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Supplemental Data

Colocalization of GWAS and eQTL

Signals Detects Target Genes

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Supplemental Data

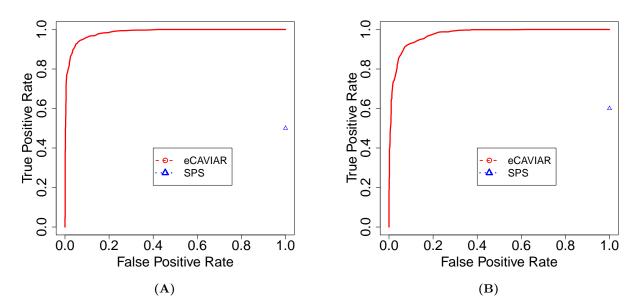


Figure S1. ROC curve computed for the same results as shown in Figures 3A and 3B where we generate the marginal statistics directly from the LD structure and we implant one causal variant in the locus. The x-axis is the false positive rate and the y-axis is the true positive rate. Panels (A) and (B) indicate the result for the case we have 50% and 80% for the causal variant, respectively.

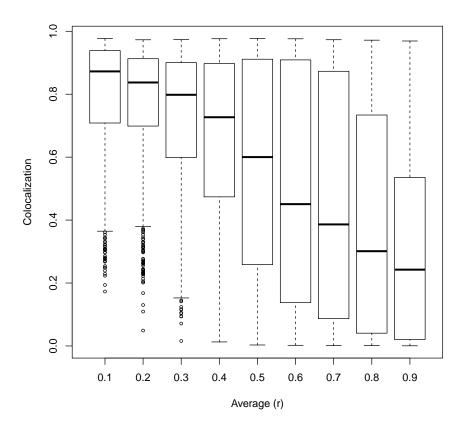


Figure S2. eCAVIAR quantification of CLPP depends on LD. We compute the probability of the same variant is causal in both GWAS and eQTL as we vary the Pearson's correlation between pair of variants. We observe that as we increase the correlation, which increase the LD complexity, the CLPP value decreases and the variance of the computed CLPP increases. We simulate datasets where we range the average Pearson's correlation between pair of variants from 0.1 to 0.9. In this setting, we implant only one causal variant in the same variant for both eQTL and GWAS. The NCP is set such that we have 50% power. Then, we simulate the marginal statistics for 1000 times.

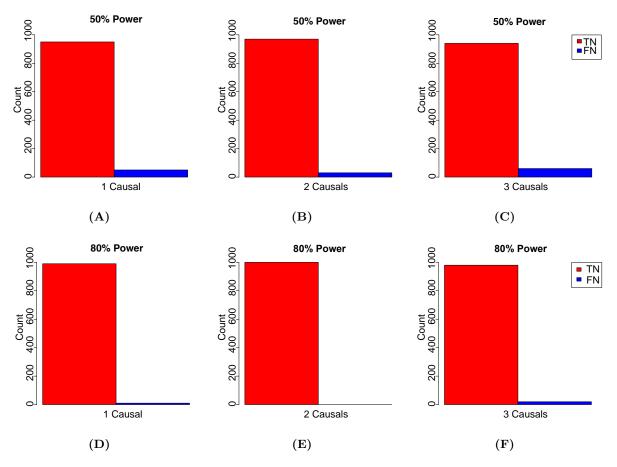
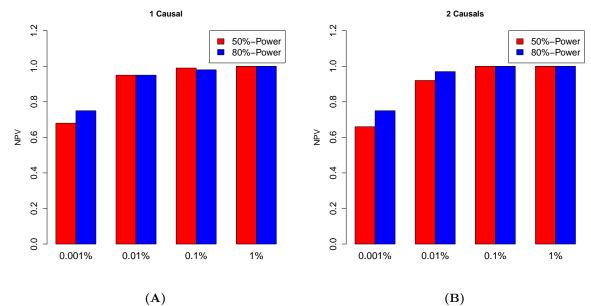


Figure S3. eCAVIAR has high true negative (TN) rate and low false negative (FN) rate. We set the non-colocalization cut-off threshold to 0.01%







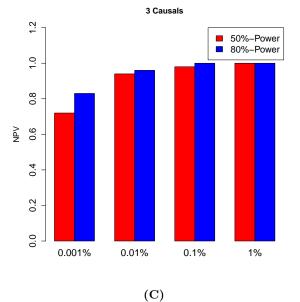


Figure S4. eCAVIAR has high negative predictive value (NPV). Panels (A), (B), and (C) illustrate the results for cases where we implant one, two, and three causal variants, respectively. We set NCP such that the statistical power at the causal variant is 50% or 80%.

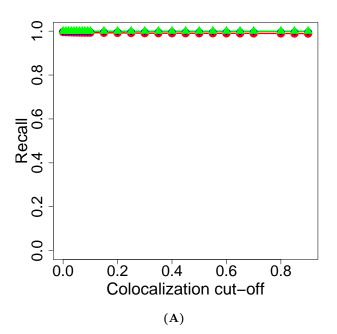


Figure S5. eCAVIAR, RTC, and COLOC have similar recall rate in the case where we have one causal variant in a locus. The x-axis is the colocalization cut-off threshold. We set the non-colocalization cut-off threshold to 0.1%.

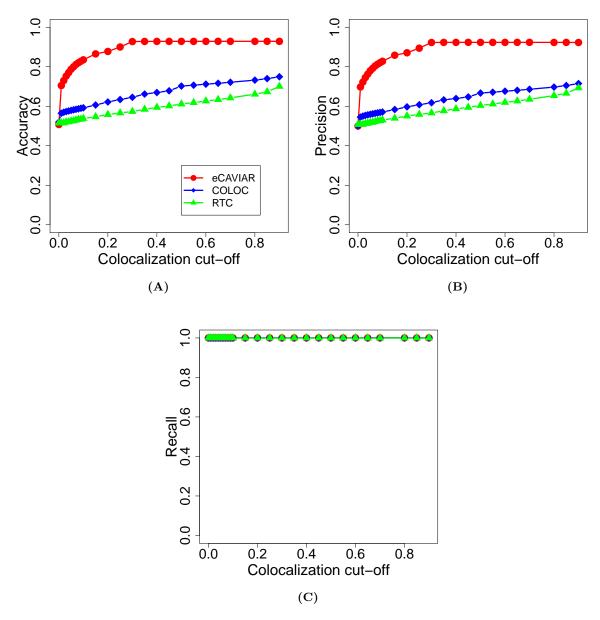


Figure S6. eCAVIAR is more accurate than existing methods for regions with one causal variant. We compared the accuracy and precision of eCAVIAR with the two existing methods (RTC and COLOC). The x-axis is the colocalization cut-off threshold. We set the non-colocalization cut-off threshold to 0.01%.

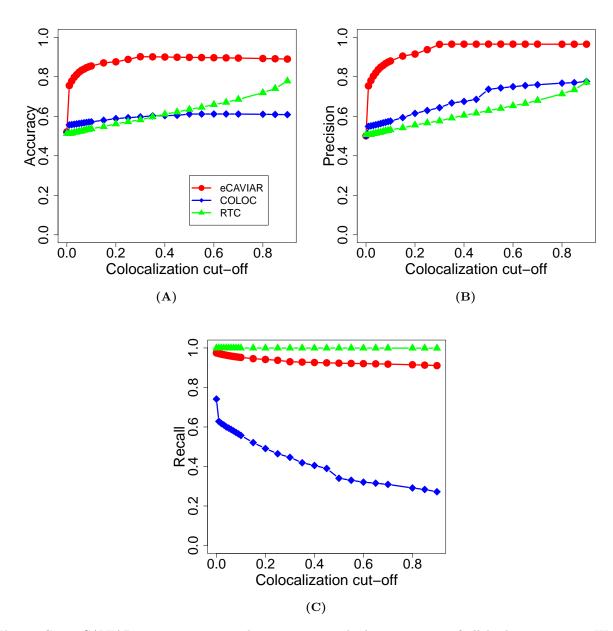


Figure S7. eCAVIAR is more accurate than existing methods in presence of allelic heterogeneity. We use similar process to generate the datasets as shown in Figure 4. However, in this case, we implant two causal variants. We set the non-colocalization cut-off threshold to 0.01%. RTC has better recall rare compared to eCAVIAR and COLOC.

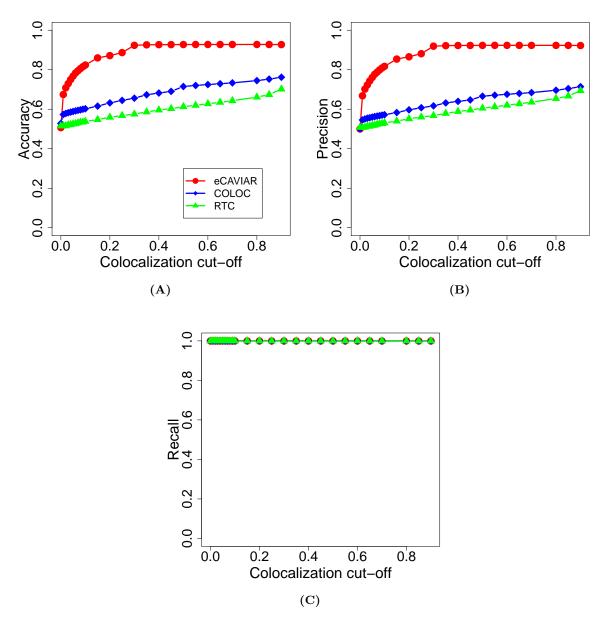


Figure S8. eCAVIAR outperforms existing methods when the causal variants are enriched with functional annotations. We implant one causal variant in our experiment.

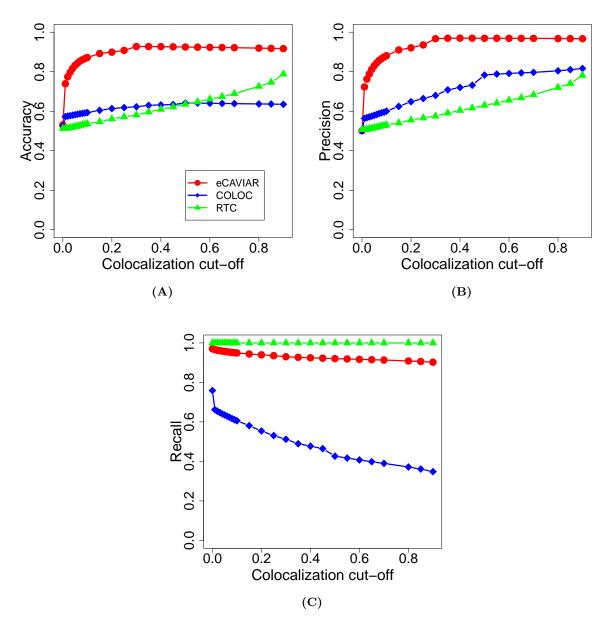


Figure S9. eCAVIAR outperforms existing methods when the causal variants are enriched with functional annotations. We implant two causal variants in our experiment.

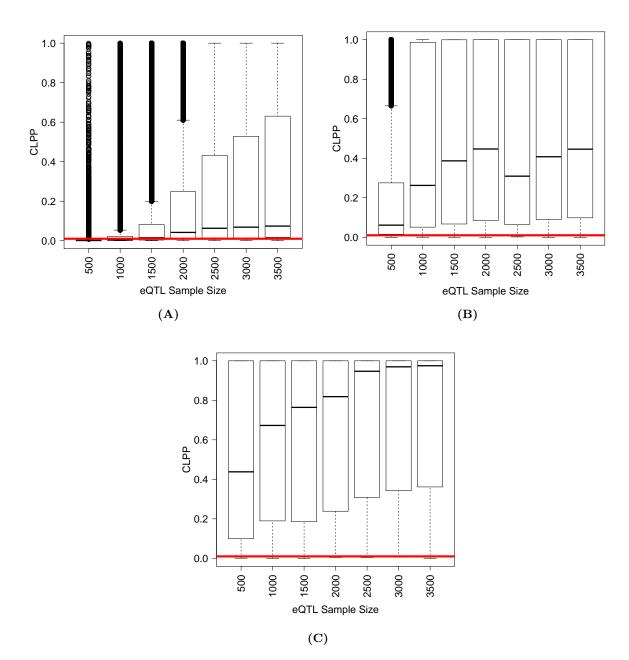


Figure S10. Effect of eQTL sample size on CLPP. The x-axis is the sample size that we vary from 500 to 3500. The y-axis is the distribution of CLPP. The red horizontal line indicates the 0.01 (1%) colocalization cut-off that is used our experiments. We set the sample size for GWAS to 5000. We simulate the effect size for colocalized variant such that it accounts for 0.01, 0.04, and 0.1 portion of the heritability as shown in panel (A), panel (B), and panel (C), respectively.

Tissue	Number of Individuals
Uterus	70
Brain Anterior cingulate cortex BA24	72
Small Intestine Terminal Ileum	77
Vagina	79
Brain Hippocampus	81
Brain Hypothalamus	81
Brain Putamen basal ganglia	82
Ovary	85
Pituitary	87
Prostate	87
Brain Cerebellar Hemisphere	89
Spleen	89
Brain Frontal Cortex BA9	92
Brain Nucleus accumbens basal ganglia	93
Brain Cortex	96
Liver	97
Brain Caudate basal ganglia	100
Brain Cerebellum	103
Cells EBV-transformed lymphocytes	114
Artery Coronary	118
Colon Sigmoid	124
Adrenal Gland	126
Esophagus Gastroesophageal Junction	127
Pancreas	149
Testis	157
Heart Atrial Appendage	159
Colon Transverse	169
Stomach	170
Breast Mammary Tissue	183
Adipose Visceral Omentum	185
Heart Left Ventricle	190
Skin Not Sun Exposed Suprapubic	196
Artery Aorta	197
Esophagus Muscularis	218
Esophagus Mucosa	241
Nerve Tibial	256
Cells Transformed fibroblasts	272
Lung	278
Thyroid	278
Artery Tibial	285
Adipose Subcutaneous	298
Skin Sun Exposed Lower leg	302
Whole Blood	338
Muscle Skeletal	361

Table S1. List of 44 tissues which are obtained from GTEx dataset. We provide the number of individuals genotyped for each tissue. Different tissues have different number of individuals.

Table S2. Potential colocalization loci for FG phenotype. Colocation loci consists of two summary statistics. One summary statistics obtained FG GWAS and the other summary statistics obtained for one of the genes in GTEx. Thus, we obtained these potential colocalization loci by considering the variants which are reported significant in previous study [?] for FG phenotype in MAGIC dataset. For each significant variant, we consider 50 variants upstream and downstream of the significant variants to define a locus. Next, we consider genes for all the tissues in GTEx data where at least one of the variants in the locus are significantly associated (corrected multiple hypothesis p-value of 10^{-5} or more significant) with its gene expression. We consider all the 44 tissues in GTEx to obtain the list of genes for each significant GWAS SNP.

rsID	Ensembl ID	Gene Name	Tissue	eQTL p-values
rs10830963	ENSG00000166002	SMCO4	Skin-Sun-Exposed-Lower-leg	2.44786305915e-05
rs10830963	ENSG00000166012	TAF1D	Brain-Nucleus-accumbens-basal-ganglia	1.7539865514e-05
rs10830963	ENSG00000180773	SLC36A4	Esophagus-Mucosa	5.2352746432e-05
rs10830963	ENSG00000182919	C11 or f54	Thyroid	8.61845523561e-06
rs10830963	ENSG00000206799	SNORA32	Skin-Sun-Exposed-Lower-leg	5.82640723736e-05
rs10830963	ENSG00000207145	SNORA18	Testis	4.2352972541e-05
rs10830963	ENSG00000255445	RP11-573M3.3	Heart-Atrial-Appendage	9.56125858181e-05
rs10885122	ENSG00000150594	ADRA2A	Brain-Nucleus-accumbens-basal-ganglia	2.57239942187e-10
rs10885122	ENSG00000150594	ADRA2A	Skin-Sun-Exposed-Lower-leg	8.13034309554e-05
rs10885122	ENSG00000203867	RBM20	Vagina	0.000126958323984
rs11071657	ENSG00000103642	LACTB	Whole-Blood	3.89333022107e-05
rs11071657	ENSG00000166104	RP11-299H22.3	Colon-Transverse	8.83954284179e-05
rs11071657	ENSG00000259498	RP11-244F12.3	Muscle-Skeletal	3.86262074991e-05
rs11071657	ENSG00000259627	RP11-244F12.2	Small-Intestine-Terminal-Ileum	2.27885479484e-05
rs11071657	ENSG00000259675	RP11-507B12.1	Vagina	6.91621612205e-05
rs11558471	ENSG00000147677	EIF3H	Spleen	3.9259893852e-05
rs11558471	ENSG00000164756	SLC30A8	Testis	2.05616636527e-05
rs11558471	ENSG00000205002	AARD	Uterus	2.36883921342e-05
rs11605924	ENSG00000121653	MAPK8IP1	Cells-EBV-transformed-lymphocytes	7.4647373768e-06
rs11605924	ENSG00000121653	MAPK8IP1	Lung	7.67362712574e-07
rs11605924	ENSG00000121653	MAPK8IP1	Skin-Not-Sun-Exposed-Suprapubic	5.74803471957e-22
rs11605924	ENSG00000121653	MAPK8IP1	Skin-Sun-Exposed-Lower-leg	4.60639977153e-17
rs11605924	ENSG00000121653	MAPK8IP1	Spleen	3.55628090171e-06
rs11605924	ENSG00000121653	MAPK8IP1	Whole-Blood	7.82629295485e-21
rs11605924	ENSG00000121671	CRY2	Testis	6.38769471885e-06
rs11605924	ENSG00000121671	CRY2	Whole-Blood	1.03433652463e-05
rs11605924	ENSG00000157613	CREB3L1	Adipose-Subcutaneous	0.000119111609495
rs11605924	ENSG00000165905	GYLTL1B	Heart-Atrial-Appendage	2.25819961703e-05
rs11605924	ENSG00000165905	GYLTL1B	Liver	1.24164721955e-05

rs11605924	ENSG00000175220	ARHGAP1	Uterus	1.90164370129e-05
rs11605924	ENSG00000175264	CHST1	Artery-Aorta	8.34252658299e-08
rs11605924	ENSG00000175264	CHST1	Thyroid	1.73199564995e-06
rs11605924	ENSG00000175274	TP53I11	Skin-Sun-Exposed-Lower-leg	0.000122950946627
rs11605924	ENSG00000180210	F2	Vagina	2.48171640639e-05
rs11605924	ENSG00000181830	SLC35C1	Artery-Tibial	2.34352827038e-06
rs11605924	ENSG00000181830	SLC35C1	Cells-Transformed-fibroblasts	6.250451226e-05
rs11605924	ENSG00000181830	SLC35C1	Esophagus-Mucosa	6.20248570017e-14
rs11605924	ENSG00000181830	SLC35C1	Heart-Left-Ventricle	7.89769212566e-05
rs11605924	ENSG00000181830	SLC35C1	Liver	2.18114395042e-06
rs11605924	ENSG00000181830	SLC35C1	Lung	3.26105205064e-05
rs11605924	ENSG00000181830	SLC35C1	Nerve-Tibial	2.348334931e-05
rs11605924	ENSG00000181830	SLC35C1	Whole-Blood	1.21176230508e-05
rs11605924	ENSG00000205106	DKFZp779M0652	Adipose-Subcutaneous	1.62392005396e-06
rs11605924	ENSG00000205106	DKFZp779M0652	Artery-Aorta	1.73680733946e-05
rs11605924	ENSG00000205106	DKFZp779M0652	Artery-Tibial	4.09586542776e-06
rs11605924	ENSG00000205106	DKFZp779 $M0652$	Brain-Hippocampus	9.98988593786e-06
rs11605924	ENSG00000205106	DKFZp779M0652	Brain-Hypothalamus	2.50756209116e-06
rs11605924	ENSG00000205106	DKFZp779 $M0652$	Brain-Nucleus-accumbens-basal-ganglia	4.27086076108e-07
rs11605924	ENSG00000205106	DKFZp779M0652	Brain-Putamen-basal-ganglia	4.81194257971e-06
rs11605924	ENSG00000205106	DKFZp779M0652	Colon-Transverse	3.77728053812e-05
rs11605924	ENSG00000205106	DKFZp779M0652	Esophagus-Mucosa	9.50866511829e-09
rs11605924	ENSG00000205106	DKFZp779M0652	Esophagus-Muscularis	1.02011808847e-08
rs11605924	ENSG00000205106	DKFZp779 $M0652$	Heart-Atrial-Appendage	1.03742684719e-05
rs11605924	ENSG00000205106	DKFZp779M0652	Heart-Left-Ventricle	1.3250774481e-08
rs11605924	ENSG00000205106	DKFZp779M0652	Liver	5.52158343567e-10
rs11605924	ENSG00000205106	DKFZp779 $M0652$	Lung	1.30661604158e-08
rs11605924	ENSG00000205106	DKFZp779 $M0652$	Nerve-Tibial	2.5524407916e-06
rs11605924	ENSG00000205106	DKFZp779M0652	Testis	1.65428551625e-10
rs11605924	ENSG00000205106	DKFZp779M0652	Thyroid	6.71470968327e-09
rs11605924	ENSG00000254427	RP11-430H10.1	Adipose-Visceral-Omentum	8.54818728207e-05
rs11605924	ENSG00000255091	RP11-495011.1	Artery-Aorta	1.87736272121e-06
rs11605924	ENSG00000255091	RP11-495011.1	Thyroid	3.08189576647e-05
rs11605924	ENSG00000255447	CTD-2210P24.6	Testis	3.8986007455e-07
rs11605924	ENSG00000255498	RP11-618K13.2	Skin-Not-Sun-Exposed-Suprapubic	4.70164342541e-07
rs11605924	ENSG00000255498	RP11-618K13.2	Skin-Sun-Exposed-Lower-leg	3.42801905208e-06

rs11605924	ENSG00000255498	RP11-618K13.2	Whole-Blood	4.42514908343e-07
rs11717195	ENSG0000065485	PDIA5	Artery-Tibial	5.02513492893e-07
rs11717195	ENSG0000065485	PDIA5	Thyroid	0.000126169254671
rs11717195	ENSG00000121542	SEC22A	Adipose-Subcutaneous	4.48128076072e-30
rs11717195	ENSG00000121542	SEC22A	Adipose-Visceral-Omentum	1.2210488429e-15
rs11717195	ENSG00000121542	SEC22A	Artery-Aorta	2.74502248777e-22
rs11717195	ENSG00000121542	SEC22A	Artery-Coronary	1.74364778894e-07
rs11717195	ENSG00000121542	SEC22A	Artery-Tibial	2.92641608999e-41
rs11717195	ENSG00000121542	SEC22A	Brain-Hippocampus	4.24221472915e-05
rs11717195	ENSG00000121542	SEC22A	Brain-Putamen-basal-ganglia	1.58717438504e-06
rs11717195	ENSG00000121542	SEC22A	Breast-Mammary-Tissue	8.14091243757e-26
rs11717195	ENSG00000121542	SEC22A	${\it Cells-EBV-transformed-lymphocytes}$	3.97453071673e-19
rs11717195	ENSG00000121542	SEC22A	Cells-Transformed-fibroblasts	1.72043210473e-21
rs11717195	ENSG00000121542	SEC22A	Colon-Sigmoid	3.32176009411e-05
rs11717195	ENSG00000121542	SEC22A	Colon-Transverse	1.32320584448e-16
rs11717195	ENSG00000121542	SEC22A	Esophagus-Gastroesophageal-Junction	6.23486130719e-08
rs11717195	ENSG00000121542	SEC22A	Esophagus-Mucosa	1.57111454052e-32
rs11717195	ENSG00000121542	SEC22A	Esophagus-Muscularis	3.02922278608e-18
rs11717195	ENSG00000121542	SEC22A	Heart-Atrial-Appendage	1.75855920111e-10
rs11717195	ENSG00000121542	SEC22A	Heart-Left-Ventricle	8.28685652528e-16
rs11717195	ENSG00000121542	SEC22A	Liver	4.55618336936e-05
rs11717195	ENSG00000121542	SEC22A	Lung	1.93513734862e-32
rs11717195	ENSG00000121542	SEC22A	Muscle-Skeletal	5.74835973093e-12
rs11717195	ENSG00000121542	SEC22A	Nerve-Tibial	8.64817990778e-31
rs11717195	ENSG00000121542	SEC22A	Ovary	7.13294372886e-08
rs11717195	ENSG00000121542	SEC22A	Pancreas	4.95208254925e-17
rs11717195	ENSG00000121542	SEC22A	Pituitary	2.83455224563e-06
rs11717195	ENSG00000121542	SEC22A	Prostate	3.8871388354e-10
rs11717195	ENSG00000121542	SEC22A	Skin-Not-Sun-Exposed-Suprapubic	2.78022705583e-24
rs11717195	ENSG00000121542	SEC22A	Skin-Sun-Exposed-Lower-leg	7.15947070871e-30
rs11717195	ENSG00000121542	SEC22A	Small-Intestine-Terminal-Ileum	9.43650095252e-05
rs11717195	ENSG00000121542	SEC22A	Spleen	3.35596740581e-12
rs11717195	ENSG00000121542	SEC22A	Stomach	4.71966998706e-09
rs11717195	ENSG00000121542	SEC22A	Testis	1.50510717516e-29
rs11717195	ENSG00000121542	SEC22A	Thyroid	2.90016216577e-26
rs11717195	ENSG00000121542	SEC22A	Uterus	2.5995012953e-05

rs11717195	ENSG00000121542	SEC 22A	Vagina	$4.99348818084 \mathrm{e}{\text{-}07}$	
rs11717195	ENSG00000121542	SEC 22A	Whole-Blood	1.01152955781e-08	
rs11717195	ENSG00000138463	DIRC2	Vagina	4.522896308e-05	
rs11717195	ENSG00000173175	ADCY5	Artery-Tibial	5.28198909019e-08	
rs11717195	ENSG00000173175	ADCY5	Colon-Sigmoid	2.41887725148e-05	
rs11717195	ENSG00000173175	ADCY5	Colon-Transverse	9.36426114519e-05	
rs11717195	ENSG00000173175	ADCY5	Esophagus-Mucosa	2.73445293874e-05	
rs11717195	ENSG00000173175	ADCY5	Esophagus-Muscularis	5.1190584425e-05	
rs11717195	ENSG00000173175	ADCY5	Heart-Atrial-Appendage	5.26218774438e-08	
rs11717195	ENSG00000173175	ADCY5	Heart-Left-Ventricle	1.9434499794e-06	
rs11717195	ENSG00000173175	ADCY5	Lung	2.25598951136e-07	
rs11717195	ENSG00000173175	ADCY5	Muscle-Skeletal	3.33007896392e-05	
rs11717195	ENSG00000173175	ADCY5	Thyroid	2.52528123532e-05	
rs11717195	ENSG00000173175	ADCY5	Uterus	9.56899015449e-06	
rs11717195	ENSG00000272678	RP11-797D24.4	Artery-Tibial	2.88661745263e-07	
rs11717195	ENSG00000272678	RP11-797D24.4	Esophagus-Mucosa	3.53458374031e-06	
rs11717195	ENSG00000272678	RP11-797D24.4	Heart-Atrial-Appendage	1.03080946347e-05	
rs11717195	ENSG00000272678	RP11-797D24.4	Heart-Left-Ventricle	4.61842170723e-08	
rs11717195	ENSG00000272678	RP11-797D24.4	Lung	2.87399425739e-06	
rs11920090	ENSG00000136603	SKIL	Ovary	2.52936436418e-05	
rs11920090	ENSG00000154310	TNIK	Pituitary	6.90410977459e-05	
rs11920090	ENSG00000163558	PRKCI	Skin-Sun-Exposed-Lower-leg	7.85139178977e-05	
rs11920090	ENSG00000163577	EIF5A2	Cells-Transformed-fibroblasts	3.45312321942e-05	
rs11920090	ENSG00000163584	RPL22L1	Whole-Blood	5.01595457402e-12	
rs11920090	ENSG00000173890	GPR160	Esophagus-Gastroesophageal-Junction	5.08237136088e-05	
rs11920090	ENSG00000242583	RP11-379K17.9	Testis	5.91788385319e-07	
rs174550	ENSG0000006118	TMEM132A	Ovary	5.65591312284e-05	
rs174550	ENSG00000110446	SLC15A3	Pituitary	6.79469593425e-05	
rs174550	ENSG00000124915	DKFZP434K028	Liver	1.52224793122e-15	
rs174550	ENSG00000124920	MYRF	Adipose-Subcutaneous	1.07163887489e-06	
rs174550	ENSG00000124920	MYRF	Artery-Tibial	1.81465783637e-06	
rs174550	ENSG00000124920	MYRF	Cells-Transformed-fibroblasts	2.05605381138e-10	
rs174550	ENSG00000124920	MYRF	Colon-Transverse	6.27262101649e-08	
rs174550	ENSG00000124920	MYRF	Liver	4.50726642799e-24	
rs174550	ENSG00000124920	MYRF	Pancreas	1.32784077095e-05	
rs174550	ENSG00000124920	MYRF	Whole-Blood	7.66855740345e-20	

rs174550	ENSG00000124942	AHNAK	Nerve-Tibial	6.84199477452e-05
rs174550	ENSG00000134780	DAGLA	Cells-Transformed-fibroblasts	4.14413546326e-05
rs174550	ENSG00000134780	DAGLA	Nerve-Tibial	2.13192293716e-06
rs174550	ENSG00000134780	DAGLA	Pancreas	8.3610269796e-06
rs174550	ENSG00000134824	FADS2	Adipose-Subcutaneous	1.47626352099e-06
rs174550	ENSG00000134824	FADS2	Adipose-Visceral-Omentum	2.86468845017e-06
rs174550	ENSG00000134824	FADS2	Artery-Aorta	3.47197673622e-11
rs174550	ENSG00000134824	FADS2	Artery-Tibial	8.53787978639e-08
rs174550	ENSG00000134824	FADS2	Breast-Mammary-Tissue	1.18893114978e-08
rs174550	ENSG00000134824	FADS2	Cells-EBV-transformed-lymphocytes	6.92791393224e-05
rs174550	ENSG00000134824	FADS2	Cells-Transformed-fibroblasts	4.37278719299e-38
rs174550	ENSG00000134824	FADS2	Colon-Sigmoid	5.75779994918e-09
rs174550	ENSG00000134824	FADS2	Colon-Transverse	3.87706737558e-23
rs174550	ENSG00000134824	FADS2	Esophagus-Gastroesophageal-Junction	1.25995286312e-10
rs174550	ENSG00000134824	FADS2	Esophagus-Muscularis	7.55772902317e-28
rs174550	ENSG00000134824	FADS2	Heart-Atrial-Appendage	1.26182123746e-08
rs174550	ENSG00000134824	FADS2	Lung	1.03018055989e-07
rs174550	ENSG00000134824	FADS2	Muscle-Skeletal	2.66948968982e-09
rs174550	ENSG00000134824	FADS2	Nerve-Tibial	4.08649453855e-16
rs174550	ENSG00000134824	FADS2	Pancreas	5.62568232448e-06
rs174550	ENSG00000134824	FADS2	Skin-Not-Sun-Exposed-Suprapubic	9.06036111248e-05
rs174550	ENSG00000134824	FADS2	Spleen	4.47894470381e-26
rs174550	ENSG00000134824	FADS2	Stomach	4.50207540976e-06
rs174550	ENSG00000134824	FADS2	Thyroid	2.18577113233e-23
rs174550	ENSG00000134824	FADS2	Whole-Blood	1.23516042549e-124
rs174550	ENSG00000134825	TMEM258	Adipose-Subcutaneous	4.06417386895e-06
rs174550	ENSG00000134825	TMEM258	Adipose-Visceral-Omentum	6.05997896336e-05
rs174550	ENSG00000134825	TMEM258	Artery-Tibial	4.30808695008e-07
rs174550	ENSG00000134825	TMEM258	Cells-EBV-transformed-lymphocytes	7.99556999059e-06
rs174550	ENSG00000134825	TMEM258	Cells-Transformed-fibroblasts	1.69327446094e-07
rs174550	ENSG00000134825	TMEM258	Colon-Transverse	1.3186362034e-05
rs174550	ENSG00000134825	TMEM258	Muscle-Skeletal	1.20748545687e-05
rs174550	ENSG00000134825	TMEM258	Nerve-Tibial	1.36798969082e-06
rs174550	ENSG00000134825	TMEM258	Ovary	9.92380755701e-05
rs174550	ENSG00000134825	TMEM258	Skin-Sun-Exposed-Lower-leg	3.3614144123e-08
rs174550	ENSG00000134825	TMEM258	Whole-Blood	1.31063979536e-07

rs174550	ENSG00000149480	MTA2	Liver	0.000112499358457
rs174550	ENSG00000149485	FADS1	Artery-Tibial	9.4043631547e-06
rs174550	ENSG00000149485	FADS1	Brain-Hippocampus	4.54093402282e-05
rs174550	ENSG00000149485	FADS1	Brain-Putamen-basal-ganglia	3.22433297405e-06
rs174550	ENSG00000149485	FADS1	Breast-Mammary-Tissue	1.74825888583e-05
rs174550	ENSG00000149485	FADS1	Colon-Sigmoid	1.33527941446e-05
rs174550	ENSG00000149485 ENSG00000149485	FADS1 FADS1	Liver	
rs174550		FADS1 FADS1	Liver Muscle-Skeletal	4.2645289605e-05 4.29712744478e-09
	ENSG00000149485			
rs174550	ENSG00000149485	FADS1	Nerve-Tibial	9.86394789532e-13
rs174550	ENSG00000149485	FADS1	Pancreas	3.71355853188e-25
rs174550	ENSG00000149485	FADS1	Stomach	1.10846086182e-10
rs174550	ENSG00000149485	FADS1	Thyroid	1.53366188198e-06
rs174550	ENSG00000149499	EML3	Skin-Not-Sun-Exposed-Suprapubic	7.6191865783e-05
rs174550	ENSG00000149506	ZP1	Adrenal-Gland	0.0001224879412
rs174550	ENSG00000162144	CYB561A3	Artery-Aorta	9.66918982602e-05
rs174550	ENSG00000162188	GNG3	Brain-Hippocampus	7.14451224699e-05
rs174550	ENSG00000162191	UBXN1	Brain-Putamen-basal-ganglia	2.14731353455e-05
rs174550	ENSG00000167987	VPS37C	Lung	8.86220323909e-06
rs174550	ENSG00000167995	BEST1	Stomach	6.31159912283e-05
rs174550	ENSG00000168000	BSCL2	Brain-Hypothalamus	1.93266607273e-05
rs174550	ENSG00000168002	POLR2G	Artery-Aorta	9.51846650203e-06
rs174550	ENSG00000207601	<i>MIR611</i>	Muscle-Skeletal	9.25132337759e-06
rs174550	ENSG00000207601	MIR611	Thyroid	9.42688703228e-05
rs174550	ENSG00000214753	HNRNPUL2	Colon-Transverse	7.55875326174e-05
rs174550	ENSG00000221968	FADS3	Liver	4.59871741566e-06
rs174550	ENSG00000255118	RP11-703H8.7	Cells-Transformed-fibroblasts	5.40761628183e-05
rs174550	ENSG00000257052	RP11-881M11.2	Skin-Sun-Exposed-Lower-leg	9.65492056964e-05
rs2191349	ENSG00000237070	LOC105375166	Colon-Transverse	8.91670843795e-05
rs4607517	ENSG00000015676	NUDCD3	Colon-Sigmoid	6.81574992936e-05
rs4607517	ENSG00000015676	NUDCD3	Colon-Transverse	0.000132421812821
rs4607517	ENSG00000058404	CAMK2B	Nerve-Tibial	1.02571474118e-06
rs4607517	ENSG00000106603	COA1	Muscle-Skeletal	9.88236081708e-05
rs4607517	ENSG00000106603	COA1	Small-Intestine-Terminal-Ileum	7.43495145463e-05
rs4607517	ENSG00000106608	URGCP	Brain-Hippocampus	0.000118741863422
rs4607517	ENSG00000106631	MYL7	Brain-Hypothalamus	2.03579173114e-07
rs4607517	ENSG00000106631	MYL7	Brain-Nucleus-accumbens-basal-ganglia	1.57407391493e-05

rs4607517	ENSG00000106633	GCK	Artery-Aorta	3.64341185775e-08
rs4607517	ENSG00000106633	GCK	Artery-Tibial	2.50853052784e-06
rs4607517	ENSG00000106633	GCK	Cells-Transformed-fibroblasts	2.59513829691e-06
rs4607517	ENSG00000106633	GCK	Colon-Sigmoid	7.91281792765e-05
rs4607517	ENSG00000106633	GCK	Ovary	2.74478432922e-05
rs4607517	ENSG00000106633	GCK	Thyroid	1.40524002295e-06
rs4607517	ENSG00000106636	YKT6	Artery-Aorta	4.64383379797e-05
rs4607517	ENSG00000106636	YKT6	Artery-Tibial	1.35586346349e-05
rs4607517	ENSG00000106636	YKT6	Muscle-Skeletal	1.3504504511e-06
rs4607517	ENSG00000106636	YKT6	Nerve-Tibial	3.77204249383e-08
rs4607517	ENSG00000106636	YKT6	Spleen	0.000110161962343
rs4607517	ENSG00000106636	YKT6	Testis	2.80798077625e-08
rs4607517	ENSG00000122678	POLM	Pituitary	0.000127963289669
rs4607517	ENSG00000214783	POLR2J4	Stomach	5.57197202642e-05
rs4607517	ENSG00000232956	SNHG15	Spleen	1.86473081598e-05
rs4607517	ENSG00000239775	AC017116.11	Artery-Tibial	1.11222953027e-05
rs4607517	ENSG00000239775	AC017116.11	Pituitary	8.06875463105e-05
rs4607517	ENSG00000239775	AC017116.11	Skin-Sun-Exposed-Lower-leg	2.66772000874e-05
rs4607517	ENSG00000272768	RP4-673M15.1	Whole-Blood	3.67370732112e-05
rs560887	ENSG00000144357	UBR3	Adipose-Subcutaneous	4.2131571751e-05
rs560887	ENSG00000152253	SPC25	Adipose-Subcutaneous	3.86223993369e-12
rs560887	ENSG00000152253	SPC25	Adipose-Visceral-Omentum	3.34610610317e-08
rs560887	ENSG00000152253	SPC25	Breast-Mammary-Tissue	1.94333203135e-09
rs560887	ENSG00000152253	SPC25	Esophagus-Gastroesophageal-Junction	6.57684285001e-05
rs560887	ENSG00000152253	SPC25	Esophagus-Muscularis	2.72793402053e-05
rs560887	ENSG00000152253	SPC25	Heart-Left-Ventricle	2.45718348331e-07
rs560887	ENSG00000152253	SPC25	Lung	1.32477367276e-10
rs560887	ENSG00000152253	SPC25	Nerve-Tibial	1.41909731656e-07
rs560887	ENSG00000152253	SPC25	Pancreas	6.27712479465e-14
rs560887	ENSG00000152253	SPC25	Stomach	7.4687660235e-08
rs560887	ENSG00000152253	SPC25	Thyroid	2.23167085387e-06
rs560887	ENSG00000152253	SPC25	Uterus	8.63781093856e-05
rs560887	ENSG00000152254	G6PC2	Adipose-Subcutaneous	2.29365890692e-44
rs560887	ENSG00000152254	G6PC2	Adipose-Visceral-Omentum	9.90863520284e-40
rs560887	ENSG00000152254	G6PC2	Breast-Mammary-Tissue	3.98413880387e-62
rs560887	ENSG00000152254	G6PC2	Colon-Transverse	3.06859548601e-09

rs560887	ENSG00000152254	G6PC2	Lung	1.77754225575e-69
rs560887	ENSG00000152254	G6PC2	Nerve-Tibial	9.62300344586e-30
rs560887	ENSG00000152254	G6PC2	Skin-Not-Sun-Exposed-Suprapubic	4.11488440163e-36
rs560887	ENSG00000152254	G6PC2	Skin-Sun-Exposed-Lower-leg	5.04823221975e-57
rs560887	ENSG00000152254	G6PC2	Stomach	5.70440304756e-21
rs560887	ENSG00000152254	G6PC2	Thyroid	1.60317874106e-34
rs560887	ENSG00000163072	NOSTRIN	Adipose-Subcutaneous	8.99742781521e-08
rs560887	ENSG00000163072	NOSTRIN	Adrenal-Gland	7.36729049722e-07
rs560887	ENSG00000163072	NOSTRIN	Artery-Tibial	3.92609755042e-06
rs560887	ENSG00000163072	NOSTRIN	Cells-EBV-transformed-lymphocytes	5.43395002041e-26
rs560887	ENSG00000163072	NOSTRIN	Colon-Transverse	3.26106663575e-06
rs560887	ENSG00000163072	NOSTRIN	Heart-Left-Ventricle	1.32053075953e-08
rs560887	ENSG00000163072	NOSTRIN	Lung	5.92637609704e-08
rs560887	ENSG00000163072	NOSTRIN	Nerve-Tibial	2.17221822661e-13
rs560887	ENSG00000163072	NOSTRIN	Pancreas	2.1316196208e-10
rs560887	ENSG00000163072	NOSTRIN	Prostate	1.59244224983e-06
rs560887	ENSG00000163072	NOSTRIN	Skin-Sun-Exposed-Lower-leg	7.2171093318e-09
rs560887	ENSG00000163072	NOSTRIN	Stomach	1.6447331884e-17
rs560887	ENSG00000163072	NOSTRIN	Testis	2.41334475961e-16
rs560887	ENSG00000163072	NOSTRIN	Thyroid	3.03601387041e-07
rs560887	ENSG00000172292	CERS6	Adipose-Visceral-Omentum	4.67914154219e-05
rs7034200	ENSG00000106686	SPATA6L	Heart-Atrial-Appendage	3.24945597547e-05
rs7034200	ENSG00000107249	GLIS3	Cells-Transformed-fibroblasts	3.45573413009e-11
rs7034200	ENSG00000107249	GLIS3	Thyroid	0.000119458479636
rs7034200	ENSG00000147853	AK3	Brain-Putamen-basal-ganglia	4.56481081455e-06
rs7034200	ENSG00000182347	PDSS1P1	Esophagus-Gastroesophageal-Junction	3.11010312751e-05
rs7034200	ENSG00000228322	RP11-358M14.2	Thyroid	4.11620980493e-05
rs7034200	ENSG00000230225	MTND5P14	Nerve-Tibial	0.000102595276953
rs7034200	ENSG00000235326	XXX	Stomach	7.05416646715e-05
rs7034200	ENSG00000237009	GLIS3-AS1	Cells-Transformed-fibroblasts	1.5014164511e-08
rs780094	ENSG0000084734	GCKR	Spleen	7.2120706573e-16
rs780094	ENSG0000084734	GCKR	Thyroid	2.79663819232e-09
rs780094	ENSG0000084774	CAD	Thyroid	5.43528025657e-06
rs780094	ENSG00000115216	NRBP1	Adipose-Subcutaneous	6.33052712903e-10
rs780094	ENSG00000115216	NRBP1	Adipose-Visceral-Omentum	9.96713371471e-06
rs780094	ENSG00000115216	NRBP1	Adrenal-Gland	3.81105034801e-05

rs780094	ENSG00000115216	NRBP1	Artery-Aorta	9.15815289491e-06
rs780094	ENSG00000115216	NRBP1	Artery-Coronary	5.11596337923e-05
rs780094	ENSG00000115216	NRBP1	Artery-Tibial	1.83676732572e-06
rs780094	ENSG00000115216	NRBP1	Cells-EBV-transformed-lymphocytes	1.46844818731e-08
rs780094	ENSG00000115216	NRBP1	Cells-Transformed-fibroblasts	4.22153030192e-09
rs780094	ENSG00000115216	NRBP1	Colon-Transverse	5.96150273596e-12
rs780094	ENSG00000115216	NRBP1	Esophagus-Mucosa	5.93300774324e-07
rs780094	ENSG00000115216	NRBP1	Esophagus-Muscularis	4.81306905877e-06
rs780094	ENSG00000115216	NRBP1	Lung	9.24479524465e-08
rs780094	ENSG00000115216	NRBP1	Nerve-Tibial	1.52854205395e-06
rs780094	ENSG00000115216	NRBP1	Skin-Not-Sun-Exposed-Suprapubic	8.45643939211e-14
rs780094	ENSG00000115216	NRBP1	Skin-Sun-Exposed-Lower-leg	5.69488821675e-16
rs780094	ENSG00000115216	NRBP1	Testis	3.61393652831e-07
rs780094	ENSG00000115216	NRBP1	Thyroid	4.76671891923e-06
rs780094	ENSG00000115216	NRBP1	Whole-Blood	1.00741937894e-11
rs780094	ENSG00000115226	FNDC4	Spleen	7.29173952836e-19
rs780094	ENSG00000115226	FNDC4	Thyroid	7.85233691869e-22
rs780094	ENSG00000115234	SNX17	Artery-Tibial	3.05445090118e-06
rs780094	ENSG00000115234	SNX17	Heart-Left-Ventricle	2.63905873533e-05
rs780094	ENSG00000115234	SNX17	Muscle-Skeletal	2.31809754253e-54
rs780094	ENSG00000115234	SNX17	Ovary	6.42403837308e-05
rs780094	ENSG00000115241	PPM1G	Adipose-Subcutaneous	1.64518363222e-07
rs780094	ENSG00000115241	PPM1G	Artery-Tibial	0.000127176868123
rs780094	ENSG00000115241	PPM1G	Esophagus-Muscularis	7.02182582115e-06
rs780094	ENSG00000115241	PPM1G	Heart-Atrial-Appendage	3.34090082833e-05
rs780094	ENSG00000115241	PPM1G	Skin-Sun-Exposed-Lower-leg	6.81873664966e-08
rs780094	ENSG00000115241	PPM1G	Thyroid	5.92296997261e-08
rs780094	ENSG00000115241	PPM1G	Uterus	6.2203422555e-05
rs780094	ENSG00000115241	PPM1G	Whole-Blood	5.08418221184e-06
rs780094	ENSG00000138002	IFT172	Cells-Transformed-fibroblasts	6.43149009709e-06
rs780094	ENSG00000138002	IFT172	Spleen	3.40766307931e-06
rs780094	ENSG00000138074	SLC5A6	Nerve-Tibial	2.59687895514e-07
rs780094	ENSG00000138085	ATRAID	Adipose-Subcutaneous	3.64204133511e-06
rs780094	ENSG00000138085	ATRAID	Cells-Transformed-fibroblasts	4.22656091471e-05
rs780094	ENSG00000138085	ATRAID	Esophagus-Mucosa	7.033199084e-05
rs780094	ENSG00000138085	ATRAID	Stomach	5.60184149184e-06

rs780094	ENSG00000157992	KRTCAP3	Adipose-Visceral-Omentum	1.19630711398e-05
rs780094	ENSG00000157992	KRTCAP3	Adrenal-Gland	1.05797393273e-09
rs780094	ENSG00000157992	KRTCAP3	Artery-Tibial	4.84310273246e-05
rs780094	ENSG00000157992	KRTCAP3	Esophagus-Mucosa	1.70988526206e-05
rs780094	ENSG00000157992	KRTCAP3	Muscle-Skeletal	3.46830497049e-09
rs780094	ENSG00000157992	KRTCAP3	Nerve-Tibial	7.76622325277e-05
rs780094	ENSG00000157992	KRTCAP3	Skin-Sun-Exposed-Lower-leg	2.65054404226e-07
rs780094	ENSG00000157992	KRTCAP3	Spleen	2.47712057827e-06
rs780094	ENSG00000157992	KRTCAP3	Testis	2.64461487969e-05
rs780094	ENSG00000157992	KRTCAP3	Thyroid	2.90732976901e-10
rs780094	ENSG00000157992	KRTCAP3	Whole-Blood	7.43521124449e-10
rs780094	ENSG00000163792	TCF23	Adipose-Visceral-Omentum	7.94940472883e-05
rs780094	ENSG00000163798	SLC4A1AP	Artery-Aorta	8.37212983625e-05
rs780094	ENSG00000163798	SLC4A1AP	Cells-Transformed-fibroblasts	5.15024966235e-05
rs780094	ENSG00000163798	SLC4A1AP	Esophagus-Mucosa	4.98374023788e-05
rs780094	ENSG00000171174	RBKS	Muscle-Skeletal	0.000105831011626
rs780094	ENSG00000176714	CCDC121	Spleen	5.68372729174e-05
rs780094	ENSG00000198522	GPN1	Muscle-Skeletal	3.37539746694e-05
rs780094	ENSG00000229122	AGBL5-IT1	Artery-Coronary	7.44343823279e-05
rs780094	ENSG00000234072	AC074117.10	Adipose-Subcutaneous	3.34633866994e-08
rs780094	ENSG00000234072	AC074117.10	Adipose-Visceral-Omentum	2.23628163175e-08
rs780094	ENSG00000234072	AC074117.10	Adrenal-Gland	1.84123088279e-06
rs780094	ENSG00000234072	AC074117.10	Artery-Aorta	1.73896132245e-05
rs780094	ENSG00000234072	AC074117.10	Artery-Coronary	2.77309460871e-07
rs780094	ENSG00000234072	AC074117.10	Artery-Tibial	1.01841800239e-08
rs780094	ENSG00000234072	AC074117.10	Breast-Mammary-Tissue	1.57199906534e-08
rs780094	ENSG00000234072	AC074117.10	Cells-Transformed-fibroblasts	1.85587761851e-22
rs780094	ENSG00000234072	AC074117.10	Colon-Sigmoid	6.10223022616e-08
rs780094	ENSG00000234072	AC074117.10	Colon-Transverse	4.29107590105e-11
rs780094	ENSG00000234072	AC074117.10	Esophagus-Gastroesophageal-Junction	3.63708645017e-06
rs780094	ENSG00000234072	AC074117.10	Esophagus-Mucosa	2.9963483337e-13
rs780094	ENSG00000234072	AC074117.10	Esophagus-Muscularis	3.66802466316e-10
rs780094	ENSG00000234072	AC074117.10	Heart-Atrial-Appendage	7.59712639918e-05
rs780094	ENSG00000234072	AC074117.10	Liver	6.4682462976e-05
rs780094	ENSG00000234072	AC074117.10	Lung	1.36528695798e-10
rs780094	ENSG00000234072	AC074117.10	Nerve-Tibial	9.65972368562e-06

rs780094	ENSG00000234072	AC074117.10	Pancreas	5.12889387587e-05
rs780094	ENSG00000234072	AC074117.10	Pituitary	9.13955202923e-12
rs780094	ENSG00000234072	AC074117.10	Prostate	1.99773648727e-06
rs780094	ENSG00000234072	AC074117.10	Stomach	3.51967258495e-06
rs780094	ENSG00000234072	AC074117.10	Testis	4.75994317682e-10
rs780094	ENSG00000234072	AC074117.10	Thyroid	6.08245717637e-20
rs780094	ENSG00000234072	AC074117.10	Vagina	3.58166998524e-06
rs780094	ENSG00000235267	AC074117.13	Breast-Mammary-Tissue	8.89875153068e-05
rs780094	ENSG00000235267	AC074117.13	Skin-Sun-Exposed-Lower-leg	3.74525571406e-05
rs780094	ENSG00000270210	RP11-373D23.3	Esophagus-Mucosa	4.63691181885e-05
rs780094	ENSG00000270210	RP11-373D23.3	Stomach	9.26771051729e-05
rs780094	ENSG00000270210	RP11-373D23.3	Thyroid	9.64224174215e-05
rs7944584	ENSG00000025434	NR1H3	Artery-Aorta	2.08009361483e-05
rs7944584	ENSG0000025434	NR1H3	Brain-Hypothalamus	1.22697126085e-07
rs7944584	ENSG0000025434	NR1H3	Brain-Nucleus-accumbens-basal-ganglia	0.000131733330946
rs7944584	ENSG0000025434	NR1H3	Cells-EBV-transformed-lymphocytes	2.29707271199e-10
rs7944584	ENSG0000025434	NR1H3	Cells-Transformed-fibroblasts	1.15434837914e-05
rs7944584	ENSG0000025434	NR1H3	Nerve-Tibial	7.20595223102e-05
rs7944584	ENSG0000030066	<i>NUP160</i>	Esophagus-Gastroesophageal-Junction	5.51250147279e-05
rs7944584	ENSG00000109919	MTCH2	Brain-Nucleus-accumbens-basal-ganglia	1.20137805688e-06
rs7944584	ENSG00000109919	MTCH2	Muscle-Skeletal	8.05066517251e-07
rs7944584	ENSG00000109919	MTCH2	Skin-Sun-Exposed-Lower-leg	8.53451748775e-05
rs7944584	ENSG00000109920	FNBP4	Cells-Transformed-fibroblasts	1.32732752938e-06
rs7944584	ENSG00000109920	FNBP4	Muscle-Skeletal	8.89603654962e-07
rs7944584	ENSG00000110492	MDK	Artery-Tibial	9.73202258852e-06
rs7944584	ENSG00000110514	MADD	Adipose-Subcutaneous	7.59998823807e-05
rs7944584	ENSG00000110514	MADD	Adipose-Visceral-Omentum	0.00012158221747
rs7944584	ENSG00000110514	MADD	Adrenal-Gland	2.2163994994e-05
rs7944584	ENSG00000110514	MADD	Artery-Tibial	8.90121801072e-08
rs7944584	ENSG00000110514	MADD	Breast-Mammary-Tissue	8.77616409944e-05
rs7944584	ENSG00000110514	MADD	Cells-Transformed-fibroblasts	4.94513410518e-09
rs7944584	ENSG00000110514	MADD	Colon-Transverse	6.3042408978e-06
rs7944584	ENSG00000110514	MADD	Esophagus-Mucosa	3.4527574667e-09
rs7944584	ENSG00000110514	MADD	Heart-Atrial-Appendage	3.17991729758e-08
rs7944584	ENSG00000110514	MADD	Heart-Left-Ventricle	2.457378901e-05
rs7944584	ENSG00000110514	MADD	Muscle-Skeletal	7.85528914203e-05

rs7944584	ENSG00000110514	MADD	Nerve-Tibial	3.26145755213e-05
rs7944584	ENSG00000110514	MADD	Pancreas	4.17482675702e-07
rs7944584	ENSG00000110514	MADD	Pituitary	1.72076652188e-11
rs7944584	ENSG00000110514	MADD	Thyroid	4.79922519912e-06
rs7944584	ENSG00000110514	MADD	Vagina	0.000122105725593
rs7944584	ENSG00000110514	MADD	Whole-Blood	2.16060782194e-05
rs7944584	ENSG00000123444	KBTBD4	Esophagus-Gastroesophageal-Junction	4.87252662217e-05
rs7944584	ENSG00000134569	LRP4	Nerve-Tibial	2.27800965124e-10
rs7944584	ENSG00000134569	LRP4	Pancreas	9.48225523427e-07
rs7944584	ENSG00000134571	MYBPC3	Heart-Atrial-Appendage	7.05925424226e-05
rs7944584	ENSG00000134571	MYBPC3	Lung	3.15496228313e-05
rs7944584	ENSG00000134571	MYBPC3	Nerve-Tibial	4.79897016865e-05
rs7944584	ENSG00000134571	MYBPC3	Whole-Blood	3.72029557362e-10
rs7944584	ENSG00000134574	DDB2	Esophagus-Mucosa	0.000125985422351
rs7944584	ENSG00000134575	ACP2	Adipose-Subcutaneous	7.68755973373e-06
rs7944584	ENSG00000134575	ACP2	Adipose-Visceral-Omentum	1.27104258893e-05
rs7944584	ENSG00000134575	ACP2	Cells-EBV-transformed-lymphocytes	1.07620921425e-08
rs7944584	ENSG00000134575	ACP2	Cells-Transformed-fibroblasts	0.000121806366566
rs7944584	ENSG00000134575	ACP2	Esophagus-Mucosa	2.73364543661e-14
rs7944584	ENSG00000134575	ACP2	Heart-Atrial-Appendage	1.20233194782e-06
rs7944584	ENSG00000134575	ACP2	Heart-Left-Ventricle	1.56687990124e-09
rs7944584	ENSG00000134575	ACP2	Lung	5.19595924101e-08
rs7944584	ENSG00000134575	ACP2	Muscle-Skeletal	0.00013144342915
rs7944584	ENSG00000134575	ACP2	Nerve-Tibial	2.61766719937e-06
rs7944584	ENSG00000134575	ACP2	Skin-Sun-Exposed-Lower-leg	1.07143020893e-17
rs7944584	ENSG00000134575	ACP2	Spleen	1.33254214111e-05
rs7944584	ENSG00000134575	ACP2	Thyroid	3.40882048091e-15
rs7944584	ENSG00000134575	ACP2	Whole-Blood	1.43425178272e-20
rs7944584	ENSG00000149179	C11 or f49	Artery-Coronary	5.67565971147e-05
rs7944584	ENSG00000149179	C11 or f49	Nerve-Tibial	0.00012048675964
rs7944584	ENSG00000149182	ARFGAP2	Artery-Tibial	6.29133475824e-05
rs7944584	ENSG00000149187	CELF1	Nerve-Tibial	2.50584748447e-05
rs7944584	ENSG00000165912	PACSIN3	Artery-Aorta	1.1604680458e-05
rs7944584	ENSG00000165912	PACSIN3	Esophagus-Mucosa	1.28324437184e-11
rs7944584	ENSG00000165912	PACSIN3	Skin-Sun-Exposed-Lower-leg	3.63199545474e-05
rs7944584	ENSG00000165912	PACSIN3	Spleen	6.6725849258e-07

rs7944584	ENSG00000165912	PACSIN3	Thyroid	7.9096234924e-07
rs7944584	ENSG00000165916	PSMC3	Adipose-Subcutaneous	2.34072925216e-05
rs7944584	ENSG00000165916	PSMC3	Cells-Transformed-fibroblasts	2.04305253074e-06
rs7944584	ENSG00000165916	PSMC3	Prostate	5.33810723515e-06
rs7944584	ENSG00000165916	PSMC3	Skin-Sun-Exposed-Lower-leg	3.11248061392e-06
rs7944584	ENSG00000165916	PSMC3	Thyroid	4.08398212744e-05
rs7944584	ENSG00000165916	PSMC3	Uterus	7.16490580163e-05
rs7944584	ENSG00000165917	RAPSN	Nerve-Tibial	8.62453385564e-05
rs7944584	ENSG00000165917	RAPSN	Skin-Sun-Exposed-Lower-leg	1.80631527423e-06
rs7944584	ENSG00000165917	RAPSN	Thyroid	1.20524036929e-07
rs7944584	ENSG00000172247	C1QTNF4	Adipose-Subcutaneous	3.56810013653e-34
rs7944584	ENSG00000172247	C1QTNF4	Artery-Aorta	1.24998255811e-09
rs7944584	ENSG00000172247	C1QTNF4	Artery-Coronary	1.33459962796e-05
rs7944584	ENSG00000172247	C1QTNF4	Artery-Tibial	4.45064032939e-11
rs7944584	ENSG00000172247	C1QTNF4	Breast-Mammary-Tissue	5.84525478966e-10
rs7944584	ENSG00000172247	C1QTNF4	Cells-EBV-transformed-lymphocytes	1.28865421506e-08
rs7944584	ENSG00000172247	C1QTNF4	Cells-Transformed-fibroblasts	5.38201805563e-10
rs7944584	ENSG00000172247	C1QTNF4	Colon-Transverse	2.04967618837e-10
rs7944584	ENSG00000172247	C1QTNF4	Esophagus-Mucosa	2.78374518782e-08
rs7944584	ENSG00000172247	C1QTNF4	Heart-Atrial-Appendage	1.13484278922e-08
rs7944584	ENSG00000172247	C1QTNF4	Liver	3.31423676529e-09
rs7944584	ENSG00000172247	C1QTNF4	Lung	3.85455422061e-22
rs7944584	ENSG00000172247	C1QTNF4	Muscle-Skeletal	1.42762821593e-17
rs7944584	ENSG00000172247	C1QTNF4	Nerve-Tibial	8.61270858773e-20
rs7944584	ENSG00000172247	C1QTNF4	Skin-Sun-Exposed-Lower-leg	1.04949880433e-28
rs7944584	ENSG00000172247	C1QTNF4	Spleen	4.24043916056e-15
rs7944584	ENSG00000172247	C1QTNF4	Thyroid	9.53527705852e-05
rs7944584	ENSG00000172247	C1QTNF4	Whole-Blood	5.13079512637e-10
rs7944584	ENSG00000247675	LRP4-AS1	Nerve-Tibial	9.25170624531e-09
rs7944584	ENSG00000247675	LRP4-AS1	Pancreas	5.93014122957e-05
rs7944584	ENSG00000271350	CTD-2384B9.1	Nerve-Tibial	2.14845905703e-05
rs7944584	ENSG00000271557	YPEL5P2	Nerve-Tibial	7.51699584443e-05

Table S3. Potential colocalization loci for FP phenotype. Colocation loci consists of two summary statistics. One summary statistics obtained FP GWAS and the other summary statistics obtained for one of the genes in GTEx. Thus, we obtained these potential colocalization loci by considering the variants which are reported significant in previous study [?] for FP phenotype in MAGIC dataset. For each significant variant, we consider genes for all the tissues in GTEx data where at least one of the variants in the locus are significantly associated (corrected multiple hypothesis p-value of 10^{-5} or more significant) with its gene expression. We consider all the 44 tissues in GTEx to obtain the list of genes for each significant GWAS SNP.

rsID	Ensembl ID	Gene Name	Tissue	eQTL p-values
rs10501320	ENSG0000025434	NR1H3	Artery-Aorta	2.08009361483e-05
rs10501320	ENSG0000025434	NR1H3	Brain-Hypothalamus	1.22697126085e-07
rs10501320	ENSG0000025434	NR1H3	Brain-Nucleus-accumbens-basal-ganglia	0.000131733330946
rs10501320	ENSG0000025434	NR1H3	Cells-Transformed-fibroblasts	1.85933794358e-05
rs10501320	ENSG0000025434	NR1H3	Nerve-Tibial	7.20595223102e-05
rs10501320	ENSG0000025434	NR1H3	Testis	2.37790021124e-07
rs10501320	ENSG0000030066	NUP160	Spleen	0.000116421914784
rs10501320	ENSG00000109919	MTCH2	Skin-Sun-Exposed-Lower-leg	8.53451748775e-05
rs10501320	ENSG00000109919	MTCH2	Testis	3.37267588819e-05
rs10501320	ENSG00000109920	FNBP4	Muscle-Skeletal	8.89603654962e-07
rs10501320	ENSG00000110492	MDK	Artery-Tibial	9.73202258852e-06
rs10501320	ENSG00000110514	MADD	Adipose-Subcutaneous	0.000112857498127
rs10501320	ENSG00000110514	MADD	Adrenal-Gland	2.2163994994e-05
rs10501320	ENSG00000110514	MADD	Artery-Aorta	0.000122351334124
rs10501320	ENSG00000110514	MADD	Artery-Tibial	8.90121801072e-08
rs10501320	ENSG00000110514	MADD	Cells-Transformed-fibroblasts	4.94513410518e-09
rs10501320	ENSG00000110514	MADD	Esophagus-Mucosa	3.4527574667e-09
rs10501320	ENSG00000110514	MADD	Heart-Atrial-Appendage	3.17991729758e-08
rs10501320	ENSG00000110514	MADD	Heart-Left-Ventricle	2.457378901e-05
rs10501320	ENSG00000110514	MADD	Muscle-Skeletal	7.85528914203e-05
rs10501320	ENSG00000110514	MADD	Nerve-Tibial	3.26145755213e-05
rs10501320	ENSG00000110514	MADD	Pancreas	4.17482675702e-07
rs10501320	ENSG00000110514	MADD	Pituitary	1.72076652188e-11
rs10501320	ENSG00000110514	MADD	Testis	2.16618984585e-07
rs10501320	ENSG00000110514	MADD	Vagina	0.000122105725593
rs10501320	ENSG00000134569	LRP4	Nerve-Tibial	6.19405876199e-31
rs10501320	ENSG00000134569	LRP4	Pancreas	4.0536511969e-16
rs10501320	ENSG00000134569	LRP4	Spleen	4.61451279373e-05
rs10501320	ENSG00000134571	MYBPC3	Heart-Atrial-Appendage	7.05925424226e-05

rs10501320	ENSG00000134571	MYBPC3	Nerve-Tibial	4.79897016865e-05	
		DDB2			
rs10501320	ENSG00000134574		Esophagus-Mucosa	0.000125985422351	
rs10501320	ENSG00000134574	DDB2	Heart-Atrial-Appendage	2.44184886821e-05	
rs10501320	ENSG00000134574	DDB2	Muscle-Skeletal	6.10776590083e-05	
rs10501320	ENSG00000134574	DDB2	Skin-Sun-Exposed-Lower-leg	2.43673646413e-10	
rs10501320	ENSG00000134575	ACP2	Adipose-Subcutaneous	7.68755973373e-06	
rs10501320	ENSG00000134575	ACP2	Esophagus-Mucosa	2.73364543661e-14	
rs10501320	ENSG00000134575	ACP2	Heart-Atrial-Appendage	8.70907835138e-07	
rs10501320	ENSG00000134575	ACP2	Heart-Left-Ventricle	1.56687990124e-09	
rs10501320	ENSG00000134575	ACP2	Lung	6.76675705027e-08	
rs10501320	ENSG00000134575	ACP2	Muscle-Skeletal	0.00013144342915	
rs10501320	ENSG00000134575	ACP2	Nerve-Tibial	2.61766719937e-06	
rs10501320	ENSG00000134575	ACP2	Skin-Not-Sun-Exposed-Suprapubic	2.55612655967e-06	
rs10501320	ENSG00000134575	ACP2	Skin-Sun-Exposed-Lower-leg	1.07143020893e-17	
rs10501320	ENSG00000134575	ACP2	Spleen	1.88165296359e-05	
rs10501320	ENSG00000134575	ACP2	Vagina	4.65323577258e-05	
rs10501320	ENSG00000149179	C11orf49	Adipose-Subcutaneous	5.67865942461e-05	
rs10501320	ENSG00000149179	C11orf49	Adrenal-Gland	1.24023663745e-05	
rs10501320	ENSG00000149179	C11orf49	Esophagus-Mucosa	8.92232838988e-05	
rs10501320	ENSG00000149179	C11orf49	Pancreas	3.3055837935e-05	
rs10501320	ENSG00000149179	C11orf49	Skin-Not-Sun-Exposed-Suprapubic	7.37619039844e-06	
rs10501320	ENSG00000149179	C11orf49	Skin-Sun-Exposed-Lower-leg	5.48944677851e-05	
rs10501320	ENSG00000149179	C11orf49	Testis	4.29749081422e-40	
rs10501320	ENSG00000149182	ARFGAP2	Artery-Tibial	6.29133475824e-05	
rs10501320	ENSG00000149187	CELF1	Prostate	1.61381245588e-05	
rs10501320	ENSG00000165912	PACSIN3	Adipose-Subcutaneous	2.28866958376e-08	
rs10501320	ENSG00000165912	PACSIN3	Artery-Aorta	1.1604680458e-05	
rs10501320	ENSG00000165912	PACSIN3	Artery-Tibial	7.62404655532e-05	
rs10501320	ENSG00000165912	PACSIN3	Cells-Transformed-fibroblasts	4.88531471913e-05	
rs10501320	ENSG00000165912	PACSIN3	Esophagus-Mucosa	1.41971075144e-16	
rs10501320	ENSG00000165912	PACSIN3	Lung	2.60370308594e-05	
rs10501320	ENSG00000165912	PACSIN3	Nerve-Tibial	9.24482319333e-06	
rs10501320	ENSG00000165912	PACSIN3	Ovary	2.39257187744e-05	
rs10501320	ENSG00000165912	PACSIN3	Skin-Not-Sun-Exposed-Suprapubic	3.11218769563e-07	
rs10501320	ENSG00000165912	PACSIN3	Skin-Sun-Exposed-Lower-leg	8.66082412293e-14	
rs10501320	ENSG00000165912	PACSIN3	Spleen	4.42663115843e-24	
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rs10501320	ENSG00000165916	PSMC3	Prostate	2.03094319084e-05
rs10501320	ENSG00000165916	PSMC3	Skin-Not-Sun-Exposed-Suprapubic	2.37308417946e-06
rs10501320	ENSG00000165916	PSMC3	Skin-Sun-Exposed-Lower-leg	7.04354880272e-05
rs10501320	ENSG00000165917	RAPSN	Skin-Sun-Exposed-Lower-leg	1.10308263023e-05
rs10501320	ENSG00000172247	C1QTNF4	Adipose-Subcutaneous	4.33882899494e-18
rs10501320	ENSG00000172247	C1QTNF4	Artery-Aorta	2.48213996867e-06
rs10501320	ENSG00000172247	C1QTNF4	Artery-Tibial	5.87064915435e-07
rs10501320	ENSG00000172247	C1QTNF4	Breast-Mammary-Tissue	3.75016221147e-06
rs10501320	ENSG00000172247	C1QTNF4	Cells-Transformed-fibroblasts	2.56325964789e-09
rs10501320	ENSG00000172247	C1QTNF4	Esophagus-Mucosa	2.49274783481e-00
rs10501320	ENSG00000172247	C1QTNF4	Liver	2.1273359988e-06
rs10501320	ENSG00000172247	C1QTNF4	Lung	3.49787806374e-14
rs10501320	ENSG00000172247	C1QTNF4	Muscle-Skeletal	8.54914979219e-1
rs10501320	ENSG00000172247	C1QTNF4	Nerve-Tibial	6.28029313008e-1
rs10501320	ENSG00000172247	C1QTNF4	Skin-Not-Sun-Exposed-Suprapubic	4.37740702424e-0
rs10501320	ENSG00000172247	C1QTNF4	Skin-Sun-Exposed-Lower-leg	3.28530648014e-2
rs10501320	ENSG00000172247	C1QTNF4	Spleen	5.88313478203e-1
rs10501320	ENSG00000175213	ZNF408	Prostate	0.00013159741462
rs10501320	ENSG00000175216	CKAP5	Esophagus-Mucosa	2.67017759846e-0
rs10501320	ENSG00000175220	ARHGAP1	Adipose-Subcutaneous	2.70869806044e-0
rs10501320	ENSG00000175220	ARHGAP1	Esophagus-Mucosa	9.82992146555e-0
rs10501320	ENSG00000175220	ARHGAP1	Muscle-Skeletal	4.12167429416e-0
rs10501320	ENSG00000175220	ARHGAP1	Nerve-Tibial	2.56649373374e-0
rs10501320	ENSG00000175220	ARHGAP1	Pancreas	2.00284121539e-0
rs10501320	ENSG00000175220	ARHGAP1	Skin-Sun-Exposed-Lower-leg	1.21286296661e-0
rs10501320	ENSG00000180210	F2	Esophagus-Gastroesophageal-Junction	2.61053625474e-0
rs10501320	ENSG00000247675	LRP4-AS1	Nerve-Tibial	3.43522823913e-2
rs10501320	ENSG00000247675	LRP4- $AS1$	Pancreas	1.183642192e-09
rs10501320	ENSG00000255520	RP11-390K5.3	Testis	1.77294876648e-0
rs10501320	ENSG00000271350	CTD-2384B9.1	Nerve-Tibial	1.46490579413e-1
rs10501320	ENSG00000271350	CTD-2384B9.1	Testis	1.85993958717e-4
rs10501320	ENSG00000271557	YPEL5P2	Nerve-Tibial	7.51699584443e-0
rs10838687	ENSG00000025434	NR1H3	Brain-Hypothalamus	1.22697126085e-0
rs10838687	ENSG00000025434	NR1H3	Brain-Nucleus-accumbens-basal-ganglia	0.00013173333094
rs10838687	ENSG00000025434	NR1H3	Nerve-Tibial	7.20595223102e-0
rs10838687	ENSG0000025434	NR1H3	Testis	2.37790021124e-0

rs10838687	ENSG0000030066	NUP160	Spleen	0.000116421914784
rs10838687	ENSG00000109919	MTCH2	Brain-Nucleus-accumbens-basal-ganglia	0.000120225415365
rs10838687	ENSG00000109919	MTCH2	Esophagus-Muscularis	4.1460701516e-05
rs10838687	ENSG00000109919	MTCH2	Testis	1.99678238221e-05
rs10838687	ENSG00000109920	FNBP4	Muscle-Skeletal	8.89603654962e-07
rs10838687	ENSG00000110492	MDK	Artery-Tibial	9.73202258852e-06
rs10838687	ENSG00000110514	MADD	Adipose-Subcutaneous	7.59998823807e-05
rs10838687	ENSG00000110514	MADD	Artery-Tibial	8.90121801072e-08
rs10838687	ENSG00000110514	MADD	Colon-Transverse	6.3042408978e-06
rs10838687	ENSG00000110514	MADD	Esophagus-Muscularis	1.15946798711e-10
rs10838687	ENSG00000110514	MADD	Heart-Left-Ventricle	2.457378901e-05
rs10838687	ENSG00000110514	MADD	Muscle-Skeletal	7.85528914203e-05
rs10838687	ENSG00000110514	MADD	Nerve-Tibial	3.26145755213e-05
rs10838687	ENSG00000110514	MADD	Pancreas	4.17482675702e-07
rs10838687	ENSG00000110514	MADD	Pituitary	1.72076652188e-11
rs10838687	ENSG00000110514	MADD	Testis	2.16618984585e-07
rs10838687	ENSG00000110514	MADD	Vagina	0.000122105725593
rs10838687	ENSG00000110514	MADD	Whole-Blood	2.16060782194e-05
rs10838687	ENSG00000123444	KBTBD4	Esophagus-Gastroesophageal-Junction	4.87252662217e-05
rs10838687	ENSG00000134569	LRP4	Nerve-Tibial	5.19553518572e-14
rs10838687	ENSG00000134569	LRP4	Pancreas	4.37416789409e-09
rs10838687	ENSG00000134571	MYBPC3	Nerve-Tibial	4.79897016865e-05
rs10838687	ENSG00000134571	MYBPC3	Whole-Blood	5.91404606882e-08
rs10838687	ENSG00000134574	DDB2	Muscle-Skeletal	9.04121978266e-05
rs10838687	ENSG00000134575	ACP2	Adipose-Subcutaneous	7.68755973373e-06
rs10838687	ENSG00000134575	ACP2	Adipose-Visceral-Omentum	8.70060974368e-06
rs10838687	ENSG00000134575	ACP2	Heart-Left-Ventricle	1.56687990124e-09
rs10838687	ENSG00000134575	ACP2	Muscle-Skeletal	0.00013144342915
rs10838687	ENSG00000134575	ACP2	Nerve-Tibial	2.61766719937e-06
rs10838687	ENSG00000134575	ACP2	Skin-Not-Sun-Exposed-Suprapubic	2.55612655967e-06
rs10838687	ENSG00000134575	ACP2	Spleen	1.33254214111e-05
rs10838687	ENSG00000134575	ACP2	Vagina	4.65323577258e-05
rs10838687	ENSG00000134575	ACP2	Whole-Blood	1.43425178272e-20
rs10838687	ENSG00000149179	C11orf49	Artery-Coronary	5.67565971147e-05
rs10838687	ENSG00000149179	C11 orf 49	Nerve-Tibial	0.00012048675964
rs10838687	ENSG00000149179	C11orf49	Testis	8.65627246025e-08

rs10838687	ENSG00000149182	ARFGAP2	Artery-Tibial	6.29133475824e-05
rs10838687	ENSG00000149187	CELF1	Prostate	1.61381245588e-05
rs10838687	ENSG00000165912	PACSIN3	Skin-Not-Sun-Exposed-Suprapubic	3.11218769563e-07
rs10838687	ENSG00000165912	PACSIN3	Spleen	6.6725849258e-07
rs10838687	ENSG00000165915	SLC39A13	Esophagus-Muscularis	1.45413413333e-05
rs10838687	ENSG00000165916	PSMC3	Adipose-Subcutaneous	2.34072925216e-05
rs10838687	ENSG00000165916	PSMC3	Prostate	7.24634191817e-06
rs10838687	ENSG00000165916	PSMC3	Skin-Not-Sun-Exposed-Suprapubic	2.37308417946e-06
rs10838687	ENSG00000165916	PSMC3	Uterus	7.16490580163e-05
rs10838687	ENSG00000165917	RAPSN	Nerve-Tibial	8.62453385564e-05
rs10838687	ENSG00000172247	C1QTNF4	Adipose-Subcutaneous	4.17124363169e-18
rs10838687	ENSG00000172247	C1QTNF4	Artery-Tibial	5.87064915435e-07
rs10838687	ENSG00000172247	C1QTNF4	Breast-Mammary-Tissue	2.92708043449e-06
rs10838687	ENSG00000172247	CIQTNF4	Colon-Transverse	8.43183897696e-07
rs10838687	ENSG00000172247	CIQTNF4	Esophagus-Muscularis	8.79210389775e-07
rs10838687	ENSG00000172247	CIQTNF4	Muscle-Skeletal	8.54914979219e-10
rs10838687	ENSG00000172247	C1QTNF4	Nerve-Tibial	6.28029313008e-15
rs10838687	ENSG00000172247	CIQINI4 C1QTNF4	Skin-Not-Sun-Exposed-Suprapubic	2.70064757209e-08
rs10838687	ENSG00000172247	CIQINI4 C1QTNF4	Spleen	5.88313478203e-10
rs10838687	ENSG00000172247	CIQTNF4	Whole-Blood	1.62499001406e-07
rs10838687	ENSG00000175220	ARHGAP1	Esophagus-Muscularis	4.79345806147e-05
rs10838687	ENSG00000175220	ARHGAP1	Nerve-Tibial	6.82296083501e-06
rs10838687	ENSG00000175220	ARHGAP1	Pancreas	3.72698860395e-05
rs10838687	ENSG00000180210	F2	Esophagus-Gastroesophageal-Junction	2.61053625474e-05
rs10838687	ENSG00000247675	LRP4-AS1	Nerve-Tibial	6.77398621043e-14
rs10838687	ENSG00000247675	LRP4-AS1	Pancreas	2.16806739278e-06
rs10838687	ENSG00000271350	CTD-2384B9.1	Nerve-Tibial	6.34706050202e-06
rs10838687	ENSG0000271350	CTD-2384B9.1	Testis	1.05240516036e-07
rs10838687	ENSG0000271557	YPEL5P2	Nerve-Tibial	7.51699584443e-05
rs11558471	ENSG00000147677	EIF3H	Spleen	3.9259893852e-05
rs11558471	ENSG00000164756	SLC30A8	Testis	2.05616636527e-05
rs11558471	ENSG00000205002	AARD	Uterus	2.36883921342e-05
rs11603334	ENSG00000110195	FOLR1	Testis	7.40531340652e-05
rs11603334	ENSG00000137522	RNF121	Adipose-Subcutaneous	0.00012619795151
rs11603334	ENSG00000168010	ATG16L2	Cells-Transformed-fibroblasts	0.000103804026962
rs11603334	ENSG0000168010	ATG16L2	Pituitary	9.57220796957e-09

rs11603334	ENSG00000186635	ARAP1	Brain-Hippocampus	2.1792203294e-07	
rs11603334	ENSG00000186635	ARAP1	Esophagus-Mucosa	2.05283456439e-31	
rs11603334	ENSG00000186635	ARAP1	Heart-Left-Ventricle	9.45571405972e-05	
rs11603334	ENSG00000186635	ARAP1	Pituitary	3.85918791654e-05	
rs11603334	ENSG00000186635	ARAP1	Skin-Not-Sun-Exposed-Suprapubic	4.26194524109e-15	
rs11603334	ENSG00000186635	ARAP1	Skin-Sun-Exposed-Lower-leg	8.6753935429e-13	
rs11603334	ENSG00000186635	ARAP1	Vagina	2.13540139623e-06	
rs11603334	ENSG00000186642	PDE2A	Pituitary	1.53022140653e-05	
rs11603334	ENSG00000214530	STARD10	Adipose-Subcutaneous	1.40157404858e-22	
rs11603334	ENSG00000214530	STARD10	Adipose-Visceral-Omentum	4.20018498505e-13	
rs11603334	ENSG00000214530	STARD10	Adrenal-Gland	1.68839708937e-05	
rs11603334	ENSG00000214530	STARD10	Artery-Coronary	3.46434620703e-07	
rs11603334	ENSG00000214530	STARD10	Artery-Tibial	1.80491817243e-19	
rs11603334	ENSG00000214530	STARD10	Brain-Hippocampus	6.91423944576e-13	
rs11603334	ENSG00000214530	STARD10	Brain-Hypothalamus	2.97445826171e-06	
rs11603334	ENSG00000214530	STARD10	Brain-Nucleus-accumbens-basal-ganglia	9.42264951173e-18	
rs11603334	ENSG00000214530	STARD10	Brain-Putamen-basal-ganglia	2.11944255236e-10	
rs11603334	ENSG00000214530	STARD10	Cells-Transformed-fibroblasts	1.10784037248e-22	
rs11603334	ENSG00000214530	STARD10	Colon-Sigmoid	9.66834209833e-05	
rs11603334	ENSG00000214530	STARD10	Esophagus-Gastroesophageal-Junction	6.01227455154e-09	
rs11603334	ENSG00000214530	STARD10	Esophagus-Mucosa	9.98256950799e-08	
rs11603334	ENSG00000214530	STARD10	Esophagus-Muscularis	3.63061000798e-21	
rs11603334	ENSG00000214530	STARD10	Heart-Atrial-Appendage	3.43706247956e-14	
rs11603334	ENSG00000214530	STARD10	Heart-Left-Ventricle	3.56243223152e-14	
rs11603334	ENSG00000214530	STARD10	Muscle-Skeletal	3.15486063502e-11	
rs11603334	ENSG00000214530	STARD10	Nerve-Tibial	8.025380066e-10	
rs11603334	ENSG00000214530	STARD10	Skin-Not-Sun-Exposed-Suprapubic	1.09794276756e-06	
rs11603334	ENSG00000214530	STARD10	Skin-Sun-Exposed-Lower-leg	1.77982862685e-12	
rs11603334	ENSG00000214530	STARD10	Spleen	6.24534239376e-06	
rs11603334	ENSG00000214530	STARD10	Testis	5.77727923768e-53	
rs11603334	ENSG00000214530	STARD10	Thyroid	1.68190571501e-21	
rs11603334	ENSG00000214530	STARD10	Whole-Blood	5.23070685799e-28	
rs11603334	ENSG00000244134	RPS12P20	Testis	3.07943415048e-05	
rs11603334	ENSG00000245148	ARAP1-AS2	Esophagus-Mucosa	3.35542315417e-09	
rs11603334	ENSG00000245148	ARAP1-AS2	Esophagus-Muscularis	0.000132342130158	
rs11603334	ENSG00000245148	ARAP1-AS2	Skin-Not-Sun-Exposed-Suprapubic	5.27971956062e-06	
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	rs11603334	ENSG00000245148	ARAP1-AS2	Skin-Sun-Exposed-Lower-leg	9.86566634234e-07
	rs11603334	ENSG00000256007	ARAP1-AS1	Esophagus-Mucosa	4.24608058891e-16
	rs11603334	ENSG00000256007	ARAP1-AS1	Skin-Not-Sun-Exposed-Suprapubic	2.9663658165e-10
	rs11603334	ENSG00000256007	ARAP1-AS1	Skin-Sun-Exposed-Lower-leg	5.59749377236e-09
	rs11603334	ENSG00000256007	ARAP1-AS1	Vagina	1.58350657457e-05
	rs11603334	ENSG00000256633	RP11-169D4.2	Pituitary	1.23716646726e-05
	rs1549318	ENSG00000137831	UACA	Adipose-Subcutaneous	7.62040217578e-05
	rs1549318	ENSG00000137831	UACA	Colon-Transverse	1.50865490222e-07
	rs1549318	ENSG00000137831	UACA	Esophagus-Muscularis	4.46725785788e-07
	rs1549318	ENSG00000137831	UACA	Heart-Left-Ventricle	5.12002585695e-06
	rs1549318	ENSG00000137831	UACA	Muscle-Skeletal	1.47756162906e-07
	rs1549318	ENSG00000137831	UACA	Nerve-Tibial	7.28332565726e-07
	rs1549318	ENSG00000166173	LARP6	Adipose-Subcutaneous	1.49001251151e-16
	rs1549318	ENSG00000166173	LARP6	Adipose-Visceral-Omentum	3.71861526378e-05
	rs1549318	ENSG00000166173	LARP6	Breast-Mammary-Tissue	3.01567496809e-10
	rs1549318	ENSG00000166173	LARP6	Cells-Transformed-fibroblasts	9.21978432466e-07
	rs1549318	ENSG00000235420	RPL29P30	Testis	1.91179363617e-17
	rs4502156	ENSG00000129003	VPS13C	Adipose-Subcutaneous	6.1378933828e-05
	rs4502156	ENSG00000129003	VPS13C	Adipose-Visceral-Omentum	0.000118069002431
	rs4502156	ENSG00000129003	VPS13C	Artery-Tibial	7.16531122802e-05
	rs4502156	ENSG00000129003	VPS13C	Breast-Mammary-Tissue	2.35245772991e-06
	rs4502156	ENSG00000129003	VPS13C	Colon-Transverse	5.8663572571e-05
	rs4502156	ENSG00000129003	VPS13C	Esophagus-Muscularis	9.70368440288e-06
	rs4502156	ENSG00000129003	VPS13C	Lung	1.04278737378e-05
	rs4502156	ENSG00000129003	VPS13C	Nerve-Tibial	5.62544360144e-13
	rs4502156	ENSG00000129003	VPS13C	Skin-Not-Sun-Exposed-Suprapubic	1.49754190539e-06
	rs4502156	ENSG00000129003	VPS13C	Spleen	4.11087561761e-06
	rs4502156	ENSG00000129003	VPS13C	Whole-Blood	8.48876714464e-14
	rs4502156	ENSG00000166104	RP11-299H22.3	Colon-Transverse	8.83954284179e-05
	rs4502156	ENSG00000198535	C2CD4A	Brain-Hippocampus	6.59057183308e-05
	rs4502156	ENSG00000198535	C2CD4A	Lung	9.4578914753e-13
	rs4502156	ENSG00000198535	C2CD4A	Nerve-Tibial	0.000108136855995
	rs4502156	ENSG00000259251	LOC101928907	Nerve-Tibial	9.94052704877e-05
	rs4502156	ENSG00000259498	RP11-244F12.3	Muscle-Skeletal	8.54339417378e-05
	rs4502156	ENSG00000259675	RP11-507B12.1	Vagina	3.22452434691e-05
	rs4502156	ENSG00000260062	GOLGA2P11	Pituitary	0.000125427812608
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rs4790333	ENSG0000070366	SMG6	Testis	2.01796250091e-06
rs4790333	ENSG0000070444	MNT	Adipose-Subcutaneous	4.53753966934e-08
rs4790333	ENSG0000070444	MNT	Adrenal-Gland	6.77383628198e-05
rs4790333	ENSG0000070444	MNT	Lung	2.08911326634e-08
rs4790333	ENSG0000070444	MNT	Nerve-Tibial	0.000123995238637
rs4790333	ENSG0000070444	MNT	Skin-Not-Sun-Exposed-Suprapubic	6.9348827878e-06
rs4790333	ENSG0000070444	MNT	Skin-Sun-Exposed-Lower-leg	1.41251332152e-10
rs4790333	ENSG0000070444	MNT	Testis	1.82199043457e-05
rs4790333	ENSG0000070444	MNT	Thyroid	2.8092498687e-05
rs4790333	ENSG0000070444	MNT	Whole-Blood	6.69149421706e-10
rs4790333	ENSG0000074660	SCARF1	Breast-Mammary-Tissue	5.81999912602e-05
rs4790333	ENSG00000108958	AC016292.3	Artery-Tibial	0.000120131601801
rs4790333	ENSG00000127804	METTL16	Muscle-Skeletal	3.64473881672e-11
rs4790333	ENSG00000132359	RAP1GAP2	Uterus	1.74458994676e-05
rs4790333	ENSG00000132376	INPP5K	Brain-Hypothalamus	0.000105311720954
rs4790333	ENSG00000132376	INPP5K	Brain-Putamen-basal-ganglia	9.25511693927e-05
rs4790333	ENSG00000132386	SERPINF1	Brain-Nucleus-accumbens-basal-ganglia	6.2230988248e-05
rs4790333	ENSG00000141258	SGSM2	Adipose-Subcutaneous	1.87167531203e-10
rs4790333	ENSG00000141258	SGSM2	Artery-Tibial	6.76272799246e-07
rs4790333	ENSG00000141258	SGSM2	Breast-Mammary-Tissue	0.000100471127705
rs4790333	ENSG00000141258	SGSM2	Esophagus-Mucosa	7.88667220572e-47
rs4790333	ENSG00000141258	SGSM2	Nerve-Tibial	2.57641700445e-11
rs4790333	ENSG00000141258	SGSM2	Pancreas	5.51821315507e-05
rs4790333	ENSG00000141258	SGSM2	Skin-Not-Sun-Exposed-Suprapubic	5.70294488579e-47
rs4790333	ENSG00000141258	SGSM2	Skin-Sun-Exposed-Lower-leg	5.39552112912e-75
rs4790333	ENSG00000141258	SGSM2	Thyroid	1.03236535251e-12
rs4790333	ENSG00000167720	SRR	Adipose-Subcutaneous	2.62770624435e-24
rs4790333	ENSG00000167720	SRR	Adipose-Visceral-Omentum	3.66599465005e-16
rs4790333	ENSG00000167720	SRR	Adrenal-Gland	2.18813555214e-21
rs4790333	ENSG00000167720	SRR	Artery-Aorta	1.5666233516e-27
rs4790333	ENSG00000167720	SRR	Artery-Tibial	3.51909532525e-27
rs4790333	ENSG00000167720	SRR	Brain-Hippocampus	1.96767876321e-06
rs4790333	ENSG00000167720	SRR	Breast-Mammary-Tissue	5.98991547683e-23
rs4790333	ENSG00000167720	SRR	Cells-Transformed-fibroblasts	1.96530464307e-58
rs4790333	ENSG00000167720	SRR	Colon-Sigmoid	4.711288685e-19
rs4790333	ENSG00000167720	SRR	Colon-Transverse	8.86943187099e-14

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rs4790333	ENSG00000167720	SRR	Esophagus-Gastroesophageal-Junction	6.39417074123e-20
rs4790333	ENSG00000167720	SRR	Esophagus-Mucosa	2.03191201428e-26
rs4790333	ENSG00000167720	SRR	Esophagus-Muscularis	4.63974738477e-35
rs4790333	ENSG00000167720	SRR	Heart-Atrial-Appendage	6.82859211342e-22
rs4790333	ENSG00000167720	SRR	Lung	2.00383443438e-27
rs4790333	ENSG00000167720	SRR	Muscle-Skeletal	1.1655201508e-40
rs4790333	ENSG00000167720	SRR	Nerve-Tibial	2.01827714245e-13
rs4790333	ENSG00000167720	SRR	Pancreas	4.61920620534e-11
rs4790333	ENSG00000167720	SRR	Prostate	1.0414967858e-05
rs4790333	ENSG00000167720	SRR	Skin-Not-Sun-Exposed-Suprapubic	1.55125329116e-08
rs4790333	ENSG00000167720	SRR	Skin-Sun-Exposed-Lower-leg	5.80578214063e-15
rs4790333	ENSG00000167720	SRR	${\it Small-Intestine-Terminal-Ileum}$	3.88146136704e-06
rs4790333	ENSG00000167720	SRR	Spleen	5.78311512656e-06
rs4790333	ENSG00000167720	SRR	Stomach	9.93468115975e-28
rs4790333	ENSG00000167720	SRR	Testis	5.68574604246e-10
rs4790333	ENSG00000167720	SRR	Thyroid	3.04088943829e-29
rs4790333	ENSG00000167720	SRR	Uterus	2.31653846137e-05
rs4790333	ENSG00000167720	SRR	Vagina	0.000128293539354
rs4790333	ENSG00000167720	SRR	Whole-Blood	2.63403824064e-12
rs4790333	ENSG00000167721	TSR1	Esophagus-Mucosa	3.19159108277e-07
rs4790333	ENSG00000167721	TSR1	Skin-Not-Sun-Exposed-Suprapubic	1.78661783865e-06
rs4790333	ENSG00000167721	TSR1	Skin-Sun-Exposed-Lower-leg	2.90773079263e-12
rs4790333	ENSG00000185924	RTN4RL1	Prostate	4.33826055896e-05
rs4790333	ENSG00000197879	MYO1C	Pituitary	4.94222032517e-05
rs4790333	ENSG00000236838	AC090617.1	Testis	4.05530839887e-11
rs4790333	ENSG00000239884	RN7SL608P	Adrenal-Gland	1.25431983335e-05
rs4790333	ENSG00000262333	HNRNPA1P16	Pancreas	4.41628579011e-05
rs4790333	ENSG00000262456	RP1-59D14.1	Adipose-Subcutaneous	1.9273076244e-05
rs4790333	ENSG00000262456	RP1-59D14.1	Lung	1.62943932639e-05
rs4790333	ENSG00000262456	RP1-59D14.1	Skin-Not-Sun-Exposed-Suprapubic	3.79534080211e-05
rs4790333	ENSG00000262456	RP1-59D14.1	Skin-Sun-Exposed-Lower-leg	3.03752617814e-09
rs4790333	ENSG00000262456	RP1-59D14.1	Whole-Blood	5.09383256186e-08
rs4790333	ENSG00000262533	RP11-667K14.4	Esophagus-Muscularis	3.34437203641e-07
rs4790333	ENSG00000262533	RP11-667K14.4	Thyroid	6.46727494394e-05
rs4790333	ENSG00000262884	LOC101927911	Adrenal-Gland	2.38261550886e-05
rs4790333	ENSG00000263345	RP1-59D14.5	Adipose-Subcutaneous	1.5906942067e-08

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rs4790333	B ENSG00000263345	RP1-59D14.5	Adipose-Visceral-Omentum	1.20925708546e-05
rs4790333	B ENSG00000263345	RP1-59D14.5	Artery-Aorta	2.64778354704e-06
rs4790333	B ENSG00000263345	RP1-59D14.5	Artery-Tibial	8.30712895499e-11
rs4790333	B ENSG00000263345	RP1-59D14.5	Breast-Mammary-Tissue	3.8474954638e-06
rs4790333	B ENSG0000263345	RP1-59D14.5	Cells-Transformed-fibroblasts	0.000103815497139
rs4790333	B ENSG00000263345	RP1-59D14.5	Colon-Sigmoid	1.0110430983e-07
rs4790333	B ENSG00000263345	RP1-59D14.5	Colon-Transverse	6.59603168599e-07
rs4790333	B ENSG00000263345	RP1-59D14.5	Esophagus-Gastroesophageal-Junction	1.17031329814e-05
rs4790333	B ENSG00000263345	RP1-59D14.5	Esophagus-Mucosa	4.69921628394e-17
rs4790333	B ENSG00000263345	RP1-59D14.5	Esophagus-Muscularis	2.28617562809e-08
rs4790333	B ENSG00000263345	RP1-59D14.5	Lung	1.14149548548e-08
rs4790333	B ENSG00000263345	RP1-59D14.5	Muscle-Skeletal	8.41241468254e-05
rs4790333	B ENSG00000263345	RP1-59D14.5	Nerve-Tibial	2.09575325565e-09
rs4790333	B ENSG00000263345	RP1-59D14.5	Pancreas	1.91973455846e-07
rs4790333	B ENSG00000263345	RP1-59D14.5	Skin-Not-Sun-Exposed-Suprapubic	6.38363784966e-20
rs4790333	B ENSG00000263345	RP1-59D14.5	Skin-Sun-Exposed-Lower-leg	2.24221376177e-41
rs4790333	B ENSG00000263345	RP1-59D14.5	Small-Intestine-Terminal-Ileum	4.41741438412e-05
rs4790333	B ENSG00000263345	RP1-59D14.5	Stomach	2.88769867873e-06
rs4790333	B ENSG00000263345	RP1-59D14.5	Thyroid	5.00865925951e-17
rs4790333	B ENSG00000272911	RP11-74E22.6	Thyroid	2.08968750769e-05
rs6235	ENSG00000164292	RHOBTB3	Muscle-Skeletal	7.32402521053e-07
rs6235	ENSG00000164308	ERAP2	Lung	6.94383289294e-06
rs6235	ENSG00000164308	ERAP2	Muscle-Skeletal	7.08184188823e-06
rs6235	ENSG00000175426	PCSK1	Adipose-Subcutaneous	1.86595194529e-05
rs6235	ENSG00000175426	PCSK1	Adipose-Visceral-Omentum	2.25873475953e-05
rs6235	ENSG00000175426	PCSK1	Artery-Aorta	7.48841603785e-05
rs6235	ENSG00000175426	PCSK1	Artery-Tibial	7.7780236049e-09
rs6235	ENSG00000175426	PCSK1	Muscle-Skeletal	1.99497206617e-05
rs6235	ENSG00000175426	PCSK1	Nerve-Tibial	2.47059964991e-05
rs6235	ENSG00000249746	RP11-254I22.3	Liver	1.15708047575e-05
rs6235	ENSG00000251000	AC008592.3	Small-Intestine-Terminal-Ileum	5.59609886091e-05
rs7903146	5 ENSG00000119927	GPAM	Uterus	0.000105881315055
rs7903146	5 ENSG00000148737	TCF7L2	Artery-Aorta	2.41309804086e-08
rs7903146	5 ENSG00000148737	TCF7L2	Cells-Transformed-fibroblasts	4.9364884775e-05
rs7903146	5 ENSG00000151532	VTI1A	Nerve-Tibial	0.000126683454415
rs7903146	5 ENSG00000165806	CASP7	Pancreas	0.000106676244722

rs7903146	ENSG00000165806	CASP7	Uterus	1.928007398e-05
rs7903146	ENSG00000197142	ACSL5	Muscle-Skeletal	8.30564397475e-05
rs7903146	ENSG00000197893	NRAP	Pancreas	5.87163857808e-05
rs7903146	ENSG00000233547	RP11-57H14.2	Adipose-Subcutaneous	8.01903050249e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Adipose-Visceral-Omentum	0.000116670281781
rs7903146	ENSG00000233547	RP11-57H14.2	Artery-Coronary	7.52881713657e-05
rs7903146	ENSG00000233547	RP11-57H14.2	Artery-Tibial	8.49093176404e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Breast-Mammary-Tissue	1.57494777667e-07
rs7903146	ENSG00000233547	RP11-57H14.2	Cells-Transformed-fibroblasts	2.20760403394e-05
rs7903146	ENSG00000233547	RP11-57H14.2	Colon-Sigmoid	1.70252542602e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Colon-Transverse	1.4331889962e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Esophagus-Gastroesophageal-Junction	3.46354862923e-08
rs7903146	ENSG00000233547	RP11-57H14.2	Esophagus-Mucosa	2.87613838631e-07
rs7903146	ENSG00000233547	RP11-57H14.2	Esophagus-Muscularis	1.21308084669e-14
rs7903146	ENSG00000233547	RP11-57H14.2	Lung	4.8139851224e-13
rs7903146	ENSG00000233547	RP11-57H14.2	Muscle-Skeletal	3.3063719769e-09
rs7903146	ENSG00000233547	RP11-57H14.2	Nerve-Tibial	7.92423442978e-33
rs7903146	ENSG00000233547	RP11-57H14.2	Ovary	6.58528159444e-07
rs7903146	ENSG00000233547	RP11-57H14.2	Pituitary	4.84808780686e-10
rs7903146	ENSG00000233547	RP11-57H14.2	Prostate	3.55202963832e-10
rs7903146	ENSG00000233547	RP11-57H14.2	${\it Skin-Not-Sun-Exposed-Suprapubic}$	3.2118971947e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Skin-Sun-Exposed-Lower-leg	8.31064590167e-07
rs7903146	ENSG00000233547	RP11-57H14.2	Testis	1.00946591504e-05
rs7903146	ENSG00000233547	RP11-57H14.2	Thyroid	4.80328361344e-11
rs7903146	ENSG00000233547	RP11-57H14.2	Uterus	2.18871343943e-05
rs7903146	ENSG00000233547	RP11-57H14.2	Vagina	7.31216360291e-06
rs7903146	ENSG00000233547	RP11-57H14.2	Whole-Blood	3.19447612908e-10
rs9727115	ENSG00000117598	PLPPR5	Testis	1.37817618778e-05
rs9727115	ENSG00000117600	PLPPR4	Cells-Transformed-fibroblasts	2.47272652341e-05
rs9727115	ENSG00000162627	SNX7	Adipose-Subcutaneous	3.10423023004e-06
rs9727115	ENSG00000162627	SNX7	Adipose-Visceral-Omentum	0.000119457641076
rs9727115	ENSG00000162627	SNX7	Artery-Aorta	3.27419942829e-14
rs9727115	ENSG00000162627	SNX7	Artery-Tibial	2.8535505536e-06
rs9727115	ENSG00000162627	SNX7	Cells-Transformed-fibroblasts	5.90947269828e-11
rs9727115	ENSG00000162627	SNX7	Esophagus-Muscularis	9.66559107291e-06
rs9727115	ENSG00000162627	SNX7	Heart-Left-Ventricle	3.72055920472e-05

rs9727115	ENSG00000162627	SNX7	Lung	3.60279212263e-10
rs9727115	ENSG00000162627	SNX7	Nerve-Tibial	7.07821490358e-10
rs9727115	ENSG00000162627	SNX7	Skin-Not-Sun-Exposed-Suprapubic	4.63114014753e-07
rs9727115	ENSG00000162627	SNX7	Skin-Sun-Exposed-Lower-leg	2.9230712805e-07
rs9727115	ENSG00000162627	SNX7	Spleen	7.04657413939e-16
rs9727115	ENSG00000162627	SNX7	Thyroid	1.1784762688e-08
rs9727115	ENSG00000162627	SNX7	Whole-Blood	1.35173975707e-15