Gene Name	Description and ID		DNA methylatio	
	<u> </u>	ES-mCpG	NSC-mCpG	CM-mCpG
NANOG	Nanog homeobox [Source:HGNC Symbol;Acc:20857]	0.04	0.06	0.63
ERPINB1	serpin peptidase inhibitor, clade B (ovalbumin), member 1 [Source:HGNC Symbol;Acc:3311]	0.16	0.13	0.64
BR1	carbonyl reductase 1 [Source:HGNC Symbol;Acc:1548]	0.10	0.15	0.53
M20D2	peptidase M20 domain containing 2 [Source:HGNC Symbol;Acc:21408]	0.19	0.29	0.66
KAP1	src kinase associated phosphoprotein 1 [Source:HGNC Symbol;Acc:15605]	0.32	0.26	0.68
	ATP-binding cassette, sub-family B (MDR/TAP), member 4 [Source:HGNC			
ABCB4	Symbol;Acc:45]	0.08	0.02	0.43
USC3	tumor suppressor candidate 3 [Source:HGNC Symbol;Acc:30242]	0.52	0.38	0.87
IORMAD2	HORMA domain containing 2 [Source:HGNC Symbol;Acc:28383]	0.47	0.39	0.81
OXL3	lysyl oxidase-like 3 [Source:HGNC Symbol;Acc:13869]	0.18	0.17	0.51
	UDP-N-acteylglucosamine pyrophosphorylase 1 [Source:HGNC			
JAP1	Symbol;Acc:12457]	0.53	0.26	0.84
PLX2	complexin 2 [Source:HGNC Symbol;Acc:2310]	0.39	0.23	0.69
CDC81 REB5	coiled-coil domain containing 81 [Source:HGNC Symbol;Acc:26281] cAMP responsive element binding protein 5 [Source:HGNC Symbol;Acc:16844]	0.3 0.13	0.16 0.07	0.59 0.4
PR113	G protein-coupled receptor 113 [Source:HGNC Symbol;Acc:18889]	0.13	0.07	0.4
MN	amnion associated transmembrane protein [Source:HGNC Symbol;Acc:14604]	0.03	0.03	0.07
RELID2	PRELI domain containing 2 [Source:HGNC Symbol;Acc:28306]	0.01	0.02	0.28
ILLIDZ	protein tyrosine phosphatase-like A domain containing 2 [Source:HGNC	0.01	0.02	0.20
TPLAD2	Symbol;Acc:20920]	0.22	0.43	0.68
ISBP1L1	heat shock factor binding protein 1-like 1 [Source:HGNC Symbol;Acc:37243]	0.57	0.53	0.82
	,,,			
TP2B2	ATPase, Ca++ transporting, plasma membrane 2 [Source:HGNC Symbol;Acc:815]	0.37	0.33	0.62
	protein kinase, cAMP-dependent, regulatory, type II, beta [Source:HGNC			
PRKAR2B	Symbol;Acc:9392]	0.4	0.52	0.77
DF15	growth differentiation factor 15 [Source:HGNC Symbol;Acc:30142]	0.35	0.21	0.6
4GNT	alpha-1,4-N-acetylglucosaminyltransferase [Source:HGNC Symbol;Acc:17968]	0.67	0.5	0.92
	major facilitator superfamily domain containing 2B [Source:HGNC			
MFSD2B	Symbol;Acc:37207]	0.27	0.5	0.75
	chromodomain helicase DNA binding protein 7 [Source:HGNC			
CHD7	Symbol;Acc:20626]	0.16	0.13	0.41
DHD	lactate dehydrogenase D [Source:HGNC Symbol;Acc:19708]	0.39	0.43	0.67
TDD C4	protein tyrosine phosphatase domain containing 1 [Source:HGNC	0.4	0.20	0.64
TPDC1	Symbol;Acc:30184]	0.4	0.39	0.64
TEA	TRAF-interacting protein with forkhead-associated domain [Source:HGNC	0.27	0.21	0.51
TFA OXD2	Symbol;Acc:19075] forkhead box D2 [Source:HGNC Symbol;Acc:3803]	0.27	0.21	0.51
UNDZ	PAS domain containing serine/threonine kinase [Source:HGNC	0.26	0.29	0.32
ASK	Symbol;Acc:17270]	0.44	0.4	0.67
IOXB2	homeobox B2 [Source:HGNC Symbol;Acc:5113]	0.13	0.15	0.38
RBM43	RNA binding motif protein 43 [Source:HGNC Symbol;Acc:24790]	0.2	0.24	0.47
DCL3	phosducin-like 3 [Source:HGNC Symbol;Acc:28860]	0.47	0.29	0.7
ERHL2	serine hydrolase-like 2 [Source:HGNC Symbol;Acc:29446]	0.02	0.12	0.34
	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin),			
ERPINA11	member 11 [Source:HGNC Symbol;Acc:19193]	0.09	0.15	0.37
RHGAP9	Rho GTPase activating protein 9 [Source:HGNC Symbol;Acc:14130]	0.78	0.76	1
	olfactory receptor, family 10, subfamily A, member 5 [Source:HGNC			
DR10A5	Symbol;Acc:15131]	0.72	0.59	0.94
RAF2	PRA1 domain family, member 2 [Source:HGNC Symbol;Acc:28911]	0.44	0.46	0.67
LCL2	phospholipase C-like 2 [Source:HGNC Symbol;Acc:9064]	0.15	0.25	0.46
//AT1A	methionine adenosyltransferase I, alpha [Source:HGNC Symbol;Acc:6903]	0.27	0.41	0.62
	major histocompatibility complex, class I, H (pseudogene) [Source:HGNC	0.00	0.47	0.00
ILA-H	Symbol;Acc:4965]	0.09	0.17	0.38
16orf54	chromosome 16 open reading frame 54 [Source:HGNC Symbol;Acc:26649]	0.64	0.65	0.86
DBP2B	odorant binding protein 2B [Source:HGNC Symbol;Acc:23381] zinc finger protein 239 [Source:HGNC Symbol;Acc:13031]	0.62	0.64	0.85
NF239 ICK	hemopoietic cell kinase [Source:HGNC Symbol;Acc:13031]	0.6 0.23	0.69 0.35	0.89 0.55
PATA13	spermatogenesis associated 13 [Source:HGNC Symbol;Acc:23222]	0.23	0.35	0.33
	GRP1 (general receptor for phosphoinositides 1)-associated scaffold protein	5.15	0.07	0.33
GRASP	[Source:HGNC Symbol;Acc:18707]	0.41	0.29	0.61
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	C1g and tumor necrosis factor related protein 1 [Source:HGNC	02	0.23	0.01
1QTNF1	Symbol;Acc:14324]	0.33	0.12	0.53
LIT1	slit homolog 1 (Drosophila) [Source:HGNC Symbol;Acc:11085]	0.17	0.17	0.37
GAT2	diacylglycerol O-acyltransferase 2 [Source:HGNC Symbol;Acc:16940]	0.11	0.15	0.35
NT5DC1	5'-nucleotidase domain containing 1 [Source:HGNC Symbol;Acc:21556]	0.45	0.47	0.66
YS1	cystin 1 [Source:HGNC Symbol;Acc:18525]	0.37	0.35	0.56
	fasciculation and elongation protein zeta 1 (zygin I) [Source:HGNC			
EZ1	Symbol;Acc:3659]	0.51	0.58	0.77
	BTB and CNC homology 1, basic leucine zipper transcription factor 2			
BACH2	[Source:HGNC Symbol;Acc:14078]	0.51	0.58	0.77
	leucine rich repeat transmembrane neuronal 3 [Source:HGNC			
RRTM3	Symbol;Acc:19410]	0.59	0.74	0.93

	major facilitator superfamily domain containing 6-like [Source:HGNC			
MFSD6L	Symbol;Acc:26656]	0.42	0.58	0.77
LRRC4	leucine rich repeat containing 4 [Source:HGNC Symbol;Acc:15586]	0.21	0.21	0.4
SLC26A4	solute carrier family 26, member 4 [Source:HGNC Symbol;Acc:8818]	0.42	0.41	0.61
SSFA2	sperm specific antigen 2 [Source:HGNC Symbol;Acc:11319]	0.3	0.26	0.49
ESRRG	estrogen-related receptor gamma [Source:HGNC Symbol;Acc:3474]	0.01	0.1	0.29
	cyclin-dependent kinase inhibitor 1B (p27, Kip1) [Source:HGNC			
CDKN1B	Symbol;Acc:1785]	0.16	0.06	0.35
STXBP5L	syntaxin binding protein 5-like [Source:HGNC Symbol;Acc:30757]	0.11	0.23	0.42
	potassium inwardly-rectifying channel, subfamily J, member 12 [Source:HGNC			
KCNJ12	Symbol;Acc:6258]	0.64	0.67	0.86
TCIDCA	T-cell, immune regulator 1, ATPase, H+ transporting, lysosomal V0 subunit A3	0.70	0.60	0.07
TCIRG1	[Source:HGNC Symbol;Acc:11647]	0.78	0.68	0.97
CXCL5	chemokine (C-X-C motif) ligand 5 [Source:HGNC Symbol;Acc:10642] ST3 beta-galactoside alpha-2,3-sialyltransferase 4 [Source:HGNC	0.44	0.55	0.74
ST3GAL4	Symbol;Acc:10864]	0.73	0.71	0.91
NOXO1	NADPH oxidase organizer 1 [Source:HGNC Symbol;Acc:19404]	0.73	0.71	0.91
SLC46A3	solute carrier family 46, member 3 [Source:HGNC Symbol;Acc:27501]	0.72	0.07	0.9
SYN2	synapsin II [Source:HGNC Symbol;Acc:11495]	0.36	0.47	0.65
MYH2	myosin, heavy chain 2, skeletal muscle, adult [Source:HGNC Symbol;Acc:7572]	0.47	0.32	0.65
TMEM171	transmembrane protein 171 [Source:HGNC Symbol;Acc:27031]	0.4	0.63	0.81
PITX2	paired-like homeodomain 2 [Source:HGNC Symbol;Acc:9005]	0.13	0.15	0.33
CLDN9	claudin 9 [Source:HGNC Symbol;Acc:2051]	0.02	0.21	0.39
GOLGA1	golgin A1 [Source:HGNC Symbol;Acc:4424]	0.12	0.13	0.31
HPS1	Hermansky-Pudlak syndrome 1 [Source:HGNC Symbol;Acc:5163]	0.42	0.39	0.6
NEURL	neuralized homolog (Drosophila) [Source:HGNC Symbol;Acc:7761]	0.37	0.41	0.59
TP53BP1	tumor protein p53 binding protein 1 [Source:HGNC Symbol;Acc:11999]	0.09	0.13	0.31
DSP	desmoplakin [Source:HGNC Symbol;Acc:3052]	0.14	0.2	0.38
	potassium channel tetramerisation domain containing 17 [Source:HGNC			
KCTD17	Symbol;Acc:25705]	0.24	0.18	0.42
TM4SF18	transmembrane 4 L six family member 18 [Source:HGNC Symbol;Acc:25181]	0.1	0.24	0.42
ZMAT1	zinc finger, matrin-type 1 [Source:HGNC Symbol;Acc:29377]	0.54	0.54	0.72
RBP4	retinol binding protein 4, plasma [Source:HGNC Symbol;Acc:9922]	0.44	0.51	0.69
	major histocompatibility complex, class II, DP alpha 1 [Source:HGNC			
HLA-DPA1	Symbol;Acc:4938]	0.69	0.7	0.87
AQP9	aquaporin 9 [Source:HGNC Symbol;Acc:643]	0.61	0.55	0.78
RUSC1	RUN and SH3 domain containing 1 [Source:HGNC Symbol;Acc:17153]	0.44	0.51	0.68
LYPD6	LY6/PLAUR domain containing 6 [Source:HGNC Symbol;Acc:28751]	0.34	0.48	0.65
ACPL2	acid phosphatase-like 2 [Source:HGNC Symbol;Acc:26303]	0.35	0.61	0.78
	ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1			
ATP8A1	[Source:HGNC Symbol;Acc:13531]	0	0	0.17
CNIKCDO	connector enhancer of kinase suppressor of Ras 2 [Source:HGNC	0.45	0.11	0.22
CNKSR2	Symbol;Acc:19701]	0.15	0.11	0.32
NUDT19	nudix (nucleoside diphosphate linked moiety X)-type motif 19 [Source:HGNC Symbol;Acc:32036]	0.07	0.15	0.32
TMEM30B	transmembrane protein 30B [Source:HGNC Symbol;Acc:27254]	0.07	0.15	0.32
TIVILIVISOD	DEAD (Asp-Glu-Ala-Asp) box polypeptide 60-like [Source:HGNC	0.04	0.13	0.52
DDX60L	Symbol;Acc:26429]	0.37	0.45	0.62
DDXOOL	calcium channel, voltage-dependent, gamma subunit 4 [Source:HGNC	0.57	0.43	0.02
CACNG4	Symbol;Acc:1408]	0.13	0.24	0.41
SHISA5	shisa homolog 5 (Xenopus laevis) [Source:HGNC Symbol;Acc:30376]	0.76	0.77	0.94
	major histocompatibility complex, class II, DR alpha [Source:HGNC			
HLA-DRA	Symbol;Acc:4947]	0.47	0.52	0.69
	protein phosphatase 3, catalytic subunit, gamma isozyme [Source:HGNC			
PPP3CC	Symbol;Acc:9316]	0.56	0.67	0.84
ACVR1C	activin A receptor, type IC [Source:HGNC Symbol;Acc:18123]	0.4	0.4	0.56
TNNI3	troponin I type 3 (cardiac) [Source:HGNC Symbol;Acc:11947]	0.72	0.71	0.88
NCKAP1L	NCK-associated protein 1-like [Source:HGNC Symbol;Acc:4862]	0.75	0.74	0.91
	tumor necrosis factor receptor superfamily, member 8 [Source:HGNC			
TNFRSF8	Symbol;Acc:11923]	0.1	0.12	0.28
	gamma-aminobutyric acid (GABA) A receptor, alpha 5 [Source:HGNC			
GABRA5	Symbol;Acc:4079]	0.59	0.63	0.79
VPREB3	pre-B lymphocyte 3 [Source:HGNC Symbol;Acc:12710]	0.69	0.73	0.89
MLKL	mixed lineage kinase domain-like [Source:HGNC Symbol;Acc:26617]	0.03	0.11	0.27
GCC2	GRIP and coiled-coil domain containing 2 [Source:HGNC Symbol;Acc:23218]	0.49	0.41	0.65
C9orf172	chromosome 9 open reading frame 172 [Source:HGNC Symbol;Acc:37284]	0.7	0.61	0.86
חאטטבי	polycystic kidney disease (polycystin) and REJ homolog (sperm receptor for egg	0.20	0.53	0.50
PKDREJ	jelly homolog, sea urchin) [Source:HGNC Symbol;Acc:9015]	0.38	0.52	0.68
SPINT3	serine peptidase inhibitor, Kunitz type, 3 [Source:HGNC Symbol;Acc:11248]	0.72	0.52	0.88
INHBE NMI	inhibin, beta E [Source:HGNC Symbol;Acc:24029]	0.28	0.51	0.67
INIVII	N-myc (and STAT) interactor [Source:HGNC Symbol;Acc:7854] A kinase (PRKA) anchor protein 2 [Source:HGNC Symbol;Acc:372]	0.04	0.38 0.04	0.54 0.2
		0.05	0.04	0.2
AKAP2			0.12	U 36
	synaptotagmin VII [Source:HGNC Symbol;Acc:11514]	0.22	0.13	0.38
AKAP2 SYT7	synaptotagmin VII [Source:HGNC Symbol;Acc:11514] transforming, acidic coiled-coil containing protein 2 [Source:HGNC	0.22		
AKAP2	synaptotagmin VII [Source:HGNC Symbol;Acc:11514]		0.13 0.14 0.42	0.38 0.35 0.58

PCDHB16	protocodharin hata 16 [CoursestICNC Combal Apart 45.45]	0	0.05	0.31
	protocadherin beta 16 [Source:HGNC Symbol;Acc:14546]	0 0.32	0.05	0.21 0.48
PRICKLE2	prickle homolog 2 (Drosophila) [Source:HGNC Symbol;Acc:20340] ribonuclease L (2',5'-oligoisoadenylate synthetase-dependent) [Source:HGNC	0.32	0.25	0.48
RNASEL	Symbol;Acc:10050]	0.43	0.31	0.59
GPBAR1	G protein-coupled bile acid receptor 1 [Source:HGNC Symbol;Acc:19680]	0.43	0.66	0.97
CTGF	connective tissue growth factor [Source:HGNC Symbol;Acc:2500]	0.18	0.17	0.33
3101	potassium voltage-gated channel, subfamily H (eag-related), member 3	0.10	0.17	0.55
CNH3	[Source:HGNC Symbol;Acc:6252]	0.3	0.29	0.45
BEGAIN	brain-enriched guanylate kinase-associated [Source:HGNC Symbol;Acc:24163]	0.6	0.61	0.76
PYCARD	PYD and CARD domain containing [Source:HGNC Symbol;Acc:16608]	0.62	0.63	0.78
OTUB2	OTU domain, ubiquitin aldehyde binding 2 [Source:HGNC Symbol;Acc:20351]	0.64	0.62	0.79
SMD1	LSM domain containing 1 [Source:HGNC Symbol;Acc:28212]	0.05	0.08	0.23
GLL1	immunoglobulin lambda-like polypeptide 1 [Source:HGNC Symbol;Acc:5870]	0.72	0.68	0.87
SOX15	SRY (sex determining region Y)-box 15 [Source:HGNC Symbol;Acc:11196]	0.57	0.61	0.76
GJB4	gap junction protein, beta 4, 30.3kDa [Source:HGNC Symbol;Acc:4286]	0.74	0.67	0.89
	olfactory receptor, family 4, subfamily C, member 11 [Source:HGNC			
OR4C11	Symbol;Acc:15167]	0.85	0.67	1
GPR85	G protein-coupled receptor 85 [Source:HGNC Symbol;Acc:4536]	0.09	0.09	0.24
ТВХ5	T-box 5 [Source:HGNC Symbol;Acc:11604]	0.03	0.02	0.18
DLEU7	deleted in lymphocytic leukemia, 7 [Source:HGNC Symbol;Acc:17567]	0.06	0.04	0.21
HOXB3	homeobox B3 [Source:HGNC Symbol;Acc:5114]	0.06	0.03	0.21
EX7	peroxisomal biogenesis factor 7 [Source:HGNC Symbol;Acc:8860]	0.3	0.34	0.49
GUCA2B	guanylate cyclase activator 2B (uroguanylin) [Source:HGNC Symbol;Acc:4683]	0.79	0.76	0.94
PDE1B	phosphodiesterase 1B, calmodulin-dependent [Source:HGNC Symbol;Acc:8775]	0.5	0.55	0.7
NF674	zinc finger protein 674 [Source:HGNC Symbol;Acc:17625]	0.55	0.49	0.7
CNPY1	canopy 1 homolog (zebrafish) [Source:HGNC Symbol;Acc:27786]	0.42	0.36	0.56
NES	nestin [Source:HGNC Symbol;Acc:7756]	0.02	0.01	0.16
NF700	zinc finger protein 700 [Source:HGNC Symbol;Acc:25292]	0.08	0.07	0.22
	solute carrier family 3 (cystine, dibasic and neutral amino acid transporters,			
	activator of cystine, dibasic and neutral amino acid transport), member 1			
SLC3A1	[Source:HGNC Symbol;Acc:11025]	0.86	0.85	1
	solute carrier family 7 (neutral amino acid transporter light chain, asc system),			
LC7A10	member 10 [Source:HGNC Symbol;Acc:11058]	0.7	0.68	0.84
EMD3	LEM domain containing 3 [Source:HGNC Symbol;Acc:28887]	0.66	0.64	0.8
RF2	interferon regulatory factor 2 [Source:HGNC Symbol;Acc:6117]	0.08	0.05	0.22
14orf93	chromosome 14 open reading frame 93 [Source:HGNC Symbol;Acc:20162]	0.1	0.13	0.27
CABP7	calcium binding protein 7 [Source:HGNC Symbol;Acc:20834]	0.64	0.61	0.78
CHKA	choline kinase alpha [Source:HGNC Symbol;Acc:1937]	0.36	0.39	0.53
	collagen-like tail subunit (single strand of homotrimer) of asymmetric			
COLQ	acetylcholinesterase [Source:HGNC Symbol;Acc:2226]	0.86	0.82	1
POA2	apolipoprotein A-II [Source:HGNC Symbol;Acc:601]	0.82	0.86	1
DLEC1	deleted in lung and esophageal cancer 1 [Source:HGNC Symbol;Acc:2899]	0.48	0.52	0.66
	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter),	0.50	0.70	0.07
SLC17A7	member 7 [Source:HGNC Symbol;Acc:16704]	0.68	0.73	0.87
ADAM8 QCF2	ADAM metallopeptidase domain 8 [Source:HGNC Symbol:Acc:215]	0.66 0.86	0.6	0.8 1
QCF2	IQ motif containing F2 [Source:HGNC Symbol;Acc:31815]	0.80	0.79	1
CE71.1	transcription factor 7-like 1 (T-cell specific, HMG-box) [Source:HGNC		0.66	0.8
CF7L1 PRRT1	Symbol;Acc:11640]	0.50		0.0
IVIVII		0.58	0.66	
MD2	proline-rich transmembrane protein 1 [Source:HGNC Symbol;Acc:13943]	0.08	0.19	0.33
	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335]	0.08 0.29	0.19 0.1	0.33 0.43
AHNAK2	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125]	0.08 0.29 0.19	0.19 0.1 0.38	0.33 0.43 0.52
AHNAK2	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539]	0.08 0.29	0.19 0.1	0.33 0.43
AHNAK2 RBM24	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC	0.08 0.29 0.19 0.66	0.19 0.1 0.38 0.03	0.33 0.43 0.52 0.8
AHNAK2 RBM24	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718]	0.08 0.29 0.19	0.19 0.1 0.38	0.33 0.43 0.52
AHNAK2 RBM24 APCDD1	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC	0.08 0.29 0.19 0.66	0.19 0.1 0.38 0.03	0.33 0.43 0.52 0.8
AHNAK2 RBM24 APCDD1 FNFSF11	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926]	0.08 0.29 0.19 0.66 0.21	0.19 0.1 0.38 0.03 0.19	0.33 0.43 0.52 0.8 0.35
AHNAK2 RBM24 APCDD1 FNFSF11 ACADL	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88]	0.08 0.29 0.19 0.66 0.21 0.13	0.19 0.1 0.38 0.03 0.19 0.17 0.22	0.33 0.43 0.52 0.8 0.35 0.31 0.36
AHNAK2 RBM24 APCDD1 FNFSF11 ACADL	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926]	0.08 0.29 0.19 0.66 0.21	0.19 0.1 0.38 0.03 0.19	0.33 0.43 0.52 0.8 0.35
AHNAK2 RBM24 APCDD1 FNFSF11 ACADL RAB21	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC	0.08 0.29 0.19 0.66 0.21 0.13	0.19 0.1 0.38 0.03 0.19 0.17 0.22	0.33 0.43 0.52 0.8 0.35 0.31 0.36
AHNAK2 RBM24 APCDD1 INFSF11 ACADL RAB21 RIPK3 C11orf70	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42
AHNAK2 RBM24 APCDD1 INFSF11 ACADL RAB21 RIPK3 C11orf70 IAPBP	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 CAPBP	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP FRIM21	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42
AHNAK2 RBM24 APCDD1 FNFSF11 ACADL RAB21 RIPK3 C11orf70 FAPBP FRIM21	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:1926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42
AHNAK2 RBM24 APCDD1 INFSF11 ACADL RAB21 RIPK3 C110rf70 FAPBP FRIM21 NPP4A	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42
AHNAK2 RBM24 APCDD1 INFSF11 ACADL RAB21 RIPK3 C11orf70 IAPBP IRIM21 NPP4A	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19
AHNAK2 RBM24 APCDD1 INFSF11 ACADL RAB21 RIPK3 C11orf70 IAPBP IRIM21 NPP4A POLE4 ARID3C	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06 0.04 0.71	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19 0.17 0.84
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2 TICAM2	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209] contactin 2 (axonal) [Source:HGNC Symbol;Acc:2172]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19 0.17 0.84 0.75
EMP3 AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2 TICAM2 TICAM2 TICAM1 LRRC32	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209] contactin 2 (axonal) [Source:HGNC Symbol;Acc:2172] toll-like receptor adaptor molecule 2 [Source:HGNC Symbol;Acc:21354]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06 0.04 0.71 0.61 0.08	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.91 0.19 0.17 0.84 0.75 0.21
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2 TICAM2 TICAM2 TICAM1 LRRC32	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11566] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209] contactin 2 (axonal) [Source:HGNC Symbol;Acc:21354] toll-like receptor adaptor molecule 1 [Source:HGNC Symbol;Acc:18348]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06 0.04 0.71 0.61 0.08 0.04	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.78 0.06 0.04 0.71 0.62 0.06 0.06	0.33 0.43 0.52 0.8 0.35 0.31 0.48 0.42 0.42 0.91 0.19 0.17 0.84 0.75 0.21 0.19
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2 TICAM2 TICAM2 TICAM1	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11366] tripartite motif containing 21 [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:11375] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209] contactin 2 (axonal) [Source:HGNC Symbol;Acc:21354] toll-like receptor adaptor molecule 1 [Source:HGNC Symbol;Acc:18348] leucine rich repeat containing 32 [Source:HGNC Symbol;Acc:4161]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06 0.04 0.71 0.61 0.08 0.04 0.2	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06 0.04 0.71 0.62 0.06 0.06 0.17	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19 0.17 0.84 0.75 0.21 0.19 0.33
AHNAK2 RBM24 APCDD1 TNFSF11 ACADL RAB21 RIPK3 C11orf70 TAPBP TRIM21 INPP4A POLE4 ARID3C CNTN2 TICAM2 TICAM2 TICAM1 LRRC32 DNLZ	epithelial membrane protein 3 [Source:HGNC Symbol;Acc:3335] AHNAK nucleoprotein 2 [Source:HGNC Symbol;Acc:20125] RNA binding motif protein 24 [Source:HGNC Symbol;Acc:21539] adenomatosis polyposis coli down-regulated 1 [Source:HGNC Symbol;Acc:15718] tumor necrosis factor (ligand) superfamily, member 11 [Source:HGNC Symbol;Acc:11926] acyl-CoA dehydrogenase, long chain [Source:HGNC Symbol;Acc:88] RAB21, member RAS oncogene family [Source:HGNC Symbol;Acc:18263] receptor-interacting serine-threonine kinase 3 [Source:HGNC Symbol;Acc:10021] chromosome 11 open reading frame 70 [Source:HGNC Symbol;Acc:28188] TAP binding protein (tapasin) [Source:HGNC Symbol;Acc:11312] inositol polyphosphate-4-phosphatase, type I, 107kDa [Source:HGNC Symbol;Acc:6074] polymerase (DNA-directed), epsilon 4, accessory subunit [Source:HGNC Symbol;Acc:18755] AT rich interactive domain 3C (BRIGHT-like) [Source:HGNC Symbol;Acc:21209] contactin 2 (axonal) [Source:HGNC Symbol;Acc:2172] toll-like receptor adaptor molecule 2 [Source:HGNC Symbol;Acc:21354] toll-like receptor adaptor molecule 1 [Source:HGNC Symbol;Acc:1348] leucine rich repeat containing 32 [Source:HGNC Symbol;Acc:4161] DNL-type zinc finger [Source:HGNC Symbol;Acc:33879]	0.08 0.29 0.19 0.66 0.21 0.13 0.14 0.17 0.33 0.22 0.17 0.78 0.06 0.04 0.71 0.61 0.08 0.04 0.2 0.32	0.19 0.1 0.38 0.03 0.19 0.17 0.22 0.04 0.34 0.28 0.28 0.78 0.06 0.04 0.71 0.62 0.06 0.06 0.07 0.36	0.33 0.43 0.52 0.8 0.35 0.31 0.36 0.31 0.48 0.42 0.42 0.91 0.19 0.17 0.84 0.75 0.21 0.19 0.33 0.49

ADCY1	adenylate cyclase 1 (brain) [Source:HGNC Symbol;Acc:232]	0.62	0.68	0.81
FMOD	fibromodulin [Source:HGNC Symbol;Acc:3774]	0.69	0.62	0.82
00514	olfactory receptor, family 5, subfamily V, member 1 [Source:HGNC	0.70	0.74	0.04
OR5V1	Symbol;Acc:13972]	0.78	0.71	0.91
FAIM2	Fas apoptotic inhibitory molecule 2 [Source:HGNC Symbol;Acc:17067]	0.74	0.65	0.87
CN10N1040	small nuclear ribonucleoprotein 48kDa (U11/U12) [Source:HGNC	0.00	0.07	0.40
SNRNP48	Symbol;Acc:21368]	0.36	0.27	0.49
A D A A ATCO	ADAM metallopeptidase with thrombospondin type 1 motif, 8 [Source:HGNC	0.60	0.50	0.04
ADAMTS8	Symbol;Acc:224]	0.68	0.59	0.81
ANP32B	acidic (leucine-rich) nuclear phosphoprotein 32 family, member B [Source:HGNC	0.08	0.18	0.31
ANP32B	Symbol;Acc:16677] ATR hinding assetts cub family C (CETR/MARR), mamber 3 [Source:HCNC]	0.08	0.18	0.31
ABCC3	ATP-binding cassette, sub-family C (CFTR/MRP), member 3 [Source:HGNC Symbol;Acc:54]	0.27	0.17	0.4
C15orf56	chromosome 15 open reading frame 56 [Source:HGNC Symbol;Acc:33868]	0.53	0.17	0.78
C1301130	2-aminoethanethiol (cysteamine) dioxygenase [Source:HGNC	0.55	0.03	0.78
ADO	Symbol;Acc:23506]	0.71	0.56	0.84
7.00	ADAM metallopeptidase with thrombospondin type 1 motif, 2 [Source:HGNC	0.71	0.50	0.04
ADAMTS2	Symbol;Acc:218]	0.31	0.48	0.61
CD72	CD72 molecule [Source:HGNC Symbol;Acc:1696]	0.87	0.69	1
CD/L	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small	0.07	0.03	
CTDSPL	phosphatase-like [Source:HGNC Symbol;Acc:16890]	0.52	0.29	0.65
	tumor necrosis factor (ligand) superfamily, member 10 [Source:HGNC	0.02	0.20	0.00
TNFSF10	Symbol;Acc:11925]	0.54	0.28	0.67
	ATP-binding cassette, sub-family F (GCN20), member 2 [Source:HGNC			
ABCF2	Symbol;Acc:71]	0.28	0.28	0.41
C11orf93	chromosome 11 open reading frame 93 [Source:HGNC Symbol;Acc:26978]	0.19	0.34	0.47
HOXB1	homeobox B1 [Source:HGNC Symbol;Acc:5111]	0.19	0.34	0.47
PAX5	paired box 5 [Source:HGNC Symbol;Acc:8619]	0.56	0.53	0.69
	Ras association (RalGDS/AF-6) domain family member 1 [Source:HGNC			
RASSF1	Symbol;Acc:9882]	0.17	0.42	0.54
STC1	stanniocalcin 1 [Source:HGNC Symbol;Acc:11373]	0.02	0.02	0.14
	protein phosphatase 1, regulatory subunit 16B [Source:HGNC			
PPP1R16B	Symbol;Acc:15850]	0.01	0.02	0.14
	POZ (BTB) and AT hook containing zinc finger 1 [Source:HGNC			
PATZ1	Symbol;Acc:13071]	0.04	0.05	0.17
ACP2	acid phosphatase 2, lysosomal [Source:HGNC Symbol;Acc:123]	0.55	0.55	0.67
GLIPR1L2	GLI pathogenesis-related 1 like 2 [Source:HGNC Symbol;Acc:28592]	0.6	0.6	0.72
	sulfotransferase family, cytosolic, 1C, member 2 [Source:HGNC			
SULT1C2	Symbol;Acc:11456]	0.19	0.18	0.31
ESYT3	extended synaptotagmin-like protein 3 [Source:HGNC Symbol;Acc:24295]	0.6	0.59	0.72
	solute carrier family 36 (proton/amino acid symporter), member 2			
SLC36A2	[Source:HGNC Symbol;Acc:18762]	0.78	0.77	0.9
NRIP2	nuclear receptor interacting protein 2 [Source:HGNC Symbol;Acc:23078]	0.49	0.5	0.62
IL6	interleukin 6 (interferon, beta 2) [Source:HGNC Symbol;Acc:6018]	0.59	0.6	0.72
ATRX	alpha thalassemia/mental retardation syndrome X-linked [Source:HGNC	0.35	0.33	0.47
CYB5D1	Symbol;Acc:886] cytochrome b5 domain containing 1 [Source:HGNC Symbol;Acc:26516]	0.35	0.33	0.47
BTBD18	BTB (POZ) domain containing 18 [Source:HGNC Symbol;Acc:37214]	0.54	0.52	0.18
DIDDI0	reticulocalbin 3, EF-hand calcium binding domain [Source:HGNC	0.54	0.52	0.00
RCN3	Symbol;Acc:21145]	0.74	0.76	0.88
MYT1L	myelin transcription factor 1-like [Source:HGNC Symbol;Acc:7623]	0.74	0.62	0.88
PACS2	phosphofurin acidic cluster sorting protein 2 [Source:HGNC Symbol;Acc:23794]	0.36	0.33	0.48
FYCO1	FYVE and coiled-coil domain containing 1 [Source:HGNC Symbol;Acc:14673]	0.09	0.06	0.21
	family with sequence similarity 155, member B [Source:HGNC	0.03	0.00	0.22
FAM155B	Symbol:Acc:307011	0.53	0.5	0.65
	ectonucleoside triphosphate diphosphohydrolase 2 [Source:HGNC			
ENTPD2	Symbol;Acc:3364]	0.55	0.59	0.71
ATPBD4	ATP binding domain 4 [Source:HGNC Symbol;Acc:30543]	0.06	0.02	0.18
	StAR-related lipid transfer (START) domain containing 8 [Source:HGNC			
STARD8	Symbol;Acc:19161]	0.6	0.55	0.72
MPZL2	myelin protein zero-like 2 [Source:HGNC Symbol;Acc:3496]	0.5	0.45	0.62
BCL6	B-cell CLL/lymphoma 6 [Source:HGNC Symbol;Acc:1001]	0.74	0.79	0.91
DFNA5	deafness, autosomal dominant 5 [Source:HGNC Symbol;Acc:2810]	0.27	0.33	0.45
RAET1E	retinoic acid early transcript 1E [Source:HGNC Symbol;Acc:16793]	0.88	0.82	1
SPAG17	sperm associated antigen 17 [Source:HGNC Symbol;Acc:26620]	0.61	0.54	0.73
C1orf116	chromosome 1 open reading frame 116 [Source:HGNC Symbol;Acc:28667]	0.77	0.7	0.89
PAK3	p21 protein (Cdc42/Rac)-activated kinase 3 [Source:HGNC Symbol;Acc:8592]	0.25	0.33	0.45
	leucine rich repeat transmembrane neuronal 1 [Source:HGNC			
LRRTM1	Symbol;Acc:19408]	0.2	0.29	0.41
FNBP1	formin binding protein 1 [Source:HGNC Symbol;Acc:17069]	0.23	0.32	0.44
	potassium voltage-gated channel, shaker-related subfamily, member 4	0 ==		
KCNA4	[Source:HGNC Symbol;Acc:6222]	0.78	0.65	0.9
	l de la la la compania de la compania del compania del compania de la compania del compania de la compania del compania de la compania de la compania de la compania de la compania del com	0.00		0
LPO	lactoperoxidase [Source:HGNC Symbol;Acc:6678]	0.83	0.69	0.95
LPO RAG1 ZNF211	lactoperoxidase [Source:HGNC Symbol;Acc:6678] recombination activating gene 1 [Source:HGNC Symbol;Acc:9831] zinc finger protein 211 [Source:HGNC Symbol;Acc:13003]	0.83 0.82 0.62	0.69 0.67 0.46	0.95 0.94 0.74

	CD79a molecule, immunoglobulin-associated alpha [Source:HGNC			
CD79A	Symbol;Acc:1698]	0.74	0.56	0.86
PXK	PX domain containing serine/threonine kinase [Source:HGNC Symbol;Acc:23326]	0.88	0.7	1
ARHGAP32	Rho GTPase activating protein 32 [Source:HGNC Symbol;Acc:17399]	0.84	0.65	0.96
PALM3	paralemmin 3 [Source:HGNC Symbol;Acc:33274]	0.48	0.29	0.6
REM2	RAS (RAD and GEM)-like GTP binding 2 [Source:HGNC Symbol;Acc:20248]	0.59	0.39	0.71
ANTXR1	anthrax toxin receptor 1 [Source:HGNC Symbol;Acc:21014]	0.54	0.34	0.66
VARS2	valyl-tRNA synthetase 2, mitochondrial [Source:HGNC Symbol;Acc:21642]	0.32	0.52	0.64
BCAS4	breast carcinoma amplified sequence 4 [Source:HGNC Symbol;Acc:14367]	0.18	0.61	0.73
CCL8 MRGPRF	chemokine (C-C motif) ligand 8 [Source:HGNC Symbol;Acc:10635] MAS-related GPR, member F [Source:HGNC Symbol;Acc:24828]	0.69	0.49	0.8 0.93
WINGPNF	transcription factor AP-2 epsilon (activating enhancer binding protein 2 epsilon)	0.62	0.55	0.95
TFAP2E	[Source:HGNC Symbol;Acc:30774]	0.34	0.35	0.46
NOSIP	nitric oxide synthase interacting protein [Source:HGNC Symbol;Acc:17946]	0.35	0.41	0.52
PRRG2	proline rich Gla (G-carboxyglutamic acid) 2 [Source:HGNC Symbol;Acc:9470]	0.35	0.41	0.52
DNPEP	aspartyl aminopeptidase [Source:HGNC Symbol;Acc:2981]	0.45	0.38	0.56
MAP6D1	MAP6 domain containing 1 [Source:HGNC Symbol;Acc:25753]	0.16	0.12	0.27
C. 153.44	split hand/foot malformation (ectrodactyly) type 1 [Source:HGNC	0.04	0.04	0.40
SHFM1 HEBP2	Symbol;Acc:10845]	0.01	0.01	0.12
IL20	heme binding protein 2 [Source:HGNC Symbol;Acc:15716] interleukin 20 [Source:HGNC Symbol;Acc:6002]	0.05	0.04	0.16 0.64
KRTAP4-7	keratin associated protein 4-7 [Source:HGNC Symbol;Acc:18898]	0.84	0.84	0.95
ACAT2	acetyl-CoA acetyltransferase 2 [Source:HGNC Symbol;Acc:94]	0.47	0.46	0.58
TC2N	tandem C2 domains, nuclear [Source:HGNC Symbol;Acc:19859]	0.37	0.36	0.48
ZNF502	zinc finger protein 502 [Source:HGNC Symbol;Acc:23718]	0.53	0.55	0.66
FCRL2	Fc receptor-like 2 [Source:HGNC Symbol;Acc:14875]	0.86	0.83	0.97
ZACN	zinc activated ligand-gated ion channel [Source:HGNC Symbol;Acc:29504]	0.79	0.76	0.9
HEXB OMD	hexosaminidase B (beta polypeptide) [Source:HGNC Symbol;Acc:4879]	0.53	0.49	0.64
MT1H	osteomodulin [Source:HGNC Symbol;Acc:8134] metallothionein 1H [Source:HGNC Symbol;Acc:7400]	0.67 0.14	0.63 0.19	0.78 0.3
MAGEE1	melanoma antigen family E, 1 [Source:HGNC Symbol;Acc:24934]	0.14	0.19	0.5
	family with sequence similarity 174, member B [Source:HGNC	0.5 .	0.55	0.5
FAM174B	Symbol;Acc:34339]	0.72	0.66	0.83
	solute carrier family 11 (proton-coupled divalent metal ion transporters),			
SLC11A1	member 1 [Source:HGNC Symbol;Acc:10907]	0.84	0.77	0.95
TRIOBP	TRIO and F-actin binding protein [Source:HGNC Symbol;Acc:17009]	0.18	0.11	0.29
TBC1D23	TBC1 domain family, member 23 [Source:HGNC Symbol;Acc:25622]	0.76	0.84	0.95
STX1B	syntaxin 1B [Source:HGNC Symbol;Acc:18539]	0.5	0.42	0.61
TP53INP1	tumor protein p53 inducible nuclear protein 1 [Source:HGNC Symbol;Acc:18022]	0.28	0.2	0.39
GGT6	gamma-glutamyltransferase 6 [Source:HGNC Symbol;Acc:26891]	0.6	0.68	0.79
PFKP	phosphofructokinase, platelet [Source:HGNC Symbol;Acc:8878]	0.38	0.28	0.49
TSLP	thymic stromal lymphopoietin [Source:HGNC Symbol;Acc:30743]	0.83	0.72	0.94
	transmembrane protein 158 (gene/pseudogene) [Source:HGNC			
TMEM158	Symbol;Acc:30293]	0.49	0.6	0.71
SNRK	SNF related kinase [Source:HGNC Symbol;Acc:30598]	0.38	0.5	0.61
C9orf9 LPGAT1	chromosome 9 open reading frame 9 [Source:HGNC Symbol;Acc:1367] lysophosphatidylglycerol acyltransferase 1 [Source:HGNC Symbol;Acc:28985]	0.48 0.13	0.6 0	0.71 0.24
C19orf26	chromosome 19 open reading frame 26 [Source:HGNC Symbol;Acc:28617]	0.13	0.42	0.67
CIJONEO	major facilitator superfamily domain containing 4 [Source:HGNC	0.50	0.42	0.07
MFSD4	Symbol;Acc:25433]	0.79	0.57	0.9
CDX4	caudal type homeobox 4 [Source:HGNC Symbol;Acc:1808]	0.43	0.66	0.77
PCBP4	poly(rC) binding protein 4 [Source:HGNC Symbol;Acc:8652]	0.31	0.06	0.42
NPB	neuropeptide B [Source:HGNC Symbol;Acc:30099]	0.22	0.49	0.6
	aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase)	0.40	0.55	0.55
AKR7A3	[Source:HGNC Symbol;Acc:390]	0.19	0.55	0.66
KRTAP19-8 NHSL1	keratin associated protein 19-8 [Source:HGNC Symbol;Acc:33898] NHS-like 1 [Source:HGNC Symbol;Acc:21021]	0.83 0.79	0.38 0.27	0.94
TNNI2	troponin I type 2 (skeletal, fast) [Source:HGNC Symbol;Acc:11946]	0.79	0.27	0.9 0.93
HRC	histidine rich calcium binding protein [Source:HGNC Symbol;Acc:5178]	0.8	0.83	0.93
NPY5R	neuropeptide Y receptor Y5 [Source:HGNC Symbol;Acc:7958]	0.46	0.58	0.68
PTGR1	prostaglandin reductase 1 [Source:HGNC Symbol;Acc:18429]	0.42	0.41	0.52
MOSPD1	motile sperm domain containing 1 [Source:HGNC Symbol;Acc:25235]	0.22	0.24	0.34
CRB2	crumbs homolog 2 (Drosophila) [Source:HGNC Symbol;Acc:18688]	0.45	0.14	0.55
ANGPT1	angiopoietin 1 [Source:HGNC Symbol;Acc:484]	0.07	0.05	0.17
PARP4	poly (ADP-ribose) polymerase family, member 4 [Source:HGNC Symbol;Acc:271]	0.21	0.18	0.31
CENANDE	sema domain, immunoglobulin domain (Ig), short basic domain, secreted,	0.00	0.13	0.22
SEMA3F GLUD2	(semaphorin) 3F [Source:HGNC Symbol;Acc:10728] glutamate dehydrogenase 2 [Source:HGNC Symbol;Acc:4336]	0.09 0.17	0.12 0.12	0.22 0.27
GLUDZ	family with sequence similarity 160, member B1 [Source:HGNC	0.17	0.12	0.27
FAM160B1	Symbol;Acc:29320]	0.09	0.03	0.19
MRC2	mannose receptor, C type 2 [Source:HGNC Symbol;Acc:16875]	0.12	0.04	0.22
DUOX1	dual oxidase 1 [Source:HGNC Symbol;Acc:3062]	0.07	0.15	0.25
DOOMI				

	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (I blood group)			
CNT2	[Source:HGNC Symbol;Acc:4204]	0.04	0.16	0.26
CL6B	B-cell CLL/lymphoma 6, member B [Source:HGNC Symbol;Acc:1002]	0.06	0.21	0.31
TD	cystic fibrosis transmembrane conductance regulator (ATP-binding cassette sub-	0.00	0.22	0.22
TR BLA00301	family C, member 7) [Source:HGNC Symbol;Acc:1884] RP11-471J12.1	0.06	0.23 0.05	0.33 0.18
ST4	bestrophin 4 [Source:HGNC Symbol;Acc:17106]	0.9	0.9	1
KRB1	bradykinin receptor B1 [Source:HGNC Symbol;Acc:1029]	0.87	0.87	0.97
	differentially expressed in FDCP 8 homolog (mouse) [Source:HGNC			
8	Symbol;Acc:25969]	0.37	0.37	0.47
S1	KiSS-1 metastasis-suppressor [Source:HGNC Symbol;Acc:6341]	0.84	0.83	0.94
	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 5 [Source:HGNC			
SNT5	Symbol;Acc:15684]	0.77	0.76	0.87
A1	chymase 1, mast cell [Source:HGNC Symbol;Acc:2097]	0.89	0.9	1
IHC23	zinc finger, DHHC-type containing 23 [Source:HGNC Symbol;Acc:28654]	0.25	0.26	0.36
H rf34	IQ motif containing H [Source:HGNC Symbol;Acc:25721]	0.84 0.64	0.82 0.62	0.94
.CN	chromosome 7 open reading frame 34 [Source:HGNC Symbol;Acc:21750] sodium leak channel, non-selective [Source:HGNC Symbol;Acc:19082]	0.62	0.64	0.74 0.74
orf27	chromosome 15 open reading frame 27 [Source:HGNC Symbol;Acc:26763]	0.24	0.26	0.36
-	lactase-like [Source:HGNC Symbol;Acc:15583]	0.63	0.65	0.75
MTSL3	ADAMTS-like 3 [Source:HGNC Symbol;Acc:14633]	0.26	0.28	0.38
	glypican 2 [Source:HGNC Symbol;Acc:4450]	0.12	0.14	0.24
2	growth arrest-specific 2 [Source:HGNC Symbol;Acc:4167]	0.66	0.63	0.76
	telomerase-associated protein 1 [Source:HGNC Symbol;Acc:11726]	0.34	0.31	0.44
15	interferon induced transmembrane protein 5 [Source:HGNC Symbol;Acc:16644]	0.8	0.84	0.94
X15	arachidonate 15-lipoxygenase [Source:HGNC Symbol;Acc:433]	0.32	0.28	0.42
	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 [Source:HGNC	0.6=	0.55	0
/L4 5	Symbol;Acc:3315]	0.67	0.63	0.77
5C1	Kruppel-like factor 5 (intestinal) [Source:HGNC Symbol;Acc:6349]	0.54	0.49	0.64 1
301	solute carrier family 35, member C1 [Source:HGNC Symbol;Acc:20197] Sp7 transcription factor [Source:HGNC Symbol;Acc:17321]	0.9 0.76	0.85 0.71	0.86
f65	chromosome 7 open reading frame 65 [Source:HGNC Symbol;Acc:34432]	0.78	0.71	0.88
103	StAR-related lipid transfer (START) domain containing 13 [Source:HGNC	0.70	0.72	0.00
RD13	Symbol;Acc:19164]	0.67	0.74	0.84
iD	crystallin, gamma D [Source:HGNC Symbol;Acc:2411]	0.86	0.77	0.96
	solute carrier family 12 (potassium/chloride transporters), member 8			
.2A8	[Source:HGNC Symbol;Acc:15595]	0.78	0.87	0.97
	solute carrier family 13 (sodium/sulfate symporters), member 4 [Source:HGNC			
3A4	Symbol;Acc:15827]	0.52	0.61	0.71
<i>1</i> 2	glutathione S-transferase mu 2 (muscle) [Source:HGNC Symbol;Acc:4634]	0.6	0.72	0.82
0R1	CD200 receptor 1 [Source:HGNC Symbol;Acc:24235]	0.9	0.77	1
	FXYD domain containing ion transport regulator 5 [Source:HGNC	0.74	0.50	0.04
5 15	Symbol; Acc: 4029]	0.74 0.09	0.59 0.27	0.84 0.37
13	cadherin 15, type 1, M-cadherin (myotubule) [Source:HGNC Symbol;Acc:1754] family with sequence similarity 71, member F2 [Source:HGNC	0.09	0.27	0.57
71F2	Symbol;Acc:27998]	0.9	0.69	1
P2	Kv channel interacting protein 2 [Source:HGNC Symbol;Acc:15522]	0.4	0.64	0.74
1R8	myotubularin related protein 8 [Source:HGNC Symbol;Acc:16825]	0.67	0.42	0.77
)1	pygopus homolog 1 (Drosophila) [Source:HGNC Symbol;Acc:30256]	0.83	0.84	0.93
)	dCMP deaminase [Source:HGNC Symbol;Acc:2710]	0.82	0.8	0.91
A4	S100 calcium binding protein A4 [Source:HGNC Symbol;Acc:10494]	0.59	0.72	0.81
01	RCSD domain containing 1 [Source:HGNC Symbol;Acc:28310]	0.42	0.58	0.67
	low density lipoprotein receptor class A domain containing 1 [Source:HGNC			
RAD1	Symbol;Acc:32069]	0.72	0.55	0.81
1	megalencephalic leukoencephalopathy with subcortical cysts 1 [Source:HGNC	0.74	0.54	0.0
1	Symbol;Acc:17082]	0.71	0.54	0.8
C8A HC15	kelch domain containing 8A [Source:HGNC Symbol;Acc:25573] zinc finger, DHHC-type containing 15 [Source:HGNC Symbol;Acc:20342]	0.71 0.46	0.25 0.46	0.8 0.55
1C15 Y1B3	guanylate cyclase 1, soluble, beta 3 [Source:HGNC Symbol;Acc:20342]	0.46	0.46	0.55
HV1	WDYHV motif containing 1 [Source:HGNC Symbol;Acc:25490]	0.33	0.37	0.46
R1	cysteine/histidine-rich 1 [Source:HGNC Symbol;Acc:17806]	0.41	0.38	0.5
554223	histocompatibility antigen-related	0.18	0.15	0.27
	solute carrier family 5 (iodide transporter), member 8 [Source:HGNC			
A8	Symbol;Acc:19119]	0.33	0.37	0.46
\ 8	EPH receptor A8 [Source:HGNC Symbol;Acc:3391]	0.19	0.14	0.28
	kinesin family member 21B [Source:HGNC Symbol;Acc:29442]	0.3	0.24	0.39
	dual specificity phosphatase 9 [Source:HGNC Symbol;Acc:3076]	0.46	0.21	0.55
P9				0.17
SP9 HR2	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231]	0.04	0.08	
SP9 HR2	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970]	0.04 0.08	0.08	0.17
SP9 HR2 D1	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970] echinoderm microtubule associated protein like 2 [Source:HGNC	0.08	0.01	0.17
SP9 HR2 D1	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970] echinoderm microtubule associated protein like 2 [Source:HGNC Symbol;Acc:18035]			
5P9 HR2 D1	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970] echinoderm microtubule associated protein like 2 [Source:HGNC Symbol;Acc:18035] interferon regulatory factor 2 binding protein 1 [Source:HGNC	0.08	0.01	0.17
SP9 HR2 D1 -2	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970] echinoderm microtubule associated protein like 2 [Source:HGNC Symbol;Acc:18035] interferon regulatory factor 2 binding protein 1 [Source:HGNC Symbol;Acc:21728]	0.08 0.02	0.01 0.02	0.17 0.11 0.09
21B SP9 HR2 O1 IL2 2BP1 K	cadherin-related family member 2 [Source:HGNC Symbol;Acc:18231] intermediate filament family orphan 1 [Source:HGNC Symbol;Acc:24970] echinoderm microtubule associated protein like 2 [Source:HGNC Symbol;Acc:18035] interferon regulatory factor 2 binding protein 1 [Source:HGNC	0.08	0.01	0.17

OTX2	orthodenticle homeobox 2 [Source:HGNC Symbol;Acc:8522]	0.01	0.02	0.11
COL4A1	collagen, type IV, alpha 1 [Source:HGNC Symbol;Acc:2202]	0.09	0.07	0.18
COL4A2	collagen, type IV, alpha 2 [Source:HGNC Symbol;Acc:2203]	0.09	0.07	0.18
ZNF672	zinc finger protein 672 [Source:HGNC Symbol;Acc:26179]	0.09	0.07	0.18
	, , , ,			
PRSS35	protease, serine, 35 [Source:HGNC Symbol;Acc:21387]	0	0.02	0.11
RFTN2	raftlin family member 2 [Source:HGNC Symbol;Acc:26402]	0.21	0.23	0.32
SH3GLB1	SH3-domain GRB2-like endophilin B1 [Source:HGNC Symbol;Acc:10833]	0.12	0.15	0.24
CYR61	cysteine-rich, angiogenic inducer, 61 [Source:HGNC Symbol;Acc:2654]	0.03	0.07	0.16
	protein phosphatase 2, regulatory subunit B", alpha [Source:HGNC			
DDDDDDA		0.15	0.1	0.24
PPP2R3A	Symbol;Acc:9307]	0.15	0.1	
DUOXA1	dual oxidase maturation factor 1 [Source:HGNC Symbol;Acc:26507]	0.13	0.23	0.32
CHIT1	chitinase 1 (chitotriosidase) [Source:HGNC Symbol;Acc:1936]	0.87	0.87	0.96
C1orf100	chromosome 1 open reading frame 100 [Source:HGNC Symbol;Acc:30435]	0.78	0.78	0.87
TMEM132A	transmembrane protein 132A [Source:HGNC Symbol;Acc:31092]	0.79	0.79	0.88
LAMP2	lysosomal-associated membrane protein 2 [Source:HGNC Symbol;Acc:6501]	0.38	0.38	0.47
FAM123B	APC membrane recruitment protein 1 [Source:HGNC Symbol;Acc:26837]	0.33	0.33	0.42
DPT	dermatopontin [Source:HGNC Symbol;Acc:3011]	0.91	0.9	1
	DnaJ (Hsp40) homolog, subfamily B, member 8 [Source:HGNC			
DNAJB8	Symbol;Acc:23699]	0.75	0.74	0.84
LYL1	lymphoblastic leukemia derived sequence 1 [Source:HGNC Symbol;Acc:6734]	0.67	0.68	0.77
CCDC152		0.53	0.54	0.63
CCDC132	coiled-coil domain containing 152 [Source:HGNC Symbol;Acc:34438]	0.55	0.34	0.03
	mal, T-cell differentiation protein 2 (gene/pseudogene) [Source:HGNC			
MAL2	Symbol;Acc:13634]	0.66	0.67	0.76
SYT14	synaptotagmin XIV [Source:HGNC Symbol;Acc:23143]	0.39	0.37	0.48
KRT73	keratin 73 [Source:HGNC Symbol;Acc:28928]	0.85	0.83	0.94
	cytochrome P450, family 1, subfamily A, polypeptide 2 [Source:HGNC			
CVD1 A 2		0.87	0.05	0.96
CYP1A2	Symbol;Acc:2596]		0.85	
NRBF2	nuclear receptor binding factor 2 [Source:HGNC Symbol;Acc:19692]	0.62	0.64	0.73
MMP10	matrix metallopeptidase 10 (stromelysin 2) [Source:HGNC Symbol;Acc:7156]	0.86	0.83	0.95
RHBDL2	rhomboid, veinlet-like 2 (Drosophila) [Source:HGNC Symbol;Acc:16083]	0.75	0.72	0.84
PDZK1IP1	PDZK1 interacting protein 1 [Source:HGNC Symbol;Acc:16887]	0.81	0.78	0.9
I DZKIII I	1 DERT Interdeding protein 1 [Jource. Tone Symbol, rec. 10007]	0.01	0.70	0.5
TT1140		0.07		0.05
TTLL10	tubulin tyrosine ligase-like family, member 10 [Source:HGNC Symbol;Acc:26693]	0.27	0.24	0.36
SEMG1	semenogelin I [Source:HGNC Symbol;Acc:10742]	0.86	0.83	0.95
CTLA4	cytotoxic T-lymphocyte-associated protein 4 [Source:HGNC Symbol;Acc:2505]	0.88	0.91	1
RNF133	ring finger protein 133 [Source:HGNC Symbol;Acc:21154]	0.87	0.9	0.99
	lymphotoxin beta (TNF superfamily, member 3) [Source:HGNC			
LTD		0.05	0.01	0.04
LTB	Symbol;Acc:6711]	0.85	0.81	0.94
PXDNL	peroxidasin homolog (Drosophila)-like [Source:HGNC Symbol;Acc:26359]	0.56	0.6	0.69
ACOT9	acyl-CoA thioesterase 9 [Source:HGNC Symbol;Acc:17152]	0.49	0.45	0.58
CSK	c-src tyrosine kinase [Source:HGNC Symbol;Acc:2444]	0.22	0.26	0.35
	APEX nuclease (apurinic/apyrimidinic endonuclease) 2 [Source:HGNC	*		0.00
ADEVA		0.39	0.25	0.48
APEX2	Symbol;Acc:17889]		0.35	
GEMIN4	gem (nuclear organelle) associated protein 4 [Source:HGNC Symbol;Acc:15717]	0.69	0.73	0.82
	peptidylprolyl cis/trans isomerase, NIMA-interacting 1 [Source:HGNC			
PIN1	Symbol;Acc:8988]	0.82	0.86	0.95
GTF2A1L	general transcription factor IIA, 1-like [Source:HGNC Symbol;Acc:30727]	0.7	0.74	0.83
MT1B	metallothionein 1B [Source:HGNC Symbol;Acc:7394]	0.86	0.81	0.95
PLS3	plastin 3 [Source:HGNC Symbol;Acc:9091]	0.52	0.47	0.61
TNS4	tensin 4 [Source:HGNC Symbol;Acc:24352]	0.7	0.75	0.84
IL13RA1	interleukin 13 receptor, alpha 1 [Source:HGNC Symbol;Acc:5974]	0.62	0.56	0.71
TMSB15B	thymosin beta 15B [Source:HGNC Symbol;Acc:28612]	0.49	0.43	0.58
CXCL6	chemokine (C-X-C motif) ligand 6 [Source:HGNC Symbol;Acc:10643]	0.42	0.48	0.57
SSTR4	somatostatin receptor 4 [Source:HGNC Symbol;Acc:11333]			0.9
		0.81	0.75	
DUSP28	dual specificity phosphatase 28 [Source:HGNC Symbol;Acc:33237]	0.7	0.76	0.85
OXT	oxytocin/neurophysin I prepropeptide [Source:HGNC Symbol;Acc:8528]	0.85	0.91	1
PLEK	pleckstrin [Source:HGNC Symbol;Acc:9070]	0.88	0.8	0.97
GSTK1	glutathione S-transferase kappa 1 [Source:HGNC Symbol;Acc:16906]	0.62	0.53	0.71
RASAL1	RAS protein activator like 1 (GAP1 like) [Source:HGNC Symbol;Acc:9873]	0.11	0.2	0.29
NAJALI		0.11	0.2	0.29
511000	ectonucleotide pyrophosphatase/phosphodiesterase 3 [Source:HGNC	0.01	0	0.5
ENPP3	Symbol;Acc:3358]	0.81	0.72	0.9
THY1	Thy-1 cell surface antigen [Source:HGNC Symbol;Acc:11801]	0.55	0.65	0.74
DNM1P35	DNM1 pseudogene 35 [Source:HGNC Symbol;Acc:35182]	0.91	0.8	1
DEFB132	defensin, beta 132 [Source:HGNC Symbol;Acc:33806]	0.62	0.51	0.71
SLC44A2	solute carrier family 44, member 2 [Source:HGNC Symbol;Acc:17292]	0.38	0.23	0.47
JLC44AZ		0.30	0.23	0.47
	solute carrier family 19 (folate transporter), member 1 [Source:HGNC			
SLC19A1	Symbol;Acc:10937]	0.52	0.32	0.61
AZI1	5-azacytidine induced 1 [Source:HGNC Symbol;Acc:29511]	0.77	0.57	0.86
	ATP-binding cassette, sub-family A (ABC1), member 6 [Source:HGNC			
ABCA6	Symbol;Acc:36]	0.71	0.91	1
	,			
IL8	interleukin 8 [Source:HGNC Symbol;Acc:6025]	0.91	0.7	1
CNTN5	contactin 5 [Source:HGNC Symbol;Acc:2175]	0.32	0.09	0.41
ZFP42	ZFP42 zinc finger protein [Source:HGNC Symbol;Acc:30949]	0.41	0.76	0.85
IL17B	interleukin 17B [Source:HGNC Symbol;Acc:5982]	0.85	0.44	0.94
DSG3	desmoglein 3 [Source:HGNC Symbol;Acc:3982]		0.44	
	• • • • • • •	0.85		0.93
TMEM63C	transmembrane protein 63C [Source:HGNC Symbol;Acc:23787]	0.47	0.46	0.55

CD34	CD24 malacula [Saurca:HCNC Sumbal-Acc:1662]	0.46	0.49	0.56
CD34 IKZF3	CD34 molecule [Source:HGNC Symbol;Acc:1662] IKAROS family zinc finger 3 (Aiolos) [Source:HGNC Symbol;Acc:13178]	0.46 0.6	0.48 0.58	0.56 0.68
C19orf35	chromosome 19 open reading frame 35 [Source:HGNC Symbol;Acc:24793]	0.85	0.58	0.08
BOLL	bol, boule-like (Drosophila) [Source:HGNC Symbol;Acc:14273]	0.69	0.71	0.79
ARRB2	arrestin, beta 2 [Source:HGNC Symbol;Acc:712]	0.56	0.59	0.67
TAS2R60	taste receptor, type 2, member 60 [Source:HGNC Symbol;Acc:20639]	0.79	0.82	0.9
TBX10	T-box 10 [Source:HGNC Symbol;Acc:11593]	0.58	0.54	0.66
LMO3	LIM domain only 3 (rhombotin-like 2) [Source:HGNC Symbol;Acc:6643]	0.53	0.57	0.65
	phytanoyl-CoA 2-hydroxylase interacting protein [Source:HGNC			
PHYHIP	Symbol;Acc:16865]	0.55	0.6	0.68
SYCP1	synaptonemal complex protein 1 [Source:HGNC Symbol;Acc:11487]	0.52	0.58	0.66
C22orf32	chromosome 22 open reading frame 32 [Source:HGNC Symbol;Acc:25055]	0.6	0.53	0.68
ZBTB4	zinc finger and BTB domain containing 4 [Source:HGNC Symbol;Acc:23847]	0.82	0.73	0.9
GLT25D2	glycosyltransferase 25 domain containing 2 [Source:HGNC Symbol;Acc:16790] solute carrier family 34 (sodium phosphate), member 3 [Source:HGNC	0.73	0.61	0.81
SLC34A3	Symbol:Acc:203051	0.82	0.7	0.9
TIAM2	T-cell lymphoma invasion and metastasis 2 [Source:HGNC Symbol;Acc:11806]	0.6	0.47	0.68
RAB17	RAB17, member RAS oncogene family [Source:HGNC Symbol; Acc:16523]	0.45	0.59	0.67
CD1D	CD1d molecule [Source:HGNC Symbol;Acc:1637]	0.53	0.7	0.78
VSIG8	chromosome 1 open reading frame 204 [Source:HGNC Symbol;Acc:27647]	0.82	0.55	0.9
	CMT duplicated region transcript 1; Uncharacterized protein			
CDRT1	[Source:UniProtKB/TrEMBL;Acc:Q9BXD7]	0.58	0.24	0.66
ZC3HAV1L	zinc finger CCCH-type, antiviral 1-like [Source:HGNC Symbol;Acc:22423]	0.37	0.73	0.81
TLX1	T-cell leukemia homeobox 1 [Source:HGNC Symbol;Acc:5056]	0.32	0.32	0.4
TLX1NB	TLX1 neighbor [Source:HGNC Symbol;Acc:37183]	0.32	0.32	0.4
FRMD4A	FERM domain containing 4A [Source:HGNC Symbol;Acc:25491]	0.41	0.41	0.49
NEKDIZ	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, zeta	0.17	0.10	0.20
NFKBIZ FNIP2	[Source:HGNC Symbol;Acc:29805] folliculin interacting protein 2 [Source:HGNC Symbol;Acc:29280]	0.17 0.31	0.18 0.32	0.26 0.4
TMEM164	transmembrane protein 164 [Source:HGNC Symbol;Acc:29280]	0.51	0.32	0.4
RARA	retinoic acid receptor, alpha [Source:HGNC Symbol;Acc:9864]	0.03	0.06	0.14
ABHD14A	abhydrolase domain containing 14A [Source:HGNC Symbol;Acc:24538]	0.09	0.06	0.17
	Usher syndrome 1C (autosomal recessive, severe) [Source:HGNC			
USH1C	Symbol;Acc:12597]	0.17	0.2	0.28
LPAR1	lysophosphatidic acid receptor 1 [Source:HGNC Symbol;Acc:3166]	0.15	0.18	0.26
RESP18	regulated endocrine-specific protein 18 [Source:HGNC Symbol;Acc:33762]	0.12	0.08	0.2
A4GALT	alpha 1,4-galactosyltransferase [Source:HGNC Symbol;Acc:18149]	0.02	0.06	0.14
ASS1	argininosuccinate synthase 1 [Source:HGNC Symbol;Acc:758]	0.41	0.46	0.54
NDUEVO	NADH dehydrogenase (ubiquinone) flavoprotein 3, 10kDa [Source:HGNC	0.42	0.03	0.2
NDUFV3 LDHC	Symbol;Acc:7719]	0.12 0.19	0.03 0.29	0.2 0.37
PDE4B	lactate dehydrogenase C [Source:HGNC Symbol;Acc:6544] phosphodiesterase 4B, cAMP-specific [Source:HGNC Symbol;Acc:8781]	0.19	0.29	0.37
CLEC14A	C-type lectin domain family 14, member A [Source:HGNC Symbol;Acc:19832]	0.12	0.25	0.33
CELF3	CUGBP, Elav-like family member 3 [Source:HGNC Symbol;Acc:11967]	0.32	0.11	0.4
LIN54	lin-54 homolog (C. elegans) [Source:HGNC Symbol;Acc:25397]	0.02	0	0.1
FBXL14	F-box and leucine-rich repeat protein 14 [Source:HGNC Symbol;Acc:28624]	0.11	0.07	0.19
MPDU1	mannose-P-dolichol utilization defect 1 [Source:HGNC Symbol;Acc:7207]	0.2	0.22	0.3
EFHD2	EF-hand domain family, member D2 [Source:HGNC Symbol;Acc:28670]	0.13	0.11	0.21
TMEM179	transmembrane protein 179 [Source:HGNC Symbol;Acc:20137]	0.23	0.16	0.31
NTAN1	N-terminal asparagine amidase [Source:HGNC Symbol;Acc:29909]	0.17	0.07	0.25
NME3	NME/NM23 nucleoside diphosphate kinase 3 [Source:HGNC Symbol;Acc:7851]	0.06	0.16	0.24
HEPH NMUR1	hephaestin [Source:HGNC Symbol;Acc:4866] neuromedin U receptor 1 [Source:HGNC Symbol;Acc:4518]	0.33 0.56	0.33 0.56	0.41 0.64
ITLN2	intelectin 2 [Source:HGNC Symbol;Acc:20599]	0.91	0.9	0.99
C20orf152	N/A	0.79	0.78	0.87
CZOOTIISZ	amine oxidase, copper containing 3 (vascular adhesion protein 1) [Source:HGNC	0.75	0.70	0.07
AOC3	Symbol;Acc:550]	0.85	0.86	0.94
GFRA4	GDNF family receptor alpha 4 [Source:HGNC Symbol;Acc:13821]	0.61	0.62	0.7
	UDP glucuronosyltransferase 2 family, polypeptide A3 [Source:HGNC			
UGT2A3	Symbol;Acc:28528]	0.85	0.86	0.94
	protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting			
PPFIA4	protein (liprin), alpha 4 [Source:HGNC Symbol;Acc:9248]	0.9	0.88	0.98
LMO1	LIM domain only 1 (rhombotin 1) [Source:HGNC Symbol;Acc:6641]	0.63	0.61	0.71
LRRC33	leucine rich repeat containing 33 [Source:HGNC Symbol;Acc:24613]	0.66	0.64	0.74
DITOD	RGD motif, leucine rich repeats, tropomodulin domain and proline-rich	0.0	0.02	1
RLTPR	containing [Source:HGNC Symbol;Acc:27089]	0.9	0.92	1
IRF8 NEB	interferon regulatory factor 8 [Source:HGNC Symbol;Acc:5358] nebulin [Source:HGNC Symbol;Acc:7720]	0.59 0.86	0.61 0.88	0.69 0.96
MYL7	myosin, light chain 7, regulatory [Source:HGNC Symbol;Acc:21719]	0.86	0.88	0.98
COL6A3	collagen, type VI, alpha 3 [Source:HGNC Symbol;Acc:2213]	0.87	0.86	0.98
RGAG4	retrotransposon gag domain containing 4 [Source:HGNC Symbol;Acc:29430]	0.56	0.53	0.64
RORC	RAR-related orphan receptor C [Source:HGNC Symbol;Acc:10260]	0.52	0.55	0.63
	transient receptor potential cation channel, subfamily V, member 2			
TRPV2	[Source:HGNC Symbol;Acc:18082]	0.62	0.65	0.73
	sodium channel, voltage-gated, type V, alpha subunit [Source:HGNC			
SCN5A	Symbol;Acc:10593]	0.61	0.65	0.73

C1R	complement component 1, r subcomponent [Source:HGNC Symbol;Acc:1246]	0.78	0.74	0.86
AAGAB	alpha- and gamma-adaptin binding protein [Source:HGNC Symbol;Acc:25662]	0.88	0.92	1
70.07.0	O-linked N-acetylglucosamine (GlcNAc) transferase [Source:HGNC	0.00	0.32	-
OGT	Symbol;Acc:8127]	0.51	0.46	0.59
TSPO	translocator protein (18kDa) [Source:HGNC Symbol;Acc:1158]	0.34	0.39	0.47
DOK3	docking protein 3 [Source:HGNC Symbol;Acc:24583]	0.89	0.84	0.97
RSPO4	R-spondin 4 [Source:HGNC Symbol;Acc:16175]	0.63	0.68	0.76
	small nuclear RNA activating complex, polypeptide 1, 43kDa [Source:HGNC			
SNAPC1	Symbol;Acc:11134]	0.51	0.45	0.59
CSTA	cystatin A (stefin A) [Source:HGNC Symbol;Acc:2481]	0.86	0.92	1
65171	tumor necrosis factor (ligand) superfamily, member 8 [Source:HGNC	0.00	0.52	-
TNFSF8	Symbol;Acc:11938]	0.74	0.8	0.88
GAL	galanin/GMAP prepropeptide [Source:HGNC Symbol;Acc:4114]	0.43	0.5	0.58
ALX3	ALX homeobox 3 [Source:HGNC Symbol;Acc:449]	0.79	0.71	0.87
	cholinergic receptor, nicotinic, alpha 2 (neuronal) [Source:HGNC			
CHRNA2	Symbol;Acc:1956]	0.66	0.58	0.74
SNX32	sorting nexin 32 [Source:HGNC Symbol;Acc:26423]	0.47	0.55	0.63
ADCY9	adenylate cyclase 9 [Source:HGNC Symbol;Acc:240]	0.57	0.65	0.73
CSMD1	CUB and Sushi multiple domains 1 [Source:HGNC Symbol;Acc:14026]	0.77	0.68	0.85
PHOSPHO1	phosphatase, orphan 1 [Source:HGNC Symbol;Acc:16815]	0.52	0.43	0.6
SLC47A1	solute carrier family 47, member 1 [Source:HGNC Symbol;Acc:25588]	0.28	0.19	0.36
LOC150568	uncharacterized	0.3	0.4	0.48
C3orf72	chromosome 3 open reading frame 72 [Source:HGNC Symbol;Acc:34428]	0.3	0.4	0.48
FOXL2	forkhead box L2 [Source:HGNC Symbol;Acc:1092]	0.3	0.4	0.48
C7	complement component 7 [Source:HGNC Symbol;Acc:1346]	0.8	0.92	1
UPB1	ureidopropionase, beta [Source:HGNC Symbol;Acc:16297]	0.4	0.52	0.6
31 DI	ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative) [Source:HGNC	J.4	0.52	0.0
ENPP4	Symbol;Acc:3359]	0.27	0.14	0.35
ZMYM3	zinc finger, MYM-type 3 [Source:HGNC Symbol;Acc:13054]	0.56	0.42	0.64
KAT2A	K(lysine) acetyltransferase 2A [Source:HGNC Symbol;Acc:4201]	0.88	0.71	0.04
AQPEP	Aminopeptidase Q [Source:UniProtKB/Swiss-Prot;Acc:Q6Q4G3]	0.44	0.71	0.30
ASPA	aspartoacylase [Source:HGNC Symbol;Acc:756]	0.44	0.68	0.7
DNAH1	dynein, axonemal, heavy chain 1 [Source:HGNC Symbol;Acc:2940]	0.87	0.58	0.95
DIVALIT	, , , , , , , , , , , , , , , , , , , ,	0.77	0.56	0.65
CNIDIII	gonadotropin-releasing hormone 1 (luteinizing-releasing hormone)	0.7	0.92	1
GNRH1	[Source:HGNC Symbol;Acc:4419]			
SCG2	secretogranin II [Source:HGNC Symbol;Acc:10575]	0.5	0.14	0.58
GRIK5	glutamate receptor, ionotropic, kainate 5 [Source:HGNC Symbol;Acc:4583]	0.68	0.29	0.76
TDANIKA	tetratricopeptide repeat and ankyrin repeat containing 1 [Source:HGNC	0.44	0.54	0.50
TRANK1	Symbol;Acc:29011]	0.11	0.51	0.59
RGMA	RGM domain family, member A [Source:HGNC Symbol;Acc:30308]	0.9	0.33	0.98
ZBTB44	zinc finger and BTB domain containing 44 [Source:HGNC Symbol;Acc:25001]	0.83	0.83	0.9
	ring finger protein 40, E3 ubiquitin protein ligase [Source:HGNC			
RNF40	Symbol;Acc:16867]	0.86	0.86	0.93
CLEC5A	C-type lectin domain family 5, member A [Source:HGNC Symbol;Acc:2054]	0.82	0.82	0.89
GYS2	glycogen synthase 2 (liver) [Source:HGNC Symbol;Acc:4707]	0.7	0.69	0.77
HHLA2	HERV-H LTR-associating 2 [Source:HGNC Symbol;Acc:4905]	0.85	0.84	0.92
CCL28	chemokine (C-C motif) ligand 28 [Source:HGNC Symbol;Acc:17700]	0.84	0.83	0.91
C6orf10	chromosome 6 open reading frame 10 [Source:HGNC Symbol;Acc:13922]	0.84	0.83	0.91
	GIPC PDZ domain containing family, member 2 [Source:HGNC			
GIPC2	Symbol;Acc:18177]	0.84	0.85	0.92
C11orf68	chromosome 11 open reading frame 68 [Source:HGNC Symbol;Acc:28801]	0.85	0.86	0.93
SFTPB	surfactant protein B [Source:HGNC Symbol;Acc:10801]	0.85	0.86	0.93
FAM165B	small integral membrane protein 11 [Source:HGNC Symbol;Acc:1293]	0.82	0.83	0.9
ZNF773	zinc finger protein 773 [Source:HGNC Symbol;Acc:30487]	0.81	0.83	0.9
CLEC9A	C-type lectin domain family 9, member A [Source:HGNC Symbol;Acc:26705]	0.83	0.85	0.92
	amiloride binding protein 1 (amine oxidase (copper-containing)) [Source:HGNC			
ABP1	Symbol;Acc:80]	0.83	0.8	0.9
COL20A1	collagen, type XX, alpha 1 [Source:HGNC Symbol;Acc:14670]	0.8	0.83	0.9
FAM120C	family with sequence similarity 120C [Source:HGNC Symbol;Acc:16949]	0.48	0.45	0.55
	benzodiazapine receptor (peripheral) associated protein 1 [Source:HGNC			
BZRAP1	Symbol;Acc:16831]	0.72	0.69	0.79
MAP2K4	mitogen-activated protein kinase kinase 4 [Source:HGNC Symbol;Acc:6844]	0.58	0.61	0.68
C10orf108	proline rich 26 [Source:HGNC Symbol;Acc:30724]	0.81	0.86	0.93
	myeloma overexpressed (in a subset of t(11;14) positive multiple myelomas)			
MYEOV	[Source:HGNC Symbol;Acc:7563]	0.77	0.82	0.89
CD300E	CD300e molecule [Source:HGNC Symbol;Acc:28874]	0.83	0.76	0.9
	guanine nucleotide binding protein (G protein), alpha 14 [Source:HGNC			
GNA14	Symbol;Acc:4382]	0.66	0.73	0.8
	protein phosphatase, Mg2+/Mn2+ dependent, 1M [Source:HGNC			
PPM1M	Symbol;Acc:26506]	0.75	0.83	0.9
MTA1	metastasis associated 1 [Source:HGNC Symbol;Acc:7410]	0.83	0.71	0.9
	vesicle-associated membrane protein 1 (synaptobrevin 1) [Source:HGNC			
VAMP1	Symbol;Acc:12642]	0.74	0.61	0.81
KRTAP10-4	keratin associated protein 10-4 [Source:HGNC Symbol;Acc:20521]	0.83	0.69	0.9
	calcium/calmodulin-dependent protein kinase I [Source:HGNC	2.25	2.00	2.5
CAMK1	Symbol;Acc:1459]	0.42	0.58	0.65
<u>-</u>	- j	J. /2	0.50	3.03

SVOP	SV2 related protein homolog (rat) [Source:HGNC Symbol;Acc:25417]	0.85	0.67	0.92
CD36	CD36 molecule (thrombospondin receptor) [Source:HGNC Symbol;Acc:1663]	0.86	0.66	0.93
CTHRC1	collagen triple helix repeat containing 1 [Source:HGNC Symbol;Acc:18831]	0.85	0.6	0.92
ATP9A	ATPase, class II, type 9A [Source:HGNC Symbol;Acc:13540]	0.72	0.45	0.79
HOXC13	homeobox C13 [Source:HGNC Symbol;Acc:5125]	0.13	0.13	0.2
	actin related protein 2/3 complex, subunit 2, 34kDa [Source:HGNC			
ARPC2	Symbol;Acc:705]	0	0	0.07
NFE2L2	nuclear factor (erythroid-derived 2)-like 2 [Source:HGNC Symbol;Acc:7782]	0	0	0.07
CLVS1	clavesin 1 [Source:HGNC Symbol;Acc:23139] C1q and tumor necrosis factor related protein 2 [Source:HGNC	0.01	0.01	0.08
C1QTNF2	Symbol;Acc:14325]	0.28	0.29	0.36
C3orf70	chromosome 3 open reading frame 70 [Source:HGNC Symbol;Acc:33731]	0.01	0	0.08
MSH5	mutS homolog 5 (E. coli) [Source:HGNC Symbol;Acc:7328]	0.04	0.03	0.11
IZUMO1	izumo sperm-egg fusion 1 [Source:HGNC Symbol;Acc:28539]	0.03	0.04	0.11
	ST3 beta-galactoside alpha-2,3-sialyltransferase 3 [Source:HGNC			
ST3GAL3	Symbol;Acc:10866]	0.16	0.15	0.23
FJX1	four jointed box 1 (Drosophila) [Source:HGNC Symbol;Acc:17166]	0.27	0.26	0.34
C3orf37	chromosome 3 open reading frame 37 [Source:HGNC Symbol;Acc:24446]	0.39	0.38	0.46
ANGPTL2	angiopoietin-like 2 [Source:HGNC Symbol;Acc:490]	0.06	0.07	0.14
IL23A	interleukin 23, alpha subunit p19 [Source:HGNC Symbol;Acc:15488]	0.41	0.39	0.48
TUBGCP5	tubulin, gamma complex associated protein 5 [Source:HGNC Symbol;Acc:18600]	0.06	0.04	0.13
SLC26A5	solute carrier family 26, member 5 (prestin) [Source:HGNC Symbol;Acc:9359]	0.15	0.13	0.13
GPR19	G protein-coupled receptor 19 [Source:HGNC Symbol;Acc:4473]	0.27	0.3	0.37
SYTL4	synaptotagmin-like 4 [Source:HGNC Symbol;Acc:15588]	0.4	0.43	0.5
	aldehyde dehydrogenase 1 family, member A2 [Source:HGNC			
ALDH1A2	Symbol;Acc:15472]	0.13	0.1	0.2
	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1			
NFKB1	[Source:HGNC Symbol;Acc:7794]	0.06	0.03	0.13
METTL7B	methyltransferase like 7B [Source:HGNC Symbol;Acc:28276]	0.16	0.19	0.26
CXCL14	chemokine (C-X-C motif) ligand 14 [Source:HGNC Symbol;Acc:10640]	0.04	0.07	0.14
05145	growth factor independent 1B transcription repressor [Source:HGNC	0.00	0.25	0.00
GFI1B	Symbol;Acc:4238] pancreatic progenitor cell differentiation and proliferation factor homolog	0.22	0.25	0.32
PPDPF	(zebrafish) [Source:HGNC Symbol;Acc:16142]	0.32	0.36	0.43
11011	alcohol dehydrogenase 1B (class I), beta polypeptide [Source:HGNC	0.32	0.50	0.43
ADH1B	Symbol;Acc:250]	0.27	0.31	0.38
KLHL14	kelch-like family member 14 [Source:HGNC Symbol;Acc:29266]	0	0.04	0.11
	small nucleolar RNA host gene 12 (non-protein coding) [Source:HGNC			
SNHG12	Symbol;Acc:30062]	0.07	0.02	0.14
FLYWCH2	FLYWCH family member 2 [Source:HGNC Symbol;Acc:25178]	0.16	0.1	0.23
	ATP-binding cassette, sub-family A (ABC1), member 7 [Source:HGNC			
ABCA7	Symbol;Acc:37]	0.18	0.1	0.25
TSPAN17	tetraspanin 17 [Source:HGNC Symbol;Acc:13594]	0.39	0.31	0.46
C15orf38 POMT1	chromosome 15 open reading frame 38 [Source:HGNC Symbol;Acc:28782]	0.38 0.12	0.46 0.03	0.53 0.19
ZFPM2	protein-O-mannosyltransferase 1 [Source:HGNC Symbol;Acc:9202] zinc finger protein, FOG family member 2 [Source:HGNC Symbol;Acc:16700]	0.12	0.03	0.19
ZIFIVIZ	phosphotyrosine interaction domain containing 1 [Source:HGNC	0.03	0.19	0.20
PID1	Symbol;Acc:26084]	0.16	0.26	0.33
ANKLE1	ankyrin repeat and LEM domain containing 1 [Source:HGNC Symbol;Acc:26812]	0.24	0.35	0.42
TASP1	taspase, threonine aspartase, 1 [Source:HGNC Symbol;Acc:15859]	0.06	0.18	0.25
	SMT3 suppressor of mif two 3 homolog 3 (S. cerevisiae) [Source:HGNC			
SUMO3	Symbol;Acc:11124]	0.13	0.3	0.37
HK1	hexokinase 1 [Source:HGNC Symbol;Acc:4922]	0.25	0.43	0.5
CBX2	chromobox homolog 2 [Source:HGNC Symbol;Acc:1552]	0.02	0.01	0.09
TMEM17	transmembrane protein 17 [Source:HGNC Symbol;Acc:26623]	0.03	0.05	0.12
TCF3	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	0.00	0.11	0.10
TCF3 RTN3	[Source:HGNC Symbol;Acc:11633] reticulon 3 [Source:HGNC Symbol;Acc:10469]	0.08 0.11	0.11 0.07	0.18 0.18
DPP6	dipeptidyl-peptidase 6 [Source:HGNC Symbol;Acc:3010]	0.11	0.07	0.18
TMEM154	transmembrane protein 154 [Source:HGNC Symbol;Acc:26489]	0.03	0.05	0.12
RRM2	ribonucleotide reductase M2 [Source:HGNC Symbol;Acc:10452]	0.11	0.03	0.18
CD80	CD80 molecule [Source:HGNC Symbol;Acc:1700]	0.13	0.14	0.21
DTX3L	deltex 3-like (Drosophila) [Source:HGNC Symbol;Acc:30323]	0.11	0.22	0.29
	poly (ADP-ribose) polymerase family, member 9 [Source:HGNC			
PARP9	Symbol;Acc:24118]	0.11	0.22	0.29
LAYN	layilin [Source:HGNC Symbol;Acc:29471]	0.87	0.87	0.94
LAX1	lymphocyte transmembrane adaptor 1 [Source:HGNC Symbol;Acc:26005]	0.89	0.89	0.96
HTN3	histatin 3 [Source:HGNC Symbol;Acc:5284]	0.93	0.93	1
C9orf66	chromosome 9 open reading frame 66 [Source:HGNC Symbol;Acc:26436]	0.77	0.77	0.84
GPR84	G protein-coupled receptor 84 [Source:HGNC Symbol:Acc:4535]	0.87	0.86	0.94
RBP5 EVPLL	retinol binding protein 5, cellular [Source:HGNC Symbol;Acc:15847] envoplakin-like [Source:HGNC Symbol;Acc:35236]	0.52 0.87	0.51 0.86	0.59 0.94
SNORD17	small nucleolar RNA, C/D box 17 [Source:HGNC Symbol;Acc:32713]	0.87	0.86	0.94
PCDHB13	protocadherin beta 13 [Source:HGNC Symbol;Acc:8684]	0.88	0.87	0.95
COL15A1	collagen, type XV, alpha 1 [Source:HGNC Symbol;Acc:2192]	0.89	0.88	0.96
			2.00	2.50

SLC12A1	solute carrier family 12 (sodium/potassium/chloride transporters), member 1 [Source:HGNC Symbol;Acc:10910]	0.87	0.88	0.95
BTBD2	BTB (POZ) domain containing 2 [Source:HGNC Symbol;Acc:15504]	0.87	0.88	0.95
CCL26	chemokine (C-C motif) ligand 26 [Source:HGNC Symbol;Acc:10625]	0.89	0.9	0.97
STAG3	stromal antigen 3 [Source:HGNC Symbol;Acc:11356]	0.39	0.4	0.47
THEM5	thioesterase superfamily member 5 [Source:HGNC Symbol;Acc:26755]	0.88	0.86	0.95
	X-prolyl aminopeptidase (aminopeptidase P) 1, soluble [Source:HGNC			
XPNPEP1	Symbol;Acc:12822]	0.62	0.6	0.69
ATL1	atlastin GTPase 1 [Source:HGNC Symbol;Acc:11231]	0.93	0.91	1
TRAPPC5	trafficking protein particle complex 5 [Source:HGNC Symbol;Acc:23067]	0.6	0.62	0.69
APOA1	apolipoprotein A-I [Source:HGNC Symbol;Acc:600]	0.9	0.87	0.97
JOSD2	Josephin domain containing 2 [Source:HGNC Symbol;Acc:28853] UDP glucuronosyltransferase 2 family, polypeptide B28 [Source:HGNC	0.76	0.73	0.83
UGT2B28	Symbol;Acc:13479]	0.86	0.89	0.96
C1S	complement component 1, s subcomponent [Source:HGNC Symbol;Acc:1247]	0.87	0.83	0.94
FLJ34503	uncharacterized	0.88	0.84	0.95
	olfactory receptor, family 2, subfamily B, member 3 [Source:HGNC			
OR2B3	Symbol;Acc:8238]	0.9	0.84	0.97
CHGB	chromogranin B (secretogranin 1) [Source:HGNC Symbol;Acc:1930]	0.28	0.21	0.35
	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,			
YWHAH	eta polypeptide [Source:HGNC Symbol;Acc:12853]	0.8	0.89	0.96
FAM131B	family with sequence similarity 131, member B [Source:HGNC Symbol;Acc:22202]	0.4	0.5	0.57
TAIVITSTB	ATP-binding cassette, sub-family D (ALD), member 2 [Source:HGNC	0.4	0.5	0.57
ABCD2	Symbol;Acc:66]	0.34	0.24	0.41
ANO2	anoctamin 2 [Source:HGNC Symbol;Acc:1183]	0.67	0.78	0.85
	olfactory receptor, family 2, subfamily G, member 2 [Source:HGNC			
OR2G2	Symbol;Acc:15007]	0.88	0.75	0.95
DGKA	diacylglycerol kinase, alpha 80kDa [Source:HGNC Symbol;Acc:2849]	0.54	0.67	0.74
GKN2	gastrokine 2 [Source:HGNC Symbol;Acc:24588]	0.64	0.5	0.71
MOGAT3	monoacylglycerol O-acyltransferase 3 [Source:HGNC Symbol;Acc:23249]	0.56	0.42	0.63
RAB27A TYMP	RAB27A, member RAS oncogene family [Source:HGNC Symbol;Acc:9766]	0.91	0.74	0.98
GRM2	thymidine phosphorylase [Source:HGNC Symbol;Acc:3148] glutamate receptor, metabotropic 2 [Source:HGNC Symbol;Acc:4594]	0.49 0.79	0.66 0.53	0.73 0.86
CELA1	chymotrypsin-like elastase family, member 1 [Source:HGNC Symbol;Acc:3308]	0.75	0.33	0.82
GMFG	glia maturation factor, gamma [Source:HGNC Symbol;Acc:4374]	0.75	0.46	0.82
ZNF208	zinc finger protein 208 [Source:HGNC Symbol;Acc:12999]	0.72	0.72	0.78
TMEM184B	transmembrane protein 184B [Source:HGNC Symbol;Acc:1310]	0.5	0.5	0.56
EPS8L3	EPS8-like 3 [Source:HGNC Symbol;Acc:21297]	0.82	0.81	0.88
CLNK	cytokine-dependent hematopoietic cell linker [Source:HGNC Symbol;Acc:17438]	0.93	0.94	1
TSPAN32 C11orf21	tetraspanin 32 [Source:HGNC Symbol;Acc:13410] chromosome 11 open reading frame 21 [Source:HGNC Symbol;Acc:13231]	0.87 0.87	0.86 0.86	0.93 0.93
C1101121	solute carrier family 14 (urea transporter), member 1 (Kidd blood group)	0.67	0.86	0.95
SLC14A1	[Source:HGNC Symbol;Acc:10918]	0.83	0.82	0.89
LAD1	ladinin 1 [Source:HGNC Symbol;Acc:6472]	0.86	0.87	0.93
VGLL3	vestigial like 3 (Drosophila) [Source:HGNC Symbol;Acc:24327]	0.47	0.48	0.54
PABPC4L	poly(A) binding protein, cytoplasmic 4-like [Source:HGNC Symbol;Acc:31955]	0.74	0.75	0.81
	olfactory receptor, family 8, subfamily H, member 1 [Source:HGNC			
OR8H1	Symbol;Acc:14824]	0.94	0.92	1
ICLD	immunoglobulin superfamily containing leucine-rich repeat [Source:HGNC	0.02	0.01	0.00
ISLR HCFC2	Symbol;Acc:6133]	0.83	0.81	0.89
	• • •		0.04	1
TICICZ	host cell factor C2 [Source:HGNC Symbol;Acc:24972]	0.92	0.94	1
	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC	0.92		
BCS1L	host cell factor C2 [Source:HGNC Symbol;Acc:24972]		0.94	0.55
	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC	0.92		
BCS1L	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020]	0.92	0.47	0.55
BCS1L TCF15 IFIT2	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409]	0.92 0.49 0.84 0.83	0.47 0.82 0.85	0.55 0.9 0.91
BCS1L TCF15 IFIT2 BEX5	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990]	0.92 0.49 0.84 0.83 0.59	0.47 0.82 0.85 0.61	0.55 0.9 0.91 0.67
BCS1L TCF15 IFIT2 BEX5 DHH	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865]	0.92 0.49 0.84 0.83 0.59 0.84	0.47 0.82 0.85 0.61 0.81	0.55 0.9 0.91 0.67 0.9
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044]	0.92 0.49 0.84 0.83 0.59 0.84 0.82	0.47 0.82 0.85 0.61 0.81 0.79	0.55 0.9 0.91 0.67 0.9 0.88
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83	0.47 0.82 0.85 0.61 0.81 0.79 0.8	0.55 0.9 0.91 0.67 0.9 0.88 0.89
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9 TLR9	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91	0.55 0.9 0.91 0.67 0.9 0.88 0.89
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83	0.47 0.82 0.85 0.61 0.81 0.79 0.8	0.55 0.9 0.91 0.67 0.9 0.88 0.89
BCS1L TCF15 IFIT2 BEX5 DHH C200rf141 CST9 TLR9 TNNC2	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:G3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9 TLR9 TNNC2 BFSP2	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 [fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:1041]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9 TLR9 TNNC2 BFSP2 C9orf117 HSD3B1	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:1041] chromosome 9 open reading frame 117 [Source:HGNC Symbol;Acc:27843]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9
BCS1L TCF15 IFIT2 BEX5 DHH C200rf141 C200rf141 TR9 TNNC2 BFSP2 C90rf117 HSD3B1 TAC4	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:27843] hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9 TIR9 TNNC2 BFSP2 C9orf117 HSD3B1 TAC4 KLK4	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:G3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:27843] hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641] kallikrein-related peptidase 4 [Source:HGNC Symbol;Acc:6365]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79 0.87	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88 0.93
BCS1L TCF15 IFIT2 BEX5 DHH C200rf141 C200rf141 TR9 TNNC2 BFSP2 C90rf117 HSD3B1 TAC4	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:G3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:27843] hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641] kallikrein-related peptidase 4 [Source:HGNC Symbol;Acc:6365] receptor accessory protein 5 [Source:HGNC Symbol;Acc:30077]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88
BCS1L TCF15 IFIT2 BEX5 DHH C200rf141 CST9 TLR9 TNNC2 BFSP2 C90rf117 HSD3B1 TAC4 KLK4 REEP5	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:1041] chromosome 9 open reading frame 117 [Source:HGNC Symbol;Acc:27843] hydroxy-delta-s-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641] kallikrein-related peptidase 4 [Source:HGNC Symbol;Acc:30077] serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin),	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79 0.87 0.74 0.87 0.78	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88 0.93 0.8 0.93 0.88
BCS1L TCF15 IFIT2 BEX5 DHH C20orf141 CST9 TLR9 TNNC2 BFSP2 C9orf117 HSD3B1 TAC4 KLK4 REEP5 SERPINA5	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:27843] hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641] kallikrein-related peptidase 4 [Source:HGNC Symbol;Acc:30077] serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 5 [Source:HGNC Symbol;Acc:8723]	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79 0.87 0.74 0.87 0.78	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82 0.84 0.71 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88 0.93 0.8 0.93 0.8 0.93
BCS1L TCF15 IFIT2 BEX5 DHH C200rf141 CST9 TLR9 TNNC2 BFSP2 C90rf117 HSD3B1 TAC4 KLK4 REEP5	host cell factor C2 [Source:HGNC Symbol;Acc:24972] BC1 (ubiquinol-cytochrome c reductase) synthesis-like [Source:HGNC Symbol;Acc:1020] transcription factor 15 (basic helix-loop-helix) [Source:HGNC Symbol;Acc:11627] interferon-induced protein with tetratricopeptide repeats 2 [Source:HGNC Symbol;Acc:5409] brain expressed, X-linked 5 [Source:HGNC Symbol;Acc:27990] desert hedgehog [Source:HGNC Symbol;Acc:2865] transmembrane protein 239 [Source:HGNC Symbol;Acc:40044] cystatin 9 (testatin) [Source:HGNC Symbol;Acc:13261] TLR9 [Source:UniProtKB/TrEMBL;Acc:C3W5P5] troponin C type 2 (fast) [Source:HGNC Symbol;Acc:11944] beaded filament structural protein 2, phakinin [Source:HGNC Symbol;Acc:1041] chromosome 9 open reading frame 117 [Source:HGNC Symbol;Acc:27843] hydroxy-delta-s-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 [Source:HGNC Symbol;Acc:5217] tachykinin 4 (hemokinin) [Source:HGNC Symbol;Acc:16641] kallikrein-related peptidase 4 [Source:HGNC Symbol;Acc:30077] serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin),	0.92 0.49 0.84 0.83 0.59 0.84 0.82 0.83 0.94 0.91 0.81 0.79 0.87 0.74 0.87 0.78	0.47 0.82 0.85 0.61 0.81 0.79 0.8 0.91 0.94 0.84 0.82	0.55 0.9 0.91 0.67 0.9 0.88 0.89 1 1 0.9 0.88 0.93 0.8 0.93 0.88

PMP22	norinhard muslin protain 22 [Saurca: HCNC Symbol Acc; 0110]	0.72	0.66	0.78
FCRLB	peripheral myelin protein 22 [Source:HGNC Symbol;Acc:9118] Fc receptor-like B [Source:HGNC Symbol;Acc:26431]	0.72	0.00	0.78
CHRM1	cholinergic receptor, muscarinic 1 [Source:HGNC Symbol;Acc:1950]	0.84	0.78	0.9
FHB	EF-hand domain family, member B [Source:HGNC Symbol;Acc:26330]	0.67	0.73	0.79
19orf69	chromosome 19 open reading frame 69 [Source:HGNC Symbol;Acc:34497]	0.69	0.61	0.75
USD3	sushi domain containing 3 [Source:HGNC Symbol;Acc:28391]	0.5	0.58	0.64
CRYAA	crystallin, alpha A [Source:HGNC Symbol;Acc:2388]	0.86	0.77	0.92
CNIAT1	guanine nucleotide binding protein (G protein), alpha transducing activity	0.07	0.70	0.93
SNAT1	polypeptide 1 [Source:HGNC Symbol;Acc:4393]	0.87	0.78	
GCK PAX8	glucokinase (hexokinase 4) [Source:HGNC Symbol;Acc:4195] paired box 8 [Source:HGNC Symbol;Acc:8622]	0.86 0.75	0.76 0.86	0.92 0.92
RIM58	tripartite motif containing 58 [Source:HGNC Symbol;Acc:24150]	0.73	0.86	0.92
HAVCR2	hepatitis A virus cellular receptor 2 [Source:HGNC Symbol;Acc:18437]	0.83	0.71	0.89
OL9A2	collagen, type IX, alpha 2 [Source:HGNC Symbol;Acc:2218]	0.59	0.72	0.78
CXCL13	chemokine (C-X-C motif) ligand 13 [Source:HGNC Symbol;Acc:10639]	0.87	0.73	0.93
CA2	oculocutaneous albinism II [Source:HGNC Symbol;Acc:8101]	0.71	0.86	0.92
AM65A	family with sequence similarity 65, member A [Source:HGNC Symbol;Acc:25836]	0.59	0.74	0.8
HIST1H4L	histone cluster 1, H4a [Source:HGNC Symbol;Acc:4781]	0.59	0.74	0.79
SLC24A5	solute carrier family 24, member 5 [Source:HGNC Symbol;Acc:20611]	0.86	0.64	0.73
	family with sequence similarity 114, member A1 [Source:HGNC			
AM114A1	Symbol;Acc:25087] malic enzyme 3, NADP(+)-dependent, mitochondrial [Source:HGNC	0.27	0.49	0.55
ME3	Symbol;Acc:6985]	0.53	0.82	0.88
GRIP1	glutamate receptor interacting protein 1 [Source:HGNC Symbol;Acc:18708]	0.83	0.3	0.89
(IAA0922	KIAA0922 [Source:HGNC Symbol;Acc:29146]	0.21	0.21	0.27
ALPL	alkaline phosphatase, liver/bone/kidney [Source:HGNC Symbol;Acc:438]	0.21	0.2	0.27
SYT3	synaptotagmin III [Source:HGNC Symbol;Acc:11511]	0.22	0.21	0.28
RBM15B	RNA binding motif protein 15B [Source:HGNC Symbol;Acc:24303]	0.21	0.19	0.27
MPRIP	myosin phosphatase Rho interacting protein [Source:HGNC Symbol;Acc:30321]	0.21	0.18	0.27
FD A B 4 1 1 1	translocation associated membrane protein 1-like 1 [Source:HGNC	0.10	0.21	0.27
TRAM1L1 NKX3-1	Symbol;Acc:28371] NK3 homeobox 1 [Source:HGNC Symbol;Acc:7838]	0.18	0.21 0.11	0.27 0.17
NKX3-1	solute carrier organic anion transporter family, member 2A1 [Source:HGNC	0.08	0.11	0.17
SLCO2A1	Symbol;Acc:10955]	0.04	0.08	0.14
	family with sequence similarity 196, member B [Source:HGNC	0.00	0.00	0.4.4
AM196B	Symbol;Acc:37271]	0.08	0.02	0.14
CCDC147	coiled-coil domain containing 147 [Source:HGNC Symbol;Acc:26676]	0.01	0.01	0.07
GDF6	growth differentiation factor 6 [Source:HGNC Symbol;Acc:4221]	0	0	0.06
24442	dishevelled associated activator of morphogenesis 2 [Source:HGNC Symbol;Acc:18143]	0.45	0.45	0.51
DAAM2 FGF8	fibroblast growth factor 8 (androgen-induced) [Source:HGNC Symbol;Acc:3686]	0.45	0.45	0.51
UI 8	ClpB caseinolytic peptidase B homolog (E. coli) [Source:HGNC	0.02	0.02	0.08
CLPB	Symbol;Acc:30664]	0.33	0.33	0.39
KIAA1755	KIAA1755 [Source:HGNC Symbol;Acc:29372]	0	0	0.06
NORD12B	small nucleolar RNA, C/D box 12B [Source:HGNC Symbol;Acc:33573]	0	0	0.06
	hydroxyprostaglandin dehydrogenase 15-(NAD) [Source:HGNC			
HPGD	Symbol;Acc:5154]	0	0	0.06
C4orf33	chromosome 4 open reading frame 33 [Source:HGNC Symbol;Acc:27025]	0	0	0.06
LC35B2	solute carrier family 35, member B2 [Source:HGNC Symbol;Acc:16872]	0.07	0.07	0.13
ACTL6B	actin-like 6B [Source:HGNC Symbol;Acc:160]	0.35	0.35	0.41
HKR1	HKR1, GLI-Kruppel zinc finger family member [Source:HGNC Symbol;Acc:4928] kelch-like 32 (Drosophila) [Source:HGNC Symbol;Acc:21221]	0.17	0.18	0.24
CLHL32 FMCO4	transmembrane and coiled-coil domains 4 [Source:HGNC Symbol;Acc:27393]	0.23	0.24 0.05	0.3 0.12
RRC3	leucine rich repeat containing 3 [Source:HGNC Symbol;Acc:27595]	0.06	0.05	0.12
SYPC	glycophorin C (Gerbich blood group) [Source:HGNC Symbol;Acc:4704]	0.00	0.03	0.12
CSNK1G3	casein kinase 1, gamma 3 [Source:HGNC Symbol;Acc:2456]	0.14	0.13	0.2
	ectonucleoside triphosphate diphosphohydrolase 3 [Source:HGNC			
NTPD3	Symbol;Acc:3365]	0.31	0.32	0.38
RLIM	ring finger protein, LIM domain interacting [Source:HGNC Symbol;Acc:13429]	0.37	0.38	0.44
CHODL	chondrolectin [Source:HGNC Symbol;Acc:17807]	0.24	0.22	0.3
RPL39L	ribosomal protein L39-like [Source:HGNC Symbol;Acc:17094]	0.06	0.04	0.12
TK32C	serine/threonine kinase 32C [Source:HGNC Symbol;Acc:21332]	0.13	0.11	0.19
EB2	zinc finger E-box binding homeobox 2 [Source:HGNC Symbol;Acc:14881]	0.03	0.01	0.09
ЛARK1	MAP/microtubule affinity-regulating kinase 1 [Source:HGNC Symbol;Acc:6896] fumarylacetoacetate hydrolase domain containing 2A [Source:HGNC	0.01	0.03	0.09
AHD2A	Symbol;Acc:24252]	0.31	0.33	0.39
STX17	syntaxin 17 [Source:HGNC Symbol;Acc:11432]	0.12	0.14	0.2
AS1L	LAS1-like (S. cerevisiae) [Source:HGNC Symbol;Acc:25726]	0.38	0.4	0.46
C11orf35	chromosome 11 open reading frame 35 [Source:HGNC Symbol;Acc:28561]	0.15	0.12	0.21
	Ras association (RalGDS/AF-6) domain family (N-terminal) member 7			
RASSF7	[Source:HGNC Symbol;Acc:1166]	0.15	0.12	0.21
SNORA26	Small nucleolar RNA SNORA26 [Source:RFAM;Acc:RF00568]	0.06	0.03	0.12
	protein tyrosine phosphatase, receptor-type, Z polypeptide 1 [Source:HGNC			
PTPRZ1	Symbol;Acc:9685]	0.06	0.03	0.12
ABR	active BCR-related [Source:HGNC Symbol;Acc:81]	0.13	0.09	0.19

GRM1	glutamate receptor, metabotropic 1 [Source:HGNC Symbol;Acc:4593]	0.26	0.22	0.32
CCS	copper chaperone for superoxide dismutase [Source:HGNC Symbol;Acc:1613]	0.26	0.22	0.32
CXorf22	chromosome X open reading frame 22 [Source:HGNC Symbol;Acc:28546]	0.41	0.45	0.51
NKAIN1	Na+/K+ transporting ATPase interacting 1 [Source:HGNC Symbol;Acc:25743]	0.41	0.02	0.31
C1orf131	chromosome 1 open reading frame 131 [Source:HGNC Symbol;Acc:25745]	0.07	0.02	0.13
GNPAT	glyceronephosphate O-acyltransferase [Source:HGNC Symbol;Acc:24416]	0.12	0.06	0.18
DCHS1	dachsous 1 (Drosophila) [Source:HGNC Symbol;Acc:13681]	0.12	0.03	0.15
SUPT3H	suppressor of Ty 3 homolog (S. cerevisiae) [Source:HGNC Symbol;Acc:11466]	0.09	0.03	0.15
	,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,, ,,		0.04	
TGFB2	transforming growth factor, beta 2 [Source:HGNC Symbol;Acc:11768]	0.04		0.16
HOXA6	homeobox A6 [Source:HGNC Symbol;Acc:5107]	0.14	0.2	0.26
KIF12	kinesin family member 12 [Source:HGNC Symbol;Acc:21495]	0.11	0.17	0.23
RPL18A	Small nucleolar RNA SNORA68 [Source:RFAM;Acc:RF00263]	0.28	0.21	0.34
USH1G	Usher syndrome 1G (autosomal recessive) [Source:HGNC Symbol;Acc:16356]	0.25	0.17	0.31
MT1G	metallothionein 1G [Source:HGNC Symbol;Acc:7399]	0.11	0.19	0.25
LDB2	LIM domain binding 2 [Source:HGNC Symbol;Acc:6533]	0.25	0.34	0.4
VDAC2	voltage-dependent anion channel 2 [Source:HGNC Symbol;Acc:12672]	0.15	0.04	0.21
COL1A1	collagen, type I, alpha 1 [Source:HGNC Symbol;Acc:2197]	0.07	0.19	0.25
S100A10	S100 calcium binding protein A10 [Source:HGNC Symbol;Acc:10487]	0.01	0.17	0.23
PITPNM3	PITPNM family member 3 [Source:HGNC Symbol;Acc:21043]	0.15	0.38	0.44
HSF1	heat shock transcription factor 1 [Source:HGNC Symbol;Acc:5224]	0.32	0.06	0.38
BTBD17	BTB (POZ) domain containing 17 [Source:HGNC Symbol;Acc:33758]	0.39	0.12	0.45
	phosphatidylinositol-specific phospholipase C, X domain containing 3			
PLCXD3	[Source:HGNC Symbol;Acc:31822]	0.21	0.23	0.29
PKD1L1	polycystic kidney disease 1 like 1 [Source:HGNC Symbol;Acc:18053]	0.91	0.91	0.97
LCN1	lipocalin 1 [Source:HGNC Symbol;Acc:6525]	0.9	0.9	0.96
ZNF735	zinc finger protein 679 [Source:HGNC Symbol;Acc:28650]	0.9	0.89	0.96
LHX8	LIM homeobox 8 [Source:HGNC Symbol;Acc:28838]	0.89	0.9	0.96
	amine oxidase, copper containing 2 (retina-specific) [Source:HGNC			
AOC2	Symbol;Acc:549]	0.91	0.92	0.98
	transglutaminase 2 (C polypeptide, protein-glutamine-gamma-			
TGM2	glutamyltransferase) [Source:HGNC Symbol;Acc:11778]	0.88	0.86	0.94
	nuclear receptor subfamily 1, group H, member 4 [Source:HGNC			
NR1H4	Symbol;Acc:7967]	0.88	0.86	0.94
ELANE	elastase, neutrophil expressed [Source:HGNC Symbol;Acc:3309]	0.76	0.74	0.82
IL24	interleukin 24 [Source:HGNC Symbol;Acc:11346]	0.9	0.92	0.98
WFDC9	WAP four-disulfide core domain 9 [Source:HGNC Symbol;Acc:20380]	0.9	0.87	0.96
DIXDC1	DIX domain containing 1 [Source:HGNC Symbol;Acc:23695]	0.76	0.79	0.85
BEST3	bestrophin 3 [Source:HGNC Symbol;Acc:17105]	0.88	0.91	0.97
MSMB	microseminoprotein, beta- [Source:HGNC Symbol;Acc:7372]	0.91	0.87	0.97
NFE2L3	nuclear factor (erythroid-derived 2)-like 3 [Source:HGNC Symbol;Acc:7783]	0.91	0.87	0.97
	olfactory receptor, family 9, subfamily A, member 2 [Source:HGNC			
OR9A2	Symbol;Acc:15093]	0.93	0.89	0.99
	transient receptor potential cation channel, subfamily A, member 1			
TRPA1	[Source:HGNC Symbol;Acc:497]	0.61	0.65	0.71
C9orf24	chromosome 9 open reading frame 24 [Source:HGNC Symbol;Acc:19919]	0.76	0.8	0.86
CORO1A	coronin, actin binding protein, 1A [Source:HGNC Symbol;Acc:2252]	0.79	0.74	0.85
FBXO41	F-box protein 41 [Source:HGNC Symbol;Acc:29409]	0.91	0.86	0.97
10041	v-ets erythroblastosis virus E26 oncogene homolog (avian) [Source:HGNC	0.51	0.00	0.57
ERG	Symbol;Acc:3446]	0.73	0.78	0.84
TMEM194B	transmembrane protein 194B [Source:HGNC Symbol;Acc:33700]	0.73	0.88	0.94
ARHGDIB	Rho GDP dissociation inhibitor (GDI) beta [Source:HGNC Symbol;Acc:679]	0.82	0.88	0.83
RASAL3	RAS protein activator like 3 [Source:HGNC Symbol;Acc:26129]	0.78	0.71	0.83
VDTOF	keratin 85 [Source:HGNC Symbol;Acc:6462]	0.00	0.04	0.05
KR185 AD∩F	apolipoprotein F [Source:HGNC Symbol;Acc:615]	0.89	0.81	0.95
APOF	chromosome 3 open reading frame 18 [Source:HGNC Symbol;Acc:24837]	0.9	0.82	0.96
C3orf18	, , , ,	0.71	0.79	0.85
C9orf152	chromosome 9 open reading frame 152 [Source:HGNC Symbol;Acc:31455]	0.88	0.78	0.94
CCBE1	collagen and calcium binding EGF domains 1 [Source:HGNC Symbol;Acc:29426]	0.41	0.51	0.57
FHL1	four and a half LIM domains 1 [Source:HGNC Symbol;Acc:3702]	0.81	0.71	0.87
THBS4	thrombospondin 4 [Source:HGNC Symbol;Acc:11788]	0.41	0.52	0.58
TDC1505	TBC1 domain family, member 8B (with GRAM domain) [Source:HGNC	0.00	0.50	0.74
TBC1D8B	Symbol;Acc:24715]	0.68	0.56	0.74
GPR173	G protein-coupled receptor 173 [Source:HGNC Symbol;Acc:18186]	0.52	0.4	0.58
COL8A2	collagen, type VIII, alpha 2 [Source:HGNC Symbol;Acc:2216]	0.41	0.53	0.59
	integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41)			
ITGA2B	[Source:HGNC Symbol;Acc:6138]	0.75	0.88	0.94
KIF3C	kinesin family member 3C [Source:HGNC Symbol;Acc:6321]	0.53	0.66	0.72
KNG1	kininogen 1 [Source:HGNC Symbol;Acc:6383]	0.56	0.43	0.62
	UDP glucuronosyltransferase 2 family, polypeptide B4 [Source:HGNC			
UGT2B4	Symbol;Acc:12553]	0.92	0.75	0.98
DENND1C	DENN/MADD domain containing 1C [Source:HGNC Symbol;Acc:26225]	0.56	0.36	0.62
SYTL3	synaptotagmin-like 3 [Source:HGNC Symbol;Acc:15587]	0.67	0.45	0.73
ULBP3	UL16 binding protein 3 [Source:HGNC Symbol;Acc:14895]	0.77	0.53	0.83
ECHDC2	enoyl CoA hydratase domain containing 2 [Source:HGNC Symbol;Acc:23408]	0.51	0.78	0.84
FGG	fibrinogen gamma chain [Source:HGNC Symbol;Acc:3694]	0.92	0.63	0.98
FAM26E	family with sequence similarity 26, member E [Source:HGNC Symbol;Acc:21568]	0.79	0.45	0.85

CPZ	carboxypeptidase Z [Source:HGNC Symbol;Acc:2333]	0.67	0.33	0.73
ADAD2	adenosine deaminase domain containing 2 [Source:HGNC Symbol;Acc:30714]	0.87	0.87	0.92
	solute carrier family 10 (sodium/bile acid cotransporter family), member 4			
SLC10A4	[Source:HGNC Symbol;Acc:22980]	0.7	0.7	0.75
PRKG2	protein kinase, cGMP-dependent, type II [Source:HGNC Symbol;Acc:9416]	0.88	0.88	0.93
ADRB2	adrenoceptor beta 2, surface [Source:HGNC Symbol;Acc:286]	0.85	0.86	0.91
	ribosomal protein S6 kinase, 90kDa, polypeptide 1 [Source:HGNC			
RPS6KA1	Symbol;Acc:10430]	0.87	0.86	0.92
BCAS1	breast carcinoma amplified sequence 1 [Source:HGNC Symbol;Acc:974]	0.95	0.94	1
	potassium channel tetramerisation domain containing 6 [Source:HGNC			
KCTD6	Symbol;Acc:22235]	0.84	0.83	0.89
BTBD16	BTB (POZ) domain containing 16 [Source:HGNC Symbol;Acc:26340]	0.85	0.86	0.91
	interferon-induced protein with tetratricopeptide repeats 1B [Source:HGNC			
IFIT1B	Symbol;Acc:23442]	0.75	0.76	0.81
TPD52	tumor protein D52 [Source:HGNC Symbol;Acc:12005]	0.7	0.71	0.76
	T cell immunoreceptor with Ig and ITIM domains [Source:HGNC			
TIGIT	Symbol;Acc:26838]	0.84	0.82	0.89
AMOT	angiomotin [Source:HGNC Symbol;Acc:17810]	0.86	0.88	0.93
CALHM2	calcium homeostasis modulator 2 [Source:HGNC Symbol;Acc:23493]	0.7	0.67	0.75
ZNF713	zinc finger protein 713 [Source:HGNC Symbol;Acc:22043]	0.95	0.92	1
BAALC	brain and acute leukemia, cytoplasmic [Source:HGNC Symbol;Acc:14333]	0.81	0.84	0.89
GLA	galactosidase, alpha [Source:HGNC Symbol;Acc:4296]	0.49	0.46	0.54
	heterogeneous nuclear ribonucleoprotein H2 (H') [Source:HGNC			
HNRNPH2	Symbol;Acc:5042]	0.49	0.46	0.54
PDZD4	PDZ domain containing 4 [Source:HGNC Symbol;Acc:21167]	0.69	0.72	0.77
	synuclein, gamma (breast cancer-specific protein 1) [Source:HGNC			
SNCG	Symbol;Acc:11141]	0.85	0.81	0.9
STARD3NL	STARD3 N-terminal like [Source:HGNC Symbol;Acc:19169]	0.91	0.95	1
	matrix metallopeptidase 12 (macrophage elastase) [Source:HGNC			
MMP12	Symbol;Acc:7158]	0.86	0.82	0.91
	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1			
SULT1A1	[Source:HGNC Symbol;Acc:11453]	0.83	0.78	0.88
TRIM38	tripartite motif containing 38 [Source:HGNC Symbol;Acc:10059]	0.85	0.8	0.9
TMC4	transmembrane channel-like 4 [Source:HGNC Symbol;Acc:22998]	0.54	0.59	0.64
PROM2	prominin 2 [Source:HGNC Symbol;Acc:20685]	0.66	0.71	0.76
	receptor (chemosensory) transporter protein 3 [Source:HGNC			
RTP3	Symbol;Acc:15572]	0.74	0.69	0.79
RHD	Rh blood group, CcEe antigens [Source:HGNC Symbol;Acc:10008]	0.7	0.75	0.8
GPA33	glycoprotein A33 (transmembrane) [Source:HGNC Symbol;Acc:4445]	0.94	0.88	0.99
	family with sequence similarity 186, member B [Source:HGNC			
FAM186B	Symbol;Acc:25296]	0.83	0.77	0.88
TTC22	tetratricopeptide repeat domain 22 [Source:HGNC Symbol;Acc:26067]	0.77	0.83	0.88
SPOCD1	SPOC domain containing 1 [Source:HGNC Symbol;Acc:26338]	0.51	0.57	0.62
	protein phosphatase 2, regulatory subunit B', beta [Source:HGNC			
PPP2R5B	Symbol;Acc:9310]	0.41	0.35	0.46
KRTAP10-2	keratin associated protein 10-2 [Source:HGNC Symbol;Acc:22967]	0.73	0.66	0.78
C1orf189	chromosome 1 open reading frame 189 [Source:HGNC Symbol;Acc:32305]	0.87	0.79	0.92
TSPAN2	tetraspanin 2 [Source:HGNC Symbol;Acc:20659]	0.88	0.8	0.93
XAF1	XIAP associated factor 1 [Source:HGNC Symbol;Acc:30932]	0.84	0.76	0.89
	glutamic-pyruvate transaminase (alanine aminotransferase) [Source:HGNC			
GPT	Symbol;Acc:4552]	0.85	0.77	0.9
	nephrosis 2, idiopathic, steroid-resistant (podocin) [Source:HGNC			
NPHS2	Symbol;Acc:13394]	0.86	0.94	0.99
	nudix (nucleoside diphosphate linked moiety X)-type motif 18 [Source:HGNC	0.00	0.5 .	0.55
NUDT18	Symbol:Acc:26194]	0.55	0.63	0.68
C10orf11	chromosome 10 open reading frame 11 [Source:HGNC Symbol;Acc:23405]	0.84	0.75	0.89
C100//11	signal-induced proliferation-associated 1 like 1 [Source:HGNC	0.04	0.75	0.03
SIPA1L1	Symbol;Acc:20284]	0.95	0.86	1
C20orf202	chromosome 20 open reading frame 202 [Source:HGNC Symbol;Acc:37254]	0.82	0.73	0.87
OIT3	oncoprotein induced transcript 3 [Source:HGNC Symbol;Acc:29953]	0.86	0.95	1
0113	endothelial cell surface expressed chemotaxis and apoptosis regulator	0.80	0.55	1
ECSCR	[Source:HGNC Symbol;Acc:35454]	0.67	0.76	0.81
KIAA0513	KIAA0513 [Source:HGNC Symbol;Acc:29058]	0.95	0.76	1
KIAAUJIJ	· · · · · · · · · · · · · · · · · · ·	0.55	0.65	1
TNESEA	tumor necrosis factor (ligand) superfamily, member 4 [Source:HGNC Symbol;Acc:11934]	0.85	0.72	0.9
TNFSF4	keratin 71 [Source:HGNC Symbol;Acc:28927]		0.72	0.9
KRT71		0.83		
RNASE13	ribonuclease, RNase A family, 13 (non-active) [Source:HGNC Symbol;Acc:25285]	0.83	0.69	0.88
C5orf20	chromosome 5 open reading frame 20 [Source:HGNC Symbol;Acc:24459]	0.86	0.71	0.91
TAS1R2	taste receptor, type 1, member 2 [Source:HGNC Symbol;Acc:14905]	0.62	0.46	0.67
FCRL6	Fc receptor-like 6 [Source:HGNC Symbol;Acc:31910]	0.88	0.72	0.93
PGAM2	phosphoglycerate mutase 2 (muscle) [Source:HGNC Symbol;Acc:8889]	0.51	0.7	0.75
DNAH2	dynein, axonemal, heavy chain 2 [Source:HGNC Symbol;Acc:2948]	0.73	0.51	0.78
IKZF1	IKAROS family zinc finger 1 (Ikaros) [Source:HGNC Symbol;Acc:13176]	0.58	0.36	0.63
	transmembrane BAX inhibitor motif containing 1 [Source:HGNC			
TMBIM1	Symbol;Acc:23410]	0.58	0.82	0.87
MORC2	MORC family CW-type zinc finger 2 [Source:HGNC Symbol;Acc:23573]	0.47	0.71	0.76

	olfactory receptor, family 52, subfamily J, member 3 [Source:HGNC			
OR52J3	Symbol;Acc:14799]	0.87	0.6	0.92
	low density lipoprotein receptor-related protein associated protein 1			
LRPAP1	[Source:HGNC Symbol;Acc:6701]	0.48	0.12	0.53
LRRC23 C11orf88	leucine rich repeat containing 23 [Source:HGNC Symbol;Acc:19138] chromosome 11 open reading frame 88 [Source:HGNC Symbol;Acc:25061]	0.48 0.14	0.08 0.15	0.53 0.2
HAND2	heart and neural crest derivatives expressed 2 [Source:HGNC Symbol;Acc:4808]	0.09	0.13	0.14
WDR63	WD repeat domain 63 [Source:HGNC Symbol;Acc:30711]	0.12	0.1	0.17
GLIPR1L1	GLI pathogenesis-related 1 like 1 [Source:HGNC Symbol;Acc:28392]	0.15	0.13	0.2
	regulatory factor X, 5 (influences HLA class II expression) [Source:HGNC			
RFX5	Symbol;Acc:9986]	0.09	0.06	0.14
SPEF1	sperm flagellar 1 [Source:HGNC Symbol;Acc:15874]	0.15	0.12	0.2
TUBA8 PHYH	tubulin, alpha 8 [Source:HGNC Symbol;Acc:12410] phytanoyl-CoA 2-hydroxylase [Source:HGNC Symbol;Acc:8940]	0.05 0.12	0.09 0.07	0.14 0.17
SLC25A42	solute carrier family 25, member 42 [Source:HGNC Symbol;Acc:28380]	0.12	0.07	0.26
NKX2-3	NK2 homeobox 3 [Source:HGNC Symbol;Acc:7836]	0.16	0.22	0.27
SH2B3	SH2B adaptor protein 3 [Source:HGNC Symbol;Acc:29605]	0.04	0.12	0.17
KSR2	kinase suppressor of ras 2 [Source:HGNC Symbol;Acc:18610]	0.12	0.21	0.26
	radical S-adenosyl methionine domain containing 1 [Source:HGNC			
RSAD1	Symbol;Acc:25634]	0.12	0.02	0.17
LBX2	ladybird homeobox 2 [Source:HGNC Symbol;Acc:15525]	0.07	0.18	0.23
PRDM13 C8orf73	PR domain containing 13 [Source:HGNC Symbol;Acc:13998] maestro heat-like repeat family member 6 [Source:HGNC Symbol;Acc:27814]	0.18 0.09	0.06 0.22	0.23 0.27
SALL4	sal-like 4 (Drosophila) [Source:HGNC Symbol;Acc:27814]	0.08	0.22	0.28
SHE	Src homology 2 domain containing E [Source:HGNC Symbol;Acc:27004]	0.05	0.05	0.1
TDRD10	tudor domain containing 10 [Source:HGNC Symbol;Acc:25316]	0.05	0.05	0.1
	ligand dependent nuclear receptor corepressor [Source:HGNC			
LCOR	Symbol;Acc:29503]	0.03	0.03	0.08
HHEX	hematopoietically expressed homeobox [Source:HGNC Symbol;Acc:4901]	0.02	0.02	0.07
GDF10	growth differentiation factor 10 [Source:HGNC Symbol;Acc:4215]	0	0	0.05
DLX4	distal-less homeobox 4 [Source:HGNC Symbol;Acc:2917] neuralized homolog 3 (Drosophila) pseudogene [Source:HGNC	0.05	0.05	0.1
NEURL3	Symbol;Acc:25162]	0.03	0.03	0.08
EVX1	even-skipped homeobox 1 [Source:HGNC Symbol;Acc:3506]	0.02	0.02	0.07
DECR1	2,4-dienoyl CoA reductase 1, mitochondrial [Source:HGNC Symbol;Acc:2753]	0.02	0.02	0.07
	family with sequence similarity 129, member A [Source:HGNC			
FAM129A	Symbol;Acc:16784]	0.11	0.1	0.16
ANK3	ankyrin 3, node of Ranvier (ankyrin G) [Source:HGNC Symbol;Acc:494]	0.03	0.02	0.08
APOA1BP GPR135	apolipoprotein A-I binding protein [Source:HGNC Symbol;Acc:18453] G protein-coupled receptor 135 [Source:HGNC Symbol;Acc:19991]	0.05 0.01	0.04 0.02	0.1 0.07
FZD5	frizzled family receptor 5 [Source:HGNC Symbol;Acc:4943]	0.01	0.02	0.07
1203	peroxisome proliferator-activated receptor alpha [Source:HGNC	0.01	0.02	0.07
PPARA	Symbol;Acc:9232]	0.03	0.01	0.08
PRDM8	PR domain containing 8 [Source:HGNC Symbol;Acc:13993]	0.03	0.01	0.08
EBF4	early B-cell factor 4 [Source:HGNC Symbol;Acc:29278]	0.09	0.11	0.16
HMOX1	heme oxygenase (decycling) 1 [Source:HGNC Symbol;Acc:5013]	0.04	0.06	0.11
PPP1CC	protein phosphatase 1, catalytic subunit, gamma isozyme [Source:HGNC Symbol;Acc:9283]	0.06	0.03	0.11
PRKCH	protein kinase C, eta [Source:HGNC Symbol;Acc:9403]	0.05	0.03	0.16
TRICH	fibroblast growth factor (acidic) intracellular binding protein [Source:HGNC	0.03	0.11	0.10
FIBP	Symbol;Acc:3705]	0.01	0.01	0.06
BICD1	bicaudal D homolog 1 (Drosophila) [Source:HGNC Symbol;Acc:1049]	0.04	0.04	0.09
SRI	sorcin [Source:HGNC Symbol;Acc:11292]	0.01	0.01	0.06
CCDC88C	coiled-coil domain containing 88C [Source:HGNC Symbol;Acc:19967]	0.04	0.03	0.09
DACCES	Ras association (RalGDS/AF-6) domain family member 2 [Source:HGNC	0	0.01	0.00
RASSF2 RRAD	Symbol;Acc:9883] Ras-related associated with diabetes [Source:HGNC Symbol;Acc:10446]	0 0.17	0.01 0.17	0.06 0.22
OSBPL5	oxysterol binding protein-like 5 [Source:HGNC Symbol;Acc:16446]	0.17	0.17	0.36
DFNB59	deafness, autosomal recessive 59 [Source:HGNC Symbol;Acc:29502]	0.13	0.12	0.18
COL23A1	collagen, type XXIII, alpha 1 [Source:HGNC Symbol;Acc:22990]	0.25	0.24	0.3
ESD	esterase D [Source:HGNC Symbol;Acc:3465]	0.06	0.07	0.12
EXOC3L2	exocyst complex component 3-like 2 [Source:HGNC Symbol;Acc:30162]	0.15	0.16	0.21
YIPF6	Yip1 domain family, member 6 [Source:HGNC Symbol;Acc:28304]	0.37	0.38	0.43
EVC	Ellis van Creveld syndrome [Source:HGNC Symbol;Acc:3497]	0.14	0.16	0.21
SQLE TMEM128	squalene epoxidase [Source:HGNC Symbol;Acc:11279] transmembrane protein 128 [Source:HGNC Symbol;Acc:28201]	0.17 0.28	0.19 0.26	0.24 0.33
ALS2CL	ALS2 C-terminal like [Source:HGNC Symbol;Acc:20605]	0.28	0.26	0.53
	HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1	J	5.10	0.01
HECW1	[Source:HGNC Symbol;Acc:22195]	0.24	0.21	0.29
	microtubule-associated protein, RP/EB family, member 3 [Source:HGNC			
MAPRE3	Symbol;Acc:6892]	0.04	0.07	0.12
CLIC6	chloride intracellular channel 6 [Source:HGNC Symbol;Acc:2065]	0.13	0.16	0.21
AVPR1B	arginine vasopressin receptor 1B [Source:HGNC Symbol;Acc:896]	0.43	0.46	0.51
RASA2 CCDC87	RAS p21 protein activator 2 [Source:HGNC Symbol;Acc:9872] coiled-coil domain containing 87 [Source:HGNC Symbol;Acc:25579]	0.17 0.32	0.13 0.37	0.22 0.42
CCDCG/	conca con domain containing or [Jource.Howe Symbol,Acc.25575]	0.52	0.37	0.42

	family with sequence similarity 150, member B [Source:HGNC			
FAM150B	Symbol;Acc:27683]	0.02	0.07	0.12
MTA3	metastasis associated 1 family, member 3 [Source:HGNC Symbol;Acc:23784]	0.18	0.24	0.29
TRIB2	tribbles homolog 2 (Drosophila) [Source:HGNC Symbol;Acc:30809]	0.13	0.19	0.24
DUSP19	dual specificity phosphatase 19 [Source:HGNC Symbol;Acc:18894]	0.27	0.21	0.32
-GR6	leucine-rich repeat containing G protein-coupled receptor 6 [Source:HGNC Symbol;Acc:19719]	0.31	0.24	0.36
LUNU	wingless-type MMTV integration site family, member 10B [Source:HGNC	0.51	0.24	0.50
WNT10B	Symbol;Acc:12775]	0.16	0.24	0.29
MBNL1	muscleblind-like splicing regulator 1 [Source:HGNC Symbol;Acc:6923]	0.46	0.38	0.51
3MBTL4	l(3)mbt-like 4 (Drosophila) [Source:HGNC Symbol;Acc:26677]	0.15	0.24	0.29
MATN2	matrilin 2 [Source:HGNC Symbol;Acc:6908]	0.05	0.14	0.19
EFNB3	ephrin-B3 [Source:HGNC Symbol;Acc:3228]	0.04	0.14	0.19
BP1	fructose-1,6-bisphosphatase 1 [Source:HGNC Symbol;Acc:3606]	0.04	0.14	0.19
TRIM45	tripartite motif containing 45 [Source:HGNC Symbol;Acc:19018]	0.16	0.05	0.21
HTR7	5-hydroxytryptamine (serotonin) receptor 7, adenylate cyclase-coupled [Source:HGNC Symbol;Acc:5302]	0.19	0.33	0.38
SUCLG1	succinate-CoA ligase, alpha subunit [Source:HGNC Symbol;Acc:11449]	0.19	0.33	0.38
OCLOI	inhibitor of Bruton agammaglobulinemia tyrosine kinase [Source:HGNC	0.24	0.03	0.23
ВТК	Symbol;Acc:17853]	0.25	0.1	0.3
	enoyl-CoA, hydratase/3-hydroxyacyl CoA dehydrogenase [Source:HGNC			
HHADH	Symbol;Acc:3247]	0.23	0.39	0.44
ABHD6	abhydrolase domain containing 6 [Source:HGNC Symbol;Acc:21398]	0.28	0.45	0.5
ISD17B8	hydroxysteroid (17-beta) dehydrogenase 8 [Source:HGNC Symbol;Acc:3554]	0.14	0.36	0.41
AFF3	AF4/FMR2 family, member 3 [Source:HGNC Symbol;Acc:6473]	0.13	0.38	0.43
NNT3	troponin T type 3 (skeletal, fast) [Source:HGNC Symbol;Acc:11950]	0.89	0.89	0.94
REC8	REC8 homolog (yeast) [Source:HGNC Symbol;Acc:16879]	0.92	0.92	0.97
AP30BP	SAP30 binding protein [Source:HGNC Symbol;Acc:30785]	0.78	0.78	0.83
RECQL5	RecQ protein-like 5 [Source:HGNC Symbol;Acc:9950]	0.79	0.79	0.84
ANKRD23 L1A	ankyrin repeat domain 23 [Source:HGNC Symbol;Acc:24470] interleukin 1, alpha [Source:HGNC Symbol;Acc:5991]	0.92 0.68	0.92 0.68	0.97 0.73
MUC4	mucin 4, cell surface associated [Source:HGNC Symbol;Acc:7514]	0.08	0.08	0.73
VIOCH	solute carrier family 36 (proton/amino acid symporter), member 3	0.52	0.52	0.57
SLC36A3	[Source:HGNC Symbol;Acc:19659]	0.92	0.92	0.97
	olfactory receptor, family 14, subfamily J, member 1 [Source:HGNC			
OR14J1	Symbol;Acc:13971]	0.9	0.9	0.95
HEPHL1	hephaestin-like 1 [Source:HGNC Symbol;Acc:30477]	0.8	0.79	0.85
	fucosyltransferase 8 (alpha (1,6) fucosyltransferase) [Source:HGNC			
UT8	Symbol;Acc:4019]	0.9	0.89	0.95
CD300C	CD300c molecule [Source:HGNC Symbol;Acc:19320]	0.89	0.88	0.94
	uncharacterized serine/threonine-protein kinase SgK494 [Source:RefSeq			
GK494	peptide;Acc:NP_001167574]	0.9	0.89	0.95
CDH26	cadherin 26 [Source:HGNC Symbol;Acc:15902]	0.93	0.92	0.98
LTA	lymphotoxin alpha (TNF superfamily, member 1) [Source:HGNC Symbol;Acc:6709]	0.91	0.9	0.96
LIA	cytochrome P450, family 3, subfamily A, polypeptide 5 [Source:HGNC	0.51	0.9	0.50
CYP3A5	Symbol;Acc:2638]	0.92	0.91	0.97
511 57 15	potassium intermediate/small conductance calcium-activated channel,	0.52	0.51	0.57
CNN4	subfamily N, member 4 [Source:HGNC Symbol;Acc:6293]	0.88	0.89	0.94
SNLY	granulysin [Source:HGNC Symbol;Acc:4414]	0.92	0.93	0.98
TTPAL	tocopherol (alpha) transfer protein-like [Source:HGNC Symbol;Acc:16114]	0.64	0.65	0.7
NF883	zinc finger protein 883 [Source:HGNC Symbol;Acc:27271]	0.67	0.68	0.73
SRRB	estrogen-related receptor beta [Source:HGNC Symbol;Acc:3473]	0.9	0.88	0.95
XOSC9	exosome component 9 [Source:HGNC Symbol;Acc:9137]	0.66	0.64	0.71
TDO2	tryptophan 2,3-dioxygenase [Source:HGNC Symbol;Acc:11708]	0.9	0.88	0.95
	solute carrier family 6 (neurotransmitter transporter, GABA), member 11	c		
SLC6A11	[Source:HGNC Symbol;Acc:11044]	0.87	0.89	0.94
GSTA3	glutathione S-transferase alpha 3 [Source:HGNC Symbol;Acc:4628]	0.87	0.89	0.94
NVNIDIN	NYN domain and retroviral integrase containing [Source:HGNC Symbol;Acc:20165]	0.8	0.77	0.85
NYNRIN MAGEA5	melanoma antigen family A, 5 [Source:HGNC Symbol;Acc:6803]	0.8	0.77	0.85
	ATP-binding cassette, sub-family A (ABC1), member 12 [Source:HGNC	0.55	0.5	0.50
ABCA12	Symbol;Acc:14637]	0.87	0.9	0.95
	leucine-rich repeats and calponin homology (CH) domain containing 1			
RCH1	[Source:HGNC Symbol;Acc:20309]	0.76	0.8	0.85
GLT1D1	glycosyltransferase 1 domain containing 1 [Source:HGNC Symbol;Acc:26483]	0.86	0.9	0.95
	SAM pointed domain containing ets transcription factor [Source:HGNC			
PDEF	Symbol;Acc:17257]	0.91	0.86	0.96
MEM176A	transmembrane protein 176A [Source:HGNC Symbol;Acc:24930]	0.77	0.72	0.82
SLP2R	glucagon-like peptide 2 receptor [Source:HGNC Symbol;Acc:4325]	0.75	0.8	0.85
JSP51	ubiquitin specific peptidase 51 [Source:HGNC Symbol;Acc:23086]	0.5	0.55	0.6
	dehydrogenase/reductase (SDR family) member 9 [Source:HGNC			
DHRS9	Symbol;Acc:16888]	0.79	0.73	0.84
DTDN7	protein tyrosine phosphatase, non-receptor type 7 [Source:HGNC	0.63	0.00	0.07
PTPN7	Symbol;Acc:9659]	0.92	0.86	0.97

	ATPase, H+ transporting, lysosomal V0 subunit a4 [Source:HGNC			
ATP6V0A4	Symbol;Acc:866]	0.79	0.73	0.84
TMEM213	transmembrane protein 213 [Source:HGNC Symbol;Acc:27220]	0.79	0.73	0.84
GOT1L1	glutamic-oxaloacetic transaminase 1-like 1 [Source:HGNC Symbol;Acc:28487]	0.92	0.86	0.97
OTUD5	OTU domain containing 5 [Source:HGNC Symbol;Acc:25402]	0.89	0.83	0.94
	olfactory receptor, family 6, subfamily V, member 1 [Source:HGNC			
OR6V1	Symbol;Acc:15090]	0.89	0.82	0.94
	epidermal growth factor receptor pathway substrate 15 [Source:HGNC			
EPS15	Symbol;Acc:3419]	0.57	0.64	0.69
TFF2	trefoil factor 2 [Source:HGNC Symbol;Acc:11756]	0.93	0.85	0.98
	olfactory receptor, family 5, subfamily AS, member 1 [Source:HGNC			
OR5AS1	Symbol;Acc:15261]	0.9	0.81	0.95
	potassium inwardly-rectifying channel, subfamily J, member 10 [Source:HGNC			
KCNJ10	Symbol;Acc:6256]	0.92	0.83	0.97
	C1q and tumor necrosis factor related protein 8 [Source:HGNC			
C1QTNF8	Symbol;Acc:31374]	0.9	0.8	0.95
	olfactory receptor, family 10, subfamily K, member 1 [Source:HGNC			
OR10K1	Symbol;Acc:14693]	0.93	0.83	0.98
	prostaglandin E receptor 1 (subtype EP1), 42kDa [Source:HGNC			
PTGER1	Symbol;Acc:9593]	0.77	0.66	0.82
CTSE	cathepsin E [Source:HGNC Symbol;Acc:2530]	0.93	0.82	0.98
	cytochrome P450, family 2, subfamily B, polypeptide 6 [Source:HGNC			
CYP2B6	Symbol;Acc:2615]	0.92	0.8	0.97
	carcinoembryonic antigen-related cell adhesion molecule 19 [Source:HGNC			
CEACAM19	Symbol;Acc:31951]	0.68	0.56	0.73
BTN3A3	butyrophilin, subfamily 3, member A3 [Source:HGNC Symbol;Acc:1140]	0.77	0.63	0.82
RXRB	retinoid X receptor, beta [Source:HGNC Symbol;Acc:10478]	0.68	0.54	0.73
ZSCAN1	zinc finger and SCAN domain containing 1 [Source:HGNC Symbol;Acc:23712]	0.53	0.67	0.72
PGLYRP4	peptidoglycan recognition protein 4 [Source:HGNC Symbol;Acc:30015]	0.8	0.65	0.85
	integrin, alpha X (complement component 3 receptor 4 subunit) [Source:HGNC			
ITGAX	Symbol;Acc:6152]	0.89	0.72	0.94
EPHX4	epoxide hydrolase 4 [Source:HGNC Symbol;Acc:23758]	0.63	0.8	0.85
TPI1	triosephosphate isomerase 1 [Source:HGNC Symbol;Acc:12009]	0.4	0.66	0.71
C16orf3	chromosome 16 open reading frame 3 [Source:HGNC Symbol;Acc:1197]	0.67	0.38	0.72
DRP2	dystrophin related protein 2 [Source:HGNC Symbol;Acc:3032]	0.77	0.44	0.82
TUB	tubby homolog (mouse) [Source:HGNC Symbol;Acc:12406]	0.8	0.28	0.85