.
pkd2	0.294	-3.407	polycystic kidney disease 2 (autosomal dominant) (PKD2), mRNA.
ythdf2	0.293	-3.41	YTH domain family, member 2 (YTHDF2), mRNA.
loc723972	0.293	-3.41	hepatopoietin PCn127 (LOC723972) on chromosome 15.
hs.137971	0.293	-3.41	AV737943 CB cDNA clone CBDAEG06 5, mRNA sequence
hs.546710	0.293	-3.411	cDNA FLJ26122 fis, clone SYN00634
wdr25	0.293	-3.416	WD repeat domain 25 (WDR25), mRNA.
u1snrnpbp	0.293	-3.417	U11/U12 snRNP 35K (U1SNRNPBP), transcript variant 3, mRNA.
mgc2747	0.293	-3.419	hypothetical protein MGC2747 (MGC2747), mRNA.
endogl1	0.292	-3.421	endonuclease G-like 1 (ENDOGL1), mRNA.

pon3	0.292	-3.421	paraoxonase 3 (PON3), mRNA.
kynu	0.292	-3.422	kynureninase (L-kynurenine hydrolase) (KYNU), transcript variant 1, mRNA.
fhl1	0.292	-3.423	four and a half LIM domains 1 (FHL1), mRNA.
fstl3	0.292	-3.426	follistatin-like 3 (secreted glycoprotein) (FSTL3), mRNA.
fjx1	0.292	-3.427	four jointed box 1 (Drosophila) (FJX1), mRNA.
fthl11	0.292	-3.429	ferritin, heavy polypeptide-like 11 (FTHL11) on chromosome 8.
b4galt1	0.291	-3.431	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1 (B4GALT1), mRNA.
serpine2	0.291	-3.434	serpin peptidase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 2 (SERPINE2), mRNA.
vps18	0.291	-3.435	vacuolar protein sorting 18 homolog (S. cerevisiae) (VPS18), mRNA.
rpl14	0.291	-3.436	ribosomal protein L14 (RPL14), transcript variant 1, mRNA.
hnrpl	0.291	-3.438	heterogeneous nuclear ribonucleoprotein L (HNRPL), transcript variant 2, mRNA.
c16orf57	0.291	-3.438	chromosome 16 open reading frame 57 (C16orf57), mRNA.
ak2	0.291	-3.439	adenylate kinase 2 (AK2), transcript variant AK2A, mRNA.
foxc1	0.291	-3.441	forkhead box C1 (FOXC1), mRNA.
mgc18216	0.29	-3.444	PREDICTED: hypothetical protein MGC18216 (MGC18216), mRNA.
f2rl1	0.29	-3.444	coagulation factor II (thrombin) receptor-like 1 (F2RL1), mRNA.
ckap1	0.29	-3.448	cytoskeleton associated protein 1 (CKAP1), mRNA.
sptlc1	0.29	-3.449	serine palmitoyltransferase, long chain base subunit 1 (SPTLC1), transcript variant 2, mRNA.
sync1	0.29	-3.449	syncoilin, intermediate filament 1 (SYNC1), mRNA.
dio2	0.29	-3.45	deiodinase, iodothyronine, type II (DIO2), transcript variant 1, mRNA.
loc653328	0.289	-3.455	PREDICTED: similar to ribosomal protein S27 (LOC653328), mRNA.
sat2	0.289	-3.457	spermidine/spermine N1-acetyltransferase 2 (SAT2), mRNA.
flj13391	0.289	-3.457	transmembrane protein 166 (TMEM166), mRNA.
fam62a	0.289	-3.462	family with sequence similarity 62 (C2 domain containing), member A (FAM62A), mRNA.
derl2	0.289	-3.462	Der1-like domain family, member 2 (DERL2), mRNA.
tmed2	0.289	-3.462	transmembrane emp24 domain trafficking protein 2 (TMED2), mRNA.
edf1	0.289	-3.465	endothelial differentiation-related factor 1 (EDF1), transcript variant alpha, mRNA.
cd99l2	0.288	-3.466	CD99 molecule-like 2 (CD99L2), transcript variant 3, mRNA.
slc38a6	0.288	-3.467	solute carrier family 38, member 6 (SLC38A6), mRNA.
serpinh1	0.288	-3.467	serpin peptidase inhibitor, clade H (heat shock protein 47), member 1, (collagen binding protein 1) (SERPINH1), mRNA.
loc440732	0.288	-3.471	PREDICTED: similar to 40S ribosomal protein S7 (S8) (LOC440732), mRNA.
ube1	0.287	-3.482	ubiquitin-activating enzyme E1 (UBE1), transcript variant 2, mRNA.
flj20254	0.287	-3.482	hypothetical protein FLJ20254 (FLJ20254), mRNA.
ciao1	0.287	-3.483	cytosolic iron-sulfur protein assembly 1 homolog (S. cerevisiae) (CIAO1), mRNA.
c14orf102	0.287	-3.485	chromosome 14 open reading frame 102 (C14orf102), transcript variant 2, mRNA.
timp4	0.287	-3.486	TIMP metallopeptidase inhibitor 4 (TIMP4), mRNA.
keap1	0.287	-3.487	kelch-like ECH-associated protein 1 (KEAP1), transcript variant 2, mRNA.
cited2	0.287	-3.488	Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2 (CITED2), mRNA.
calu	0.286	-3.491	calumenin (CALU), mRNA.
ier5	0.286	-3.491	immediate early response 5 (IER5), mRNA.
tmpit	0.286	-3.495	transmembrane protein 120A (TMEM120A), mRNA.
hebp1	0.286	-3.496	heme binding protein 1 (HEBP1), mRNA.

rps19	0.286	-3.496	ribosomal protein S19 (RPS19), mRNA.
eif4e2	0.286	-3.496	eukaryotic translation initiation factor 4E family member 2 (EIF4E2), mRNA.
usp49	0.286	-3.5	ubiquitin specific peptidase 49 (USP49), mRNA.
h3f3b	0.286	-3.501	H3 histone, family 3B (H3.3B) (H3F3B), mRNA.
exosc9	0.286	-3.502	exosome component 9 (EXOSC9), mRNA.
bcyrn1	0.285	-3.504	brain cytoplasmic RNA 1, Bc1 analog (mouse) (BCYRN1) on chromosome 2.
ranbp1	0.285	-3.505	RAN binding protein 1 (RANBP1), mRNA.
c10orf9	0.285	-3.507	cyclin Y (CCNY), transcript variant 2, mRNA.
arhgap21	0.285	-3.509	Rho GTPase activating protein 21 (ARHGAP21), mRNA.
rsad1	0.285	-3.511	radical S-adenosyl methionine domain containing 1 (RSAD1), mRNA.
snf8	0.284	-3.516	SNF8, ESCRT-II complex subunit, homolog (S. cerevisiae) (SNF8), mRNA.
slc25a11	0.284	-3.517	solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11 (SLC25A11), mRNA.
ptpns1	0.284	-3.523	protein tyrosine phosphatase, non-receptor type substrate 1 (PTPNS1), mRNA.
slit3	0.283	-3.531	slit homolog 3 (Drosophila) (SLIT3), mRNA.
retsat	0.283	-3.531	retinol saturase (all-trans-retinol 13,14-reductase) (RETSAT), mRNA.
mrpl50	0.283	-3.532	mitochondrial ribosomal protein L50 (MRPL50), nuclear gene encoding mitochondrial protein, mRNA.
cyp24a1	0.283	-3.532	cytochrome P450, family 24, subfamily A, polypeptide 1 (CYP24A1), nuclear gene encoding mitochondrial protein, mRNA.
mtmr15	0.283	-3.534	myotubularin related protein 15 (MTMR15), mRNA.
loc389541	0.283	-3.536	similar to CG14977-PA (LOC389541), mRNA.
sqstm1	0.282	-3.542	sequestosome 1 (SQSTM1), mRNA.
eif1b	0.282	-3.547	eukaryotic translation initiation factor 1B (EIF1B), mRNA.
dstn	0.282	-3.547	destrin (actin depolymerizing factor) (DSTN), transcript variant 1, mRNA.
sdc4	0.282	-3.55	syndecan 4 (SDC4), mRNA.
fundc1	0.281	-3.554	FUN14 domain containing 1 (FUNDC1), mRNA.
lrpap1	0.281	-3.559	low density lipoprotein receptor-related protein associated protein 1 (LRPAP1), mRNA.
tnfrsf10d	0.281	-3.559	tumor necrosis factor receptor superfamily, member 10d, decoy with truncated death domain (TNFRSF10D), mRNA.
dusp19	0.281	-3.56	dual specificity phosphatase 19 (DUSP19), mRNA.
iqgap1	0.281	-3.564	IQ motif containing GTPase activating protein 1 (IQGAP1), mRNA.
ndst1	0.281	-3.565	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1 (NDST1), mRNA.
atp6ap2	0.28	-3.571	ATPase, H+ transporting, lysosomal accessory protein 2 (ATP6AP2), mRNA.
rpl23	0.28	-3.576	ribosomal protein L23 (RPL23), mRNA.
fkbp2	0.279	-3.581	FK506 binding protein 2, 13kDa (FKBP2), transcript variant 1, mRNA.
polr2h	0.279	-3.582	polymerase (RNA) II (DNA directed) polypeptide H (POLR2H), mRNA.
c16orf28	0.279	-3.582	chromosome 16 open reading frame 28 (C16orf28), mRNA.
lmcd1	0.279	-3.583	LIM and cysteine-rich domains 1 (LMCD1), mRNA.
fam98a	0.279	-3.586	family with sequence similarity 98, member A (FAM98A), mRNA.
eif3s10	0.279	-3.59	eukaryotic translation initiation factor 3, subunit 10 theta, 150/170kDa (EIF3S10), mRNA.
loc644799	0.278	-3.592	PREDICTED: hypothetical protein LOC644799, transcript variant 1 (LOC644799), mRNA.
ube4b	0.278	-3.593	ubiquitination factor E4B (UFD2 homolog, yeast) (UBE4B), mRNA.
hapln3	0.278	-3.595	hyaluronan and proteoglycan link protein 3 (HAPLN3), mRNA.
znf503	0.278	-3.595	zinc finger protein 503 (ZNF503), mRNA.
tm7sf3	0.278	-3.596	transmembrane 7 superfamily member 3 (TM7SF3), mRNA.

csrp1	0.278	-3.6	cysteine and glycine-rich protein 1 (CSRP1), mRNA.
antxr1	0.278	-3.601	anthrax toxin receptor 1 (ANTXR1), transcript variant 2, mRNA.
polr2g	0.277	-3.606	polymerase (RNA) II (DNA directed) polypeptide G (POLR2G), mRNA.
lrp10	0.277	-3.611	low density lipoprotein receptor-related protein 10 (LRP10), mRNA.
lpp	0.276	-3.617	LIM domain containing preferred translocation partner in lipoma (LPP), mRNA.
arid5b	0.276	-3.618	AT rich interactive domain 5B (MRF1-like) (ARID5B), mRNA.
gnl3	0.276	-3.625	guanine nucleotide binding protein-like 3 (nucleolar) (GNL3), transcript variant 2, mRNA.
ccbe1	0.276	-3.625	collagen and calcium binding EGF domains 1 (CCBE1), mRNA.
phpt1	0.276	-3.625	phosphohistidine phosphatase 1 (PHPT1), mRNA.
tmem185b	0.276	-3.626	transmembrane protein 185B (TMEM185B) on chromosome 2.
sars2	0.276	-3.627	seryl-tRNA synthetase 2, mitochondrial (SARS2), mRNA.
mcm4	0.276	-3.628	MCM4 minichromosome maintenance deficient 4 (S. cerevisiae) (MCM4), transcript variant 1, mRNA.
tm4sf20	0.275	-3.633	transmembrane 4 L six family member 20 (TM4SF20), mRNA.
prtfdc1	0.275	-3.635	phosphoribosyl transferase domain containing 1 (PRTFDC1), mRNA.
cdk4	0.275	-3.639	cyclin-dependent kinase 4 (CDK4), mRNA.
galnt11	0.274	-3.643	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (GalNAc-T11) (GALNT11), mRNA.
sfrs1	0.274	-3.644	splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor) (SFRS1), mRNA.
mgc72080	0.274	-3.644	MGC72080 pseudogene (MGC72080) on chromosome 7.
pqbp1	0.274	-3.65	polyglutamine binding protein 1 (PQBP1), transcript variant 1, mRNA.
tmem17	0.274	-3.652	transmembrane protein 17 (TMEM17), mRNA.
psme2	0.274	-3.653	proteasome (prosome, macropain) activator subunit 2 (PA28 beta) (PSME2), mRNA.
glrx	0.273	-3.657	glutaredoxin (thioltransferase) (GLRX), mRNA.
loc442578	0.273	-3.657	similar to Cohesin subunit SA-3 (Stromal antigen 3) (Stromalin 3) (SCC3 homolog 3) (LOC442578), mRNA.
pde7b	0.273	-3.657	phosphodiesterase 7B (PDE7B), mRNA.
foxf2	0.273	-3.658	forkhead box F2 (FOXF2), mRNA.
set	0.272	-3.671	SET translocation (myeloid leukemia-associated) (SET), mRNA.
tmepai	0.272	-3.676	transmembrane, prostate androgen induced RNA (TMEPAI), transcript variant 4, mRNA.
ubiad1	0.272	-3.679	UbiA prenyltransferase domain containing 1 (UBIAD1), mRNA.
nomo1	0.272	-3.681	NODAL modulator 1 (NOMO1), mRNA.
rpl7a	0.272	-3.681	ribosomal protein L7a (RPL7A), mRNA.
vamp2	0.271	-3.691	vesicle-associated membrane protein 2 (synaptobrevin 2) (VAMP2), mRNA.
olfml2a	0.271	-3.696	olfactomedin-like 2A (OLFML2A), mRNA.
creb1	0.27	-3.698	cAMP responsive element binding protein 1 (CREB1), transcript variant A, mRNA.
bles03	0.27	-3.701	chromosome 11 open reading frame 68 (C11orf68), mRNA.
loc283412	0.269	-3.712	PREDICTED: similar to 60S ribosomal protein L29 (Cell surface heparin binding protein HIP) (LOC283412), mRNA.
hs.554507	0.269	-3.713	primary neuroblastoma cDNA, clone:Nbla10527, full insert sequence
gdi2	0.269	-3.719	GDP dissociation inhibitor 2 (GDI2), mRNA.
rbbp7	0.269	-3.72	retinoblastoma binding protein 7 (RBBP7), mRNA.
kiaa1967	0.269	-3.723	KIAA1967 (KIAA1967), transcript variant 1, mRNA.
rab13	0.268	-3.725	RAB13, member RAS oncogene family (RAB13), mRNA.
ppgb	0.268	-3.731	protective protein for beta-galactosidase (galactosialidosis) (PPGB), mRNA.
c21orf63	0.268	-3.738	chromosome 21 open reading frame 63 (C21orf63), mRNA.

hs.524171	0.267	-3.743	cDNA clone IMAGE:5264735
nploc4	0.267	-3.744	nuclear protein localization 4 homolog (S. cerevisiae) (NPLOC4), mRNA.
loc440737	0.267	-3.744	PREDICTED: similar to 60S ribosomal protein L35 (LOC440737), mRNA.
xrcc2	0.267	-3.745	X-ray repair complementing defective repair in Chinese hamster cells 2 (XRCC2), mRNA.
hspc111	0.267	-3.751	hypothetical protein HSPC111 (HSPC111), mRNA.
tle1	0.267	-3.752	transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila) (TLE1), mRNA.
sema3e	0.266	-3.762	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E (SEMA3E), mRNA.
lrrfip1	0.265	-3.768	leucine rich repeat (in FLII) interacting protein 1 (LRRFIP1), mRNA.
c14orf166	0.265	-3.769	chromosome 14 open reading frame 166 (C14orf166), mRNA.
tpp1	0.265	-3.77	tripeptidyl peptidase I (TPP1), mRNA.
immt	0.265	-3.77	inner membrane protein, mitochondrial (mitofilin) (IMMT), mRNA.
ns4atp2	0.265	-3.772	SAP30-like (SAP30L), mRNA.
slc35e1	0.264	-3.786	solute carrier family 35, member E1 (SLC35E1), mRNA.
mgc2749	0.264	-3.789	chromosome 19 open reading frame 50 (C19orf50), mRNA.
tmem156	0.264	-3.79	transmembrane protein 156 (TMEM156), mRNA.
cdh4	0.264	-3.79	cadherin 4, type 1, R-cadherin (retinal) (CDH4), mRNA.
tor1a	0.264	-3.793	torsin family 1, member A (torsin A) (TOR1A), mRNA.
hak	0.263	-3.798	heart alpha-kinase (HAK), mRNA.
mgc29891	0.263	-3.799	hypothetical protein MGC29891 (MGC29891), mRNA.
rhoc	0.263	-3.802	ras homolog gene family, member C (RHOC), mRNA.
cct3	0.263	-3.803	chaperonin containing TCP1, subunit 3 (gamma) (CCT3), transcript variant 1, mRNA.
slc16a12	0.263	-3.805	solute carrier family 16, member 12 (monocarboxylic acid transporter 12) (SLC16A12), mRNA.
kdelc2	0.262	-3.811	KDEL (Lys-Asp-Glu-Leu) containing 2 (KDELC2), mRNA.
ptdss2	0.262	-3.814	phosphatidylserine synthase 2 (PTDSS2), mRNA.
dkfzp564k14	0.261	-3.825	implantation-associated protein (DKFZp564K142), mRNA.
fkbp14	0.261	-3.826	FK506 binding protein 14, 22 kDa (FKBP14), mRNA.
fthl12	0.261	-3.826	ferritin, heavy polypeptide-like 12 (FTHL12) on chromosome 9.
dmn	0.261	-3.834	desmuslin (DMN), transcript variant A, mRNA.
loc651913	0.261	-3.836	PREDICTED: hypothetical protein LOC651913 (LOC651913), mRNA.
slc39a10	0.261	-3.837	solute carrier family 39 (zinc transporter), member 10 (SLC39A10), mRNA.
calr	0.26	-3.846	calreticulin (CALR), mRNA.
gpr1	0.26	-3.848	G protein-coupled receptor 1 (GPR1), mRNA.
bloc1s1	0.26	-3.85	biogenesis of lysosome-related organelles complex-1, subunit 1 (BLOC1S1), mRNA.
mrps21	0.26	-3.852	mitochondrial ribosomal protein S21 (MRPS21), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
tatdn2	0.259	-3.863	TatD DNase domain containing 2 (TATDN2), mRNA.
ssbp1	0.259	-3.864	single-stranded DNA binding protein 1 (SSBP1), mRNA.
frg1	0.259	-3.864	FSHD region gene 1 (FRG1), mRNA.
ccnb1	0.259	-3.868	cyclin B1 (CCNB1), mRNA.
igfbp3	0.259	-3.868	insulin-like growth factor binding protein 3 (IGFBP3), transcript variant 1, mRNA.
muc13	0.258	-3.869	mucin 13, cell surface associated (MUC13), mRNA.
ccr6	0.258	-3.872	chemokine (C-C motif) receptor 6 (CCR6), transcript variant 2, mRNA.
tdp1	0.258	-3.874	tyrosyl-DNA phosphodiesterase 1 (TDP1), transcript variant 1, mRNA.

bcl3	0.258	-3.881	B-cell CLL/lymphoma 3 (BCL3), mRNA.
timm9	0.257	-3.892	translocase of inner mitochondrial membrane 9 homolog (yeast) (TIMM9), mRNA.
dtwd2	0.256	-3.906	DTW domain containing 2 (DTWD2), mRNA.
loc653314	0.256	-3.906	PREDICTED: similar to ribosomal protein L19, transcript variant 5 (LOC653314), mRNA.
mdc1	0.256	-3.908	mediator of DNA damage checkpoint 1 (MDC1), mRNA.
ociad1	0.256	-3.909	OCIA domain containing 1 (OCIAD1), mRNA.
gabrb3	0.255	-3.916	gamma-aminobutyric acid (GABA) A receptor, beta 3 (GABRB3), transcript variant 1, mRNA.
c20orf29	0.255	-3.92	chromosome 20 open reading frame 29 (C20orf29), mRNA.
znf313	0.255	-3.921	zinc finger protein 313 (ZNF313), mRNA.
dusp14	0.255	-3.922	dual specificity phosphatase 14 (DUSP14), mRNA.
hspc268	0.255	-3.928	hypothetical protein HSPC268 (HSPC268), mRNA.
mtif3	0.254	-3.94	mitochondrial translational initiation factor 3 (MTIF3), mRNA.
ergic1	0.253	-3.956	endoplasmic reticulum-golgi intermediate compartment (ERGIC) 1 (ERGIC1), transcript variant 2, mRNA.
cdk6	0.253	-3.956	cyclin-dependent kinase 6 (CDK6), mRNA.
pygl	0.253	-3.957	phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI) (PYGL), mRNA.
c6orf115	0.252	-3.963	PREDICTED: chromosome 6 open reading frame 115 (C6orf115), mRNA.
raet1g	0.252	-3.963	retinoic acid early transcript 1G (RAET1G), mRNA.
fam119a	0.252	-3.966	family with sequence similarity 119, member A (FAM119A), mRNA.
alpp	0.252	-3.967	alkaline phosphatase, placental (Regan isozyme) (ALPP), mRNA.
stom	0.251	-3.979	stomatin (STOM), transcript variant 1, mRNA.
hla-a	0.251	-3.98	major histocompatibility complex, class I, A (HLA-A), mRNA.
eif4g2	0.251	-3.986	eukaryotic translation initiation factor 4 gamma, 2 (EIF4G2), mRNA.
fthl8	0.251	-3.991	ferritin, heavy polypeptide-like 8 (FTHL8) on chromosome X.
clcf1	0.25	-3.994	cardiotrophin-like cytokine factor 1 (CLCF1), mRNA.
cxorf26	0.249	-4.013	chromosome X open reading frame 26 (CXorf26), mRNA.
c15orf48	0.249	-4.021	chromosome 15 open reading frame 48 (C15orf48), transcript variant 1, mRNA.
gabra5	0.248	-4.027	gamma-aminobutyric acid (GABA) A receptor, alpha 5 (GABRA5), mRNA.
hs.534061	0.247	-4.044	full-length cDNA clone XCL0BB001ZD04 of Neuroblastoma of (human)
msc	0.246	-4.058	musculin (activated B-cell factor-1) (MSC), mRNA.
hspa2	0.246	-4.059	heat shock 70kDa protein 2 (HSPA2), mRNA.
loc399900	0.244	-4.09	hypothetical gene supported by AK093779 (LOC399900), mRNA.
ltv1	0.243	-4.108	LTV1 homolog (S. cerevisiae) (LTV1), mRNA.
crip1	0.243	-4.117	cysteine-rich protein 1 (intestinal) (CRIP1), mRNA.
c17orf79	0.242	-4.139	chromosome 17 open reading frame 79 (C17orf79), mRNA.
pappa	0.241	-4.145	pregnancy-associated plasma protein A, pappalysin 1 (PAPPA), mRNA.
hist2h2aa3	0.241	-4.153	histone cluster 2, H2aa3 (HIST2H2AA3), mRNA.
csrp2	0.24	-4.161	cysteine and glycine-rich protein 2 (CSRP2), mRNA.
pdcd7	0.24	-4.164	programmed cell death 7 (PDCD7), mRNA.
tspan3	0.24	-4.17	tetraspanin 3 (TSPAN3), transcript variant 1, mRNA.
c5orf28	0.238	-4.195	chromosome 5 open reading frame 28 (C5orf28), mRNA.
nlrp8	0.238	-4.208	NLR family, pyrin domain containing 8 (NLRP8), mRNA.
sar1a	0.238	-4.21	SAR1 gene homolog A (S. cerevisiae) (SAR1A), mRNA.

nt5dc2	0.237	-4.218	5'-nucleotidase domain containing 2 (NT5DC2), mRNA.
znf486	0.237	-4.225	PREDICTED: zinc finger protein 486 (ZNF486), mRNA.
becn1	0.237	-4.226	beclin 1 (coiled-coil, myosin-like BCL2 interacting protein) (BECN1), mRNA.
hs.564109	0.236	-4.239	AGENCOURT_8673490 Lupski_sciatic_nerve cDNA clone IMAGE:6199918 5, mRNA sequence
fat	0.236	-4.24	FAT tumor suppressor homolog 1 (Drosophila) (FAT), mRNA.
ahnak	0.235	-4.255	AHNAK nucleoprotein (AHNAK), transcript variant 2, mRNA.
col4a2	0.235	-4.256	collagen, type IV, alpha 2 (COL4A2), mRNA.
ndufb10	0.235	-4.258	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10, 22kDa (NDUFB10), mRNA.
col5a1	0.235	-4.261	collagen, type V, alpha 1 (COL5A1), mRNA.
loc729603	0.234	-4.266	calcium binding protein P22 pseudogene (LOC729603) on chromosome 6.
nutf2	0.234	-4.28	nuclear transport factor 2 (NUTF2), mRNA.
ncoa5	0.232	-4.301	nuclear receptor coactivator 5 (NCOA5), mRNA.
c20orf3	0.232	-4.304	chromosome 20 open reading frame 3 (C20orf3), mRNA.
basp1	0.232	-4.315	brain abundant, membrane attached signal protein 1 (BASP1), mRNA.
decr1	0.23	-4.356	2,4-dienoyl CoA reductase 1, mitochondrial (DECR1), nuclear gene encoding mitochondrial protein, mRNA.
osgin1	0.228	-4.378	oxidative stress induced growth inhibitor 1 (OSGIN1), transcript variant 1, mRNA.
il10	0.228	-4.384	interleukin 10 (IL10), mRNA.
hist1h2bk	0.228	-4.39	histone cluster 1, H2bk (HIST1H2BK), mRNA.
ncor2	0.228	-4.394	nuclear receptor co-repressor 2 (NCOR2), transcript variant 1, mRNA.
hhex	0.227	-4.397	hematopoietically expressed homeobox (HHEX), mRNA.
znf69	0.227	-4.401	zinc finger protein 69 (ZNF69), mRNA.
pvrl3	0.227	-4.404	poliovirus receptor-related 3 (PVRL3), mRNA.
rab3b	0.227	-4.407	RAB3B, member RAS oncogene family (RAB3B), mRNA.
zmat3	0.226	-4.431	zinc finger, matrin type 3 (ZMAT3), transcript variant 2, mRNA.
psmc5	0.225	-4.437	proteasome (prosome, macropain) 26S subunit, ATPase, 5 (PSMC5), mRNA.
sox4	0.225	-4.444	SRY (sex determining region Y)-box 4 (SOX4), mRNA.
strap	0.225	-4.444	serine/threonine kinase receptor associated protein (STRAP), mRNA.
ube2g2	0.225	-4.445	ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast) (UBE2G2), transcript variant 2, mRNA.
fahd1	0.224	-4.457	fumarylacetoacetate hydrolase domain containing 1 (FAHD1), transcript variant 2, mRNA.
gcn1l1	0.224	-4.465	GCN1 general control of amino-acid synthesis 1-like 1 (yeast) (GCN1L1), mRNA.
mgc4093	0.222	-4.495	hypothetical protein MGC4093 (MGC4093), mRNA.
c3orf59	0.221	-4.521	chromosome 3 open reading frame 59 (C3orf59), mRNA.
sgk	0.221	-4.533	serum/glucocorticoid regulated kinase (SGK), mRNA.
rpl23a	0.221	-4.535	ribosomal protein L23a (RPL23A), mRNA.
eid2b	0.219	-4.573	EP300 interacting inhibitor of differentiation 2B (EID2B), mRNA.
cirbp	0.217	-4.611	cold inducible RNA binding protein (CIRBP), mRNA.
col5a2	0.215	-4.655	collagen, type V, alpha 2 (COL5A2), mRNA.
mnt	0.214	-4.663	MAX binding protein (MNT), mRNA.
lrap	0.214	-4.672	leukocyte-derived arginine aminopeptidase (LRAP), mRNA.
cfdp1	0.214	-4.674	craniofacial development protein 1 (CFDP1), mRNA.
mrpl38	0.213	-4.688	mitochondrial ribosomal protein L38 (MRPL38), nuclear gene encoding mitochondrial protein, mRNA.
tgm2	0.212	-4.706	transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase) (TGM2), transcript variant 2, mRNA.

porcn	0.212	-4.708	porcupine homolog (Drosophila) (PORCN), transcript variant C, mRNA.
alcam	0.212	-4.717	activated leukocyte cell adhesion molecule (ALCAM), mRNA.
man2b2	0.212	-4.722	mannosidase, alpha, class 2B, member 2 (MAN2B2), mRNA.
kcnj6	0.211	-4.729	potassium inwardly-rectifying channel, subfamily J, member 6 (KCNJ6), mRNA.
pcyox1	0.21	-4.759	prenylcysteine oxidase 1 (PCYOX1), mRNA.
c14orf24	0.21	-4.764	chromosome 14 open reading frame 24 (C14orf24), mRNA.
nbpf10	0.21	-4.772	PREDICTED: neuroblastoma breakpoint family, member 10, transcript variant 7 (NBPF10), mRNA.
glipr1	0.209	-4.776	GLI pathogenesis-related 1 (glioma) (GLIPR1), mRNA.
rabepk	0.209	-4.785	Rab9 effector protein with kelch motifs (RABEPK), mRNA.
col4a1	0.209	-4.785	collagen, type IV, alpha 1 (COL4A1), mRNA.
zbed5	0.208	-4.807	zinc finger, BED-type containing 5 (ZBED5), mRNA.
rcp9	0.208	-4.807	calcitonin gene-related peptide-receptor component protein (RCP9), mRNA.
fstl1	0.207	-4.828	follistatin-like 1 (FSTL1), mRNA.
tnfsf7	0.205	-4.868	tumor necrosis factor (ligand) superfamily, member 7 (TNFSF7), mRNA.
fxyd5	0.204	-4.906	FXYD domain containing ion transport regulator 5 (FXYD5), transcript variant 2, mRNA.
4-Mar	0.201	-4.969	membrane-associated ring finger (C3HC4) 4 (MARCH4), mRNA.
c16orf53	0.201	-4.97	chromosome 16 open reading frame 53 (C16orf53), mRNA.
meg3	0.199	-5.038	PREDICTED: maternally expressed 3, transcript variant 74 (MEG3), misc RNA.
acta2	0.196	-5.105	actin, alpha 2, smooth muscle, aorta (ACTA2), mRNA.
vtn	0.196	-5.108	vitronectin (serum spreading factor, somatomedin B, complement S-protein) (VTN), mRNA.
znf430	0.195	-5.115	zinc finger protein 430 (ZNF430), mRNA.
thbs1	0.195	-5.139	thrombospondin 1 (THBS1), mRNA.
cspg2	0.194	-5.162	versican (VCAN), mRNA.
hist1h1c	0.193	-5.19	histone cluster 1, H1c (HIST1H1C), mRNA.
wsb1	0.192	-5.205	WD repeat and SOCS box-containing 1 (WSB1), transcript variant 3, mRNA.
pebp1	0.192	-5.222	phosphatidylethanolamine binding protein 1 (PEBP1), mRNA.
spock	0.191	-5.226	sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) (SPOCK), mRNA.
rbp1	0.191	-5.24	retinol binding protein 1, cellular (RBP1), mRNA.
loc613037	0.191	-5.249	nuclear pore complex interacting protein pseudogene (LOC613037) on chromosome 16.
flj46309	0.19	-5.256	hypothetical protein LOC649598 (FLJ46309), mRNA.
nop5/nop58	0.189	-5.278	nucleolar protein NOP5/NOP58 (NOP5/NOP58), mRNA.
ris1	0.185	-5.394	Ras-induced senescence 1 (RIS1), mRNA.
loc441087	0.185	-5.406	hypothetical gene supported by AK125735 (LOC441087), mRNA.
loc645895	0.182	-5.482	PREDICTED: hypothetical protein LOC645895 (LOC645895), mRNA.
vat1	0.182	-5.507	vesicle amine transport protein 1 homolog (T californica) (VAT1), mRNA.
c14orf85	0.181	-5.534	chromosome 14 open reading frame 85 (C14orf85) on chromosome 14.
chp	0.178	-5.606	calcium binding protein P22 (CHP), mRNA.
hs.371609	0.173	-5.778	cDNA clone IMAGE:5261213
c14orf4	0.173	-5.78	chromosome 14 open reading frame 4 (C14orf4), mRNA.
tagln	0.166	-6.031	transgelin (TAGLN), transcript variant 1, mRNA.
pofut1	0.165	-6.055	protein O-fucosyltransferase 1 (POFUT1), transcript variant 2, mRNA.
htra1	0.161	-6.205	HtrA serine peptidase 1 (HTRA1), mRNA.

flj44124	0.157	-6.383	hypothetical protein LOC641737 (FLJ44124), mRNA.
cryab	0.154	-6.473	crystallin, alpha B (CRYAB), mRNA.
psmd12	0.152	-6.588	proteasome (prosome, macropain) 26S subunit, non-ATPase, 12 (PSMD12), mRNA. XM_946055 XM_946058
gls	0.151	-6.616	glutaminase (GLS), mRNA.
apoe	0.149	-6.695	apolipoprotein E (APOE), mRNA.
arl16	0.145	-6.898	PREDICTED: ADP-ribosylation factor-like 16 (ARL16), mRNA.
cyp1b1	0.143	-6.997	cytochrome P450, family 1, subfamily B, polypeptide 1 (CYP1B1), mRNA.
hmox1	0.14	-7.124	heme oxygenase (decycling) 1 (HMOX1), mRNA.
c8orf45	0.135	-7.408	chromosome 8 open reading frame 45 (C8orf45), mRNA.
nnmt	0.134	-7.488	nicotinamide N-methyltransferase (NNMT), mRNA.
abca1	0.133	-7.532	ATP-binding cassette, sub-family A (ABC1), member 1 (ABCA1), mRNA.
igfl3	0.123	-8.127	IGF-like family member 3 (IGFL3), mRNA.
anpep	0.109	-9.187	alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA.