

150521975	ADAMTSL4	0	0%	0%	0%	0%	0%	Genome Browser
chr3:186560463-186560538	ADIPOQ	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:134204162-134204574	ANAPC13,CEP63	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:41517959-41518026	ANK1,MIR486	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:47157986-47157990	ANXA8,ANXA8L1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:44656967-44657088	ARL17A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:131307601-131308779	ASAP1-IT1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:149570057-149570312	ATP6V0E2,ATP6V0E2-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:77166194-77166332	ATP7A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:35838320-35838404	ATPBD4-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:19281034-19281414	B9D1,MAPK7	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:41869863-41869870	B9D2,TMEM91	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:11913694-11922689	BCAR4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:103457657-103457740	BVM	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:105766783-105767329	BRF1,PACS2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:115933774-115934364	C10orf118,MIR2110	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:11865397-11865469	C10orf47	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:89787171-89787392	C16orf7,ZNF276	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:209957723-209957890	C1orf74	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:77020251-77020576	C1QTNF1,LOC100507410	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:20008631-20008769	C22orf25	0	0%	0%	0%	0%	0%	Show in Genome Browser

chr22:24856421-24890783	C22orf45	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:74020029-74020088	C6orf147	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:1126443-1126880	C7orf50,GPER	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:1062569-1062662	C7orf50,MIR339	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:104145192-104145597	C8orf56	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:144788865-144789739	CCDC166	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:43282776-43282860	CCDC23,ERMAP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:81040103-81040338	CENPN,CMC2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:123651344-123651790	CETN4P	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:73587802-73587890	CHCHD8,PAAF1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:19207989-19257925	CHODL-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:136553399-136553739	CHRM2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:103790135-103790344	CISD2	0	0.47%	0%	0%	0%	0%	Show in Genome Browser
chr17:36887013-36887048	CISD3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:92822810-92824778	CLLU1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:46691041-46692557	CN5H6.4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:15206869-15247466	COL6A4P1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:57481254-57481369	COQ9,CIAPIN1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:244998639-244999058	COX20	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:83316607-83316728	CPEB1,LOC283692	0	0%	0%	0%	0%	0%	Show in Genome Browser

chr2:211482295-211484597	CPS1,CPS1-IT1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:70933103-70945467	CSN1S2AP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:71001765-71012421	CSN1S2BP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:48483867-48495951	CTXN2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:53578621-53578694	DANCR	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:20911526-20911556	DCUN1D3,LYRM1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:20067834-20067906	DGCR8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:95624025-95624195	DICER1-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:31640783-31641309	DKFZp566F0947	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:10829080-10829179	DNM2,MIR638	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:111682249-111682383	DRAM2,CEPT1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:8086641-8086766	EEF1E1,EEF1E1-MUTED,SCARNA27	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:107461817-107462135	ELMOD1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:62379908-62380212	EML3,ROM1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:143105817-143105985	EPHA1-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:65222384-65222634	ERBB2IP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:54666153-54666834	ESRG,CACNA2D3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:38869437-38869455	FAM114A1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:89630940-89644466	FAM13A-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:41994576-41995355	FAM215A	0	0%	0%	0%	0%	0%	Show in Genome Browser

chr10:120863611-120863628	FAM45B,FAM45A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:11979496-11979559	FAM66D	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:98795434-98795746	FARP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:11714376-11714431	FBXO44,FBXO2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:175204561-175204827	FBXO8,CEP44	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:100003674-100004281	FKSG29	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:38665790-38665796	FLJ13197,KLF3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:2535656-2539552	FLJ35024	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:130628921-130793562	FLJ43663	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:90042560-90043818	FOXN3-AS2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:69048010-69051601	FTLP10	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:117748014-117748145	FXYD6	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:49207116-49207134	GAGE2A,GAGE2C,GAGE2D,GAGE8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:49197563-49197581	GAGE2D,GAGE8,GAGE2A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:45670583-45670980	GATM,LOC145663	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:247687982-247689235	GCSAML-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:52739463-52739856	GLT8D1,SPCS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:31633657-31633985	GPANK1,CSNK2B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:241375115-241375501	GPC1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:25447001-25448832	GPR158-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr5:68856051-68856073	GTF2H2C,GTF2H2D	0	0%	0%	0%	0%	0%	Genome Browser
chr20:61732644-61733466	HAR1B,HAR1A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:122654824-122654971	HAS2-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:23426125-23426351	HAUS4,MIR4707	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:152760411-152760571	HAUS7,BGN	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:30282046-30282259	HCG18,HCG17	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:30294131-30294620	HCG18,TRIM39	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:31439006-31440184	HCG26	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:80376453-80376616	HEXDC,C17orf101	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:54678664-54679028	HNRNPA1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:54268917-54270114	HPVC1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:133572745-133573726	HPYR1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:77724582-77726678	HSBP1L1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:113818551-113848339	HTR2C	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:125753509-125753912	HYLS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:144325474-144329226	HYMAI,PLAGL1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:153059701-153059967	IDH3G,SSR4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:10762539-10764548	ILF3-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:61874562-61874762	IPO11,LRRRC70	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:73164159-73164300	JPX	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:44526787-								Show in

44526886	KATNAL2	0	0%	0%	0%	0%	0%	Genome Browser
chr17:21279699-21280225	KCNJ12	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:2891263-2892094	KCNQ1DN	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:197476140-197476423	KIAA0226,FYTTD1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:127780320-127780535	KIAA0408,SOGA3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:136983261-136983338	KLHL3,MIR874	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:53342655-53342722	KRT8,KRT18	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:32201360-32202051	KRTAP7-1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:113558007-113558119	LARP7,C4orf21	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:156544096-156638483	LEKR1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:110721062-110721127	LIMS3-LOC440895,LOC100288570	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:43099462-43108628	LINC00111	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:40110879-40145401	LINC00114	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:761587-762902	LINC00115	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:36096105-36107053	LINC00160	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:29912375-29912676	LINC00161	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:46419129-46424642	LINC00162	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:102197694-102197805	LINC00239	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:102133333-102133769	LINC00263	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:61816009-61816260	LINC00305,LOC284294	0	0%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr21:35552978-35562220	LINC00310	0	0%	0%	0%	0%	0%	Genome Browser
chr3:8613468-8615580	LINC00312	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:22114914-22175426	LINC00320	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:133417649-133427710	LINC00326	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:75084725-75085031	LINC00338,SEC14L1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:21666675-22056919	LINC00340	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:95873606-95876427	LINC00341	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:72124149-72130448	LINC00472	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:166401017-166401038	LINC00473,LINC00602	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:18548073-18550201	LINC00493	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:83536046-83536234	LINC00575	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:29578778-29583939	LINC00589	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:219841006-219841644	LINC00608	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:107560511-107574076	LINC00635	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:18774309-18774626	LINC00652	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:28283908-28284140	LINC00662	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:72569012-72569025	LOC100093631,GTF2IP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:151814217-151814507	LOC100132111	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:70351169-70352448	LOC100133985	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:36802245-36802694	LOC100134317	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:25990435-	LOC100134868	0	0%	0%	0%	0%	0%	Show in Genome

25990525								Browser
chr10:93110514-93110636	LOC100188947	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:54495993-54496481	LOC100240734	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:26987146-26988085	LOC100270746,LINC00240	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:133452739-133454881	LOC100272217	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:149108877-149184371	LOC100272228	0	2.06%	0%	0%	0%	0%	Show in Genome Browser
chr6:46714699-46726941	LOC100287718	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:39735502-39735731	LOC100288846,CTAGE5	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:70233000-70233828	LOC100289511,SRSF5	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:106828643-106959485	LOC100302640	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:107218007-107220236	LOC100422737	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:8653778-8654077	LOC100506207,HULC	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:131148324-131156430	LOC100507203	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:43602791-43602992	LOC100652772,NNT	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:71954578-71956418	LOC145474	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:95995024-96051074	LOC145820	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:246952919-246953000	LOC149134	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:46452727-46452825	LOC150381	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:24141465-24143037	LOC152024	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:137696892-137699799	LOC158696	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:81805991-	LOC219347	0	0%	0%	0%	0%	0%	Show in Genome

81820689								Browser
chr10:10976904-10994126	LOC254312	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:53440810-53440847	LOC283335,TENC1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:56135944-56225006	LOC283856	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:74240612-74271783	LOC284276	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:31986547-31986664	LOC284551	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:170240547-170248073	LOC284688	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:26188822-26188914	LOC284801,MIR663A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:173007232-173010911	LOC285593	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:9641428-9765805	LOC285692	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:35699269-35703141	LOC285847	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:158493826-158493877	LOC340017	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:12623572-12668910	LOC340357	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:40058524-40058580	LOC344967,N4BP2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:84898604-84898920	LOC388152	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:37049239-37055146	LOC388796	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:65486868-65489185	LOC401463	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:89102316-89103330	LOC439994	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:89101999-89102167	LOC439994,LOC728190	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:89623367-89625125	LOC440173	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:4845379-	LOC440335	0	0%	0%	0%	0%	0%	Show in Genome

4845450								Browser
chr3:9439108-9439174	LOC440944	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:65183598-65183967	LOC643542,DSEL	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:783034-787490	LOC643837	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:56686237-56703430	LOC644145	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:128491328-128491415	LOC727677	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:100441812-100442114	LOC728012,MCHR2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:88998426-89096646	LOC728190	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:97515409-97515523	LOC728558,ENTPD1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:20004935-20005468	LOC729609	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:2323004-2330880	LOC730811,MYT1L	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:65225723-65226320	LOC84989	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:128226678-128229427	LOC90246	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:86247339-86250991	LOC90784	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:78285576-78286567	LOC91450	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:62646439-62780873	LOC92249	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:48989836-48998255	LPAR6	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:145751603-145752343	LRRC24,C8orf82	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:150022930-150023099	LRRC61	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:248902718-248903151	LYPD8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:43972846-43972879	MAPT	0	0%	0%	0%	0%	0%	Show in Genome

								Browser
chr17:43920723-43921527	MAPT-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:101361107-101365422	MEG8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:3477262-3477354	MEGF6,MIR551A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:133677408-133678029	MGC16121	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:1953202-1953302	MIR132	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:98511627-98511727	MIR137HG	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:1102484-1102578	MIR200B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:135061015-135061124	MIR202	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:568089-568198	MIR210HG	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:95054743-95054875	MIR3651,IARS,SNORA84	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:29729152-29729247	MIR3653,AP1B1,SNORD125	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:54291959-54292027	MIR373	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:219866367-219866430	MIR375	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:114058017-114058126	MIR448	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:110271155-110271212	MIR4497,TRPV4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:20513295-20513527	MIR4741,RBBP8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr21:29115712-29118351	MIR5009,LINC00113	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:11370451-11370544	MIR572	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:189547711-189547798	MIR944,TP63	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:52302295-52302377	MIRLET7G,WDR82	0	0%	0%	0%	0%	0%	Show in Genome Browser

chr1:1567560-1567753	MMP23B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:1593971-1594298	MOB2,LOC338651	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:33078645-33083532	N4BP2L2,N4BP2L2-IT2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:100395525-100395704	NCBP1,TSTD2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:4524555-4524718	NMRAL1,HMOX2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:65736786-65736915	NOL11,SNORA38B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:95087588-95087740	NOL8,CENPP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:79860072-79860402	NPB	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:132316094-132316167	NPHP3-ACAD11,CCRL1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:6052304-6052531	NPHP4,KCNAB2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:93771701-93771745	NUDT4,NUDT4P1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:56182375-56182590	NUPR1L	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:56702971-56718902	OACYLP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:41590239-41591794	OIP5-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:247693435-247693891	OR2C3,GCSAML-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:27294194-27294331	OST4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:163148164-163148311	PACRG,PARK2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:40532035-40532296	PAK6	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:112403072-112403146	PALM2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:31839357-31839509	PAX6,DKFZp686K1684	0	0%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr2:193614571-193641621	PCGEM1	0	0%	0%	0%	0%	0%	Genome Browser
chr15:77516252-77517746	PEAK1,LINC00597	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:9673900-9675447	PER4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:18118499-18118695	PET117	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:111927141-111931075	PGCP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:71320616-71357573	PIP5K1B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:15511279-15511464	PIR	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:16974582-16974783	PLCL2,MIR3714	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:6526129-6526151	PLEKHG5,TNFRSF25	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:72500192-72500309	PMS2L2,SPDYE8P	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:105627708-105627858	POPDC3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:75544420-75544497	POR	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:241395418-241395505	PP14571,GPC1,MIR149	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:10216965-10217070	PPAN,PPAN-P2RY11	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:45909467-45909549	PPP1R13L,CD3EAP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:42636619-42637486	PPP1R2P9	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:281385-282052	PPP2R3B-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:13359819-13359832	PRAMEF5,PRAMEF6	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:12976450-12976507	PRAMEF8	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:129872519-129872616	PRDM10,LINC00167	0	0%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr19:43853208-43853700	PRG1	0	0%	0%	0%	0%	0%	Genome Browser
chr7:766338-766846	PRKAR1B,HEATR2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:45072688-45073120	PRR5	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:50354416-50354756	PTOV1,PTOV1-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr18:8360818-8367032	PTPRM,LOC100192426	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:125773065-125773116	PUS3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:24811803-24825477	PWRN1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:117886850-117887105	RAD21-AS1,MIR3610	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr11:36616093-36619812	RAG2,C11orf74	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:17584787-17585107	RAI1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrY:24549623-24549729	RBM1YF,RBM1YJ	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:31926581-31926668	RDBP,SKIV2L	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:183605208-183605636	RGL1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:122231058-122231145	RHOF	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:35657750-35658014	RMRP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:97858799-97927544	RMST	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:51483892-51484276	RNASEH2B,RNASEH2B-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:42751397-42751866	RNF170,MIR4469	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:104097322-104097655	RNPC3,AMY2B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:43011251-43011399	RNU12	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:80529139-	RNU5E-1,CKMT2,RNU5D-1	0	0%	0%	0%	0%	0%	Show in Genome Browser

80529196								Browser
chr6:80451636-80451669	RNY4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:92764482-92764521	RPAP2,GLMN	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:17970727-17970730	RPL18A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:109459346-109541613	RPL34-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:20811230-20811570	RPPH1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:82824834-82824865	RPS17L	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:55459628-55459634	RPS27A,CLHC1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:200332821-200337479	SATB2,SATB2-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:50963901-50964034	SCO2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:72406599-72410422	SENP8,MYO9A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:10828608-10836877	SFTA1P	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:78194025-78194199	SGSH,SLC26A11	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:80457236-80457302	SH3BGRL,HMGN5	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:42396298-42396342	SLC20A2,C8orf40	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:53904044-53904721	SLC25A3P1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:107296961-107300651	SLC26A4-AS1	0	0.75%	0%	0%	0%	0%	Show in Genome Browser
chr7:107301080-107302083	SLC26A4,SLC26A4-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:27434888-27434898	SLC5A6,ATRAID	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:11034420-11034615	SLC6A1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:135465205-135465470	SMAD5-AS1	0	0%	0%	0%	0%	0%	Show in Genome

								Browser
chr5:70248837-70248841	SMN2,SMN1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:50643459-50643577	SNAR-D	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:47333842-47333961	SNAR-E	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:51108220-51108342	SNAR-F	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:190595719-190595839	SNAR-I	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:95999253-95999398	SNHG10	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:70563405-70563498	SNORD111B,SF3B3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:25509674-25509725	SNORD115-45	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:33952763-33952852	SNORD121A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:33934286-33934378	SNORD121B,UBAP2	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:9548939-9549026	SNORD123,LOC100505806	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr14:71865054-71865123	SNORD56B	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:24734861-24735071	SPATA13	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr13:36920362-36920646	SPG20,SPG20OS	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr10:135381692-135382876	SPRNP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:116592503-116593380	ST7-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:116594164-116594388	ST7-AS1,ST7-OT4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:116824179-116831815	ST7-OT3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:110848083-110848157	STARD4,STARD4-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:26232880-26232993	STMN1,MIR3917	0	0%	0%	0%	0%	0%	Show in Genome Browser

chr6:132929365-132930441	TAAR3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:9833900-9834226	TADA3,ARPC4	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr12:114845619-114846247	TBX5-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:153523383-153523438	TEX28	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:56685791-56685827	TGS1,TMEM68	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:149380037-149380168	TIGD6,HMGXB3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:3355433-3355643	TIGD7,ZNF75A	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr16:8619503-8622226	TMEM114	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:134832766-134833022	TMEM140	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:122685740-122685971	TMEM155,PP12613	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr3:99904668-99913030	TMEM30C	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:95582894-95583173	TMEM56	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr6:138188581-138188631	TNFAIP3,LOC100130476	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:151129105-151129198	TNFAIP8L2,TNFAIP8L2-SCNM1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr15:43802568-43802707	TP53BP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr17:7589389-7590757	TP53,WRAP53	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr4:165875600-165875920	TRIM61	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrY:22631758-22681114	TTY10	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrY:14774298-14804152	TTY15	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrY:7567398-7567528	TTY16	0	0%	0%	0%	0%	0%	Show in Genome Browser

chrY:8551411-8551473	TTY18	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrY:9167490-9172441	TTY20	0	0%	0%	0%	0%	0%	Show in Genome Browser
chrX:84189157-84189895	UBE2DNL	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:15945312-15946230	UCA1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr2:27531028-27531130	UCN	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr7:43965731-43965996	URGCP	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr9:79791679-79791822	VPS13A-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:140090861-140090958	VTRNA1-1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:140105744-140105831	VTRNA1-3	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:38273464-38273472	YRDC,C1orf122	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr22:32340479-32341336	YWHAH,C22orf24	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr5:76382623-76414346	ZBED3-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:77523115-77585268	ZFH4-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:62338794-62338911	ZGPAT,ARFRP1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:146228197-146228202	ZNF252P,ZNF252P-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:38085642-38085673	ZNF571,ZNF540	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr8:182129-182199	ZNF596,RPL23AP53	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr19:37329239-37329284	ZNF790	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr20:47894558-47894714	ZNFX1,ZNFX1-AS1	0	0%	0%	0%	0%	0%	Show in Genome Browser
chr1:2482431-2484284	LOC115110	0.01	1.34%	0%	0%	0%	0%	Show in Genome Browser
chr12:90102732-								Show in

90105728	LOC338758	0.01	1.9%	0%	0%	0%	0%	Genome Browser
chr10:81585658-81587355	LOC642361	0.01	1.06%	0%	0%	0%	0%	Show in Genome Browser
chrX:102942917-102943086	MORF4L2-AS1	0.01	1.76%	0%	0%	0%	0%	Show in Genome Browser
chr7:101928405-101928549	SH2B2	0.01	1.37%	0%	0%	0%	0%	Show in Genome Browser
chr6:94429881-94485057	TSG1	0.01	6.44%	0%	0%	0%	0%	Show in Genome Browser
chr8:63890420-63897459	UG0898H09,NKAIN3	0.01	1.27%	0%	0%	0%	0%	Show in Genome Browser
chr12:105724414-105765295	C12orf75	0.02	7.23%	0%	0%	0%	0%	Show in Genome Browser
chr7:135347221-135361159	C7orf73	0.02	6.25%	0%	0%	0%	0%	Show in Genome Browser
chr8:104152921-104153285	C8orf56,BAALC	0.02	2.73%	0%	0%	0%	0%	Show in Genome Browser
chr4:15683352-15692062	FAM200B	0.02	4.11%	0%	0%	0%	0%	Show in Genome Browser
chr13:111516334-111522655	LINC00346	0.02	2.84%	0%	0%	0%	0%	Show in Genome Browser
chr3:101395989-101398055	LOC100009676	0.02	2.27%	0%	0%	0%	0%	Show in Genome Browser
chr19:29456837-29460055	LOC148145	0.02	2.79%	0%	0%	0%	0%	Show in Genome Browser
chr10:80703084-80826343	LOC283050	0.02	8.71%	0%	0%	0%	0%	Show in Genome Browser
chr4:6675821-6676433	LOC93622	0.02	2.93%	0%	0%	0%	0%	Show in Genome Browser
chr6:134142285-134175130	MGC34034	0.02	5.11%	0%	0%	0%	0%	Show in Genome Browser
chr15:25068794-25165271	SNRPN	0.02	1.56%	0%	0%	0%	0%	Show in Genome Browser
chrY:8572513-8573324	TTY19	0.02	2.97%	0%	0%	0%	0%	Show in Genome Browser
chr3:55542336-55545304	ERC2	0.03	3.03%	0%	0%	0%	0%	Show in Genome Browser
chr5:127357246-127418766	FLJ33630	0.03	4.98%	0%	0%	0%	0%	Show in Genome Browser
chr6:31021984-								Show in

31027653	HCG22	0.03	3.17%	0%	0%	0%	0%	Genome Browser
chr16:19727778-19729479	IQCK,C16orf88	0.03	5.45%	0%	0%	0%	0%	Show in Genome Browser
chr18:39060238-39100561	KC6	0.03	5.86%	0%	0%	0%	0%	Show in Genome Browser
chr21:42513427-42519991	LINC00323	0.03	6.53%	0%	0%	0%	0%	Show in Genome Browser
chr6:166337536-166354254	LINC00473	0.03	4.69%	0%	0%	0%	0%	Show in Genome Browser
chr4:124573940-124851518	LOC285419	0.03	7.4%	0%	0%	0%	0%	Show in Genome Browser
chr1:78695283-78808324	MGC27382	0.03	1.7%	0%	0%	0%	0%	Show in Genome Browser
chr11:125703211-125709966	PATE4	0.03	7.77%	0%	0%	0%	0%	Show in Genome Browser
chr15:79575146-79590580	ANKRD34C	0.04	8.3%	0%	0%	0%	0%	Show in Genome Browser
chr16:71879899-71891231	ATXN1L	0.04	8.26%	0%	0%	0%	0%	Show in Genome Browser
chr18:5143672-5145754	C18orf42	0.04	4.32%	0%	0%	0%	0%	Show in Genome Browser
chr1:221503270-221509638	C1orf140	0.04	7.64%	0%	0%	0%	0%	Show in Genome Browser
chr9:115867006-115873957	FAM225B	0.04	9.47%	0%	0%	0%	0%	Show in Genome Browser
chr4:7099151-7105103	FLJ36777	0.04	8.41%	0%	0%	0%	0%	Show in Genome Browser
chr3:192959568-192961760	HRASLS,MGC2889	0.04	4.1%	0%	0%	0%	0%	Show in Genome Browser
chr13:91543208-91578851	LINC00410	0.04	11.03%	0%	0%	0%	0%	Show in Genome Browser
chr3:107642478-107647751	LINC00636	0.04	7.49%	0%	0%	0%	0%	Show in Genome Browser
chr19:19867183-19887222	LINC00663	0.04	7.96%	0%	0%	0%	0%	Show in Genome Browser
chr18:36786888-37115579	LINC00669	0.04	7.86%	0%	0%	0%	0%	Show in Genome Browser
chr3:122605360-122611262	LOC100129550	0.04	4.57%	0%	0%	0%	0%	Show in Genome Browser
chr5:180256959-								Show in

180258229	LOC100859930,LOC729678	0.04	4.48%	0%	0%	0%	0%	Genome Browser
chr15:69854059-69863779	LOC145837	0.04	2.84%	0%	0%	0%	0%	Show in Genome Browser
chr5:180258736-180262725	LOC729678	0.04	1.17%	0%	0%	0%	0%	Show in Genome Browser
chr2:109743784-109746575	SH3RF3,SH3RF3-AS1	0.04	7.79%	0%	0%	0%	0%	Show in Genome Browser
chr1:3652550-3656951	TP73,TP73-AS1	0.04	3.72%	0%	0%	0%	0%	Show in Genome Browser
chr12:72647287-72658401	TRHDE-AS1	0.04	4.14%	0%	0%	0%	0%	Show in Genome Browser
chr15:69096160-69099440	ANP32A-IT1	0.05	5.48%	0%	0%	0%	0%	Show in Genome Browser
chr3:138823027-138844003	BPESC1	0.05	5.12%	0%	0%	0%	0%	Show in Genome Browser
chr14:70036532-70037918	C14orf162	0.05	5.91%	0%	0%	0%	0%	Show in Genome Browser
chr5:138727635-138730885	C5orf65	0.05	5.53%	0%	0%	0%	0%	Show in Genome Browser
chr6:11094266-11138964	C6orf228	0.05	9.6%	0%	0%	0%	0%	Show in Genome Browser
chr2:218148748-218465552	DIRC3	0.05	9.46%	0%	0%	0%	0%	Show in Genome Browser
chr5:111755280-111756673	FLJ11235	0.05	5.38%	0%	0%	0%	0%	Show in Genome Browser
chr4:38614322-38637679	FLJ13197	0.05	11.22%	0%	0%	0%	0%	Show in Genome Browser
chr7:43152198-43153990	HECW1	0.05	5.92%	0%	0%	0%	0%	Show in Genome Browser
chr4:21844964-21854811	KCNIP4,KCNIP4-IT1	0.05	5.03%	0%	0%	0%	0%	Show in Genome Browser
chr16:432241-442960	LOC100134368	0.05	11.47%	0%	0%	0%	0%	Show in Genome Browser
chr10:4698348-4704604	LOC100216001,LOC338588	0.05	10.49%	0%	0%	0%	0%	Show in Genome Browser
chr3:94657107-94710168	LOC255025	0.05	8.41%	0%	0%	0%	0%	Show in Genome Browser
chr19:19292686-19293381	MEF2BNB,MEF2BNB-MEF2B	0.05	5.89%	0%	0%	0%	0%	Show in Genome Browser
chr8:146228286-146231431	ZNF252P-AS1	0.05	5.72%	0%	0%	0%	0%	Show in Genome

								Browser
chr2:38358247-38408991	CYP1B1-AS1	0.06	10.93%	0%	0%	0%	0%	Show in Genome Browser
chr1:29443331-29445939	EPB41,TMEM200B	0.06	6.28%	0%	0%	0%	0%	Show in Genome Browser
chr11:64889562-64889660	FAU,MRPL49	0.06	6.06%	0%	0%	0%	0%	Show in Genome Browser
chr7:141404138-141438030	FLJ40852	0.06	12.39%	0%	0%	0%	0%	Show in Genome Browser
chr15:50641137-50646370	GABPB1-AS1,FLJ10038	0.06	6.87%	0%	0%	0%	0%	Show in Genome Browser
chr12:54356098-54361178	HOTAIR	0.06	12.4%	0%	0%	0%	0%	Show in Genome Browser
chr7:1614351-1629259	KIAA1908	0.06	12.11%	0%	0%	0%	0%	Show in Genome Browser
chr20:22541194-22559280	LINC00261	0.06	9.69%	0%	0%	0%	0%	Show in Genome Browser
chr1:204337558-204338847	LINC00628	0.06	6.97%	0%	0%	0%	0%	Show in Genome Browser
chr20:34633542-34638882	LINC00657	0.06	6.74%	0%	0%	0%	0%	Show in Genome Browser
chr7:152161209-152162628	LOC100128822	0.06	6.33%	0%	0%	0%	0%	Show in Genome Browser
chr15:57178368-57210697	LOC145783	0.06	8.1%	0%	0%	0%	0%	Show in Genome Browser
chr15:90048162-90060668	LOC283761	0.06	9.36%	0%	0%	0%	0%	Show in Genome Browser
chr20:37842424-37853391	LOC339568	0.06	12.89%	0%	0%	0%	0%	Show in Genome Browser
chr12:8509562-8543348	LOC389634	0.06	9.26%	0%	0%	0%	0%	Show in Genome Browser
chr7:130126900-130129808	MEST,MESTIT1	0.06	4.7%	0%	0%	0%	0%	Show in Genome Browser
chr11:133766330-133771635	MIR4697,LOC283174	0.06	5.1%	0%	0%	0%	0%	Show in Genome Browser
chr11:57510987-57518708	TMX2-CTNND1,BTBD18	0.06	12.83%	0%	0%	0%	0%	Show in Genome Browser
chrX:26576454-26579168	VENTXP1	0.06	6.62%	0%	0%	0%	0%	Show in Genome Browser
chr8:77593515-77595510	ZFHX4,ZFHX4-AS1	0.06	7.61%	0%	0%	0%	0%	Show in Genome Browser

chr3:55691245-55693497	ERC2-IT1	0.07	7.1%	0%	0%	0%	0%	Show in Genome Browser
chr20:61885893-61892963	FLJ16779	0.07	7.55%	0%	0%	0%	0%	Show in Genome Browser
chr22:18518526-18520734	FLJ41941	0.07	7.33%	0%	0%	0%	0%	Show in Genome Browser
chr20:61726845-61727535	HAR1B	0.07	7.67%	0%	0%	0%	0%	Show in Genome Browser
chr12:121407641-121410095	HNF1A-AS1	0.07	7.33%	0%	0%	0%	0%	Show in Genome Browser
chr14:93533797-93538476	ITPK1,ITPK1-AS1	0.07	6.98%	0%	0%	0%	0%	Show in Genome Browser
chr13:44596471-44604598	LINC00284	0.07	14.6%	0%	0%	0%	0%	Show in Genome Browser
chr13:30914409-30948036	LINC00426	0.07	17.65%	0%	0%	0%	0%	Show in Genome Browser
chr12:94853779-94856342	LOC144486	0.07	7.02%	0%	0%	0%	0%	Show in Genome Browser
chr7:158801045-158818928	LOC154822	0.07	7.69%	0%	0%	0%	0%	Show in Genome Browser
chr10:12875133-12877544	LOC283070	0.07	6.42%	0%	0%	0%	0%	Show in Genome Browser
chr19:57321445-57323847	ZIM2,PEG3,PEG3-AS1	0.07	7.49%	0%	0%	0%	0%	Show in Genome Browser
chrX:1520355-1521118	ASMTL-AS1	0.08	7.98%	0%	0%	0%	0%	Show in Genome Browser
chr15:25361692-25367621	IPW	0.08	11.57%	0%	0%	0%	0%	Show in Genome Browser
chr10:1205708-1210612	LINC00200	0.08	12.13%	0%	0%	0%	0%	Show in Genome Browser
chr6:170200927-170202969	LINC00574	0.08	8.81%	0%	0%	0%	0%	Show in Genome Browser
chr10:79686570-79689582	LOC100128292	0.08	8.13%	0%	0%	0%	0%	Show in Genome Browser
chr9:123605320-123616644	LOC253039	0.08	11.1%	0%	0%	0%	0%	Show in Genome Browser
chr9:130890808-130892911	LOC389791	0.08	8.26%	0%	0%	0%	0%	Show in Genome Browser
chr17:43973149-43976163	MAPT-IT1	0.08	8.95%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr19:19281014-19288269	MEF2BNB-MEF2B	0.08	14.05%	0%	0%	0%	0%	Genome Browser
chr17:14207057-14209061	MGC12916,HS3ST3B1	0.08	8.67%	0%	0%	0%	0%	Show in Genome Browser
chr15:42491769-42491864	MIR627,VPS39	0.08	8.33%	0%	0%	0%	0%	Show in Genome Browser
chr11:125658006-125661493	PATE3	0.08	14.87%	0%	0%	0%	0%	Show in Genome Browser
chr2:233412779-233415226	TIGD1,MIR5001	0.08	8.74%	0%	0%	0%	0%	Show in Genome Browser
chr3:38492520-38494716	ACVR2B-AS1	0.09	8.05%	0%	0%	0%	0%	Show in Genome Browser
chr8:22497884-22499722	FLJ14107,BIN3	0.09	8.75%	0%	0%	0%	0%	Show in Genome Browser
chr4:80782444-80784399	GDEP	0.09	9.2%	0%	0%	0%	0%	Show in Genome Browser
chr5:150226085-150228229	IRGM	0.09	6.64%	0%	0%	0%	0%	Show in Genome Browser
chr2:33050510-33151756	LINC00486	0.09	3.28%	0%	0%	0%	0%	Show in Genome Browser
chr3:151980413-151985828	MBNL1,LOC401093	0.09	7.93%	0%	0%	0%	0%	Show in Genome Browser
chr5:179233388-179233486	MGAT4B,SQSTM1	0.09	9.09%	0%	0%	0%	0%	Show in Genome Browser
chr19:57352270-57359922	MIMT1	0.09	9.86%	0%	0%	0%	0%	Show in Genome Browser
chr6:137303297-137311497	NHEG1	0.09	13.84%	0%	0%	0%	0%	Show in Genome Browser
chr16:67977377-67977850	SLC12A4,LCAT	0.09	9.28%	0%	0%	0%	0%	Show in Genome Browser
chr9:115876700-115882124	FAM225A	0.1	10.5%	0%	0%	0%	0%	Show in Genome Browser
chr10:114067937-114116353	GUCY2GP	0.1	7.05%	0%	0%	0%	0%	Show in Genome Browser
chr2:74730444-74732177	LBX2-AS1,PCGF1	0.1	10.38%	0%	0%	0%	0%	Show in Genome Browser
chr14:56247854-56260174	LINC00520	0.1	15.82%	0%	0%	0%	0%	Show in Genome Browser
chr13:114451484-114454062	LINC00552	0.1	10.46%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr7:139877061-139879439	LOC100134229	0.1	9.2%	0%	0%	0%	0%	Genome Browser
chr15:93113433-93115493	LOC100144604	0.1	4.95%	0%	0%	0%	0%	Show in Genome Browser
chr8:125954250-125957988	LOC157381	0.1	12.33%	0%	0%	0%	0%	Show in Genome Browser
chr12:9800643-9806404	LOC374443	0.1	12.24%	0%	0%	0%	0%	Show in Genome Browser
chr5:135527157-135528851	LOC389332	0.1	8.14%	0%	0%	0%	0%	Show in Genome Browser
chr1:45769582-45771290	LOC400752	0.1	8.77%	0%	0%	0%	0%	Show in Genome Browser
chr1:165451586-165470864	LOC400794	0.1	8.54%	0%	0%	0%	0%	Show in Genome Browser
chr2:108439520-108442583	LOC729121	0.1	5.69%	0%	0%	0%	0%	Show in Genome Browser
chrX:151883119-151884565	MAGEA2	0.1	14.34%	0%	0%	0%	0%	Show in Genome Browser
chr4:174451609-174462980	NBLA00301	0.1	8.41%	0%	0%	0%	0%	Show in Genome Browser
chr9:79398621-79402465	PCA3,PRUNE2	0.1	17.03%	0%	0%	0%	0%	Show in Genome Browser
chr19:47778142-47778979	PRR24	0.1	10.73%	0%	0%	0%	0%	Show in Genome Browser
chr16:583946-586142	SOLH	0.1	6.54%	0%	0%	0%	0%	Show in Genome Browser
chr9:92254698-92334674	UNQ6494	0.1	7.93%	0%	0%	0%	0%	Show in Genome Browser
chr4:184018174-184020352	WWC2-AS2	0.1	10.6%	0%	0%	0%	0%	Show in Genome Browser
chr19:45041045-45060150	CEACAM22P	0.11	8.9%	0%	0%	0%	0%	Show in Genome Browser
chr10:124516210-124556266	FLJ46361	0.11	10.83%	0%	0%	0%	0%	Show in Genome Browser
chr17:46800542-46800675	HOXB-AS5	0.11	11.94%	0%	0%	0%	0%	Show in Genome Browser
chr9:98782016-98783563	LINC00092	0.11	11.62%	0%	0%	0%	0%	Show in Genome Browser
chr3:10801169-10805877	LINC00606	0.11	1.98%	0%	0%	0%	0%	Show in Genome Browser
chr13:28710981-	PAN3-AS1	0.11	11.52%	0%	0%	0%	0%	Show in Genome

28712542								Browser
chr6:41072849-41106464	ADCY10P1	0.12	13.34%	0%	0%	0%	0%	Show in Genome Browser
chr7:149564785-149568885	ATP6V0E2-AS1	0.12	13.55%	0%	0%	0%	0%	Show in Genome Browser
chr14:96505662-96560133	C14orf132	0.12	16.3%	0%	0%	0%	0%	Show in Genome Browser
chr14:65877313-65879335	FUT8,FUT8-AS1	0.12	12.6%	0%	0%	0%	0%	Show in Genome Browser
chr8:122651586-122652408	HAS2,HAS2-AS1	0.12	19.38%	0%	0%	0%	0%	Show in Genome Browser
chr6:135963331-136011975	LINC00271	0.12	25.87%	0%	0%	0%	0%	Show in Genome Browser
chr18:5243593-5246503	LINC00667	0.12	12.36%	0%	0%	0%	0%	Show in Genome Browser
chr1:151810339-151813033	LOC100132111,C2CD4D	0.12	9.25%	0%	0%	0%	0%	Show in Genome Browser
chr9:35909480-35911617	LOC158376	0.12	17.54%	0%	0%	0%	0%	Show in Genome Browser
chr1:24528954-24538180	LOC284632	0.12	20.37%	0%	0%	0%	0%	Show in Genome Browser
chr20:25121434-25129426	LOC284798	0.12	11.74%	0%	0%	0%	0%	Show in Genome Browser
chr6:143875467-143890476	LOC285740	0.12	18.66%	0%	0%	0%	0%	Show in Genome Browser
chr22:46402496-46406657	LOC730668	0.12	12.3%	0%	0%	0%	0%	Show in Genome Browser
chr15:40643234-40648632	PHGR1	0.12	2.47%	0%	0%	0%	0%	Show in Genome Browser
chr10:89618919-89623194	PTEN,KLLN	0.12	11.71%	0%	0%	0%	0%	Show in Genome Browser
chr2:128458597-128461405	SFT2D3	0.12	11.46%	0%	0%	0%	0%	Show in Genome Browser
chr20:584639-590910	TCF15	0.12	12.73%	0%	0%	0%	0%	Show in Genome Browser
chr17:1173858-1174564	BHLHA9	0.13	13.71%	0%	0%	0%	0%	Show in Genome Browser
chr16:4784289-4784699	C16orf71,ANKS3	0.13	9%	0%	0%	0%	0%	Show in Genome Browser
chrX:49106609-	CCDC22,FOXP3	0.13	13.14%	0%	0%	0%	0%	Show in Genome

49106897								Browser
chr10:8092413-8095447	FLJ45983	0.13	11.43%	0%	0%	0%	0%	Show in Genome Browser
chr17:6543957-6544221	KIAA0753,TXNDC17	0.13	13.96%	0%	0%	0%	0%	Show in Genome Browser
chr11:129872731-129875381	LINC00167	0.13	12.33%	0%	0%	0%	0%	Show in Genome Browser
chr18:5236724-5238028	LINC00526	0.13	13.79%	0%	0%	0%	0%	Show in Genome Browser
chr1:3816968-3832009	LOC100133612	0.13	18.36%	0%	0%	0%	0%	Show in Genome Browser
chr10:127262940-127267012	LOC100169752	0.13	8.2%	0%	0%	0%	0%	Show in Genome Browser
chr4:100871636-100873620	LOC256880	0.13	12.69%	0%	0%	0%	0%	Show in Genome Browser
chr11:121959811-121962448	MIR100HG	0.13	9.59%	0%	0%	0%	0%	Show in Genome Browser
chr4:85890878-85928168	WDFY3-AS2	0.13	23.26%	0%	0%	0%	0%	Show in Genome Browser
chr5:95192582-95195834	C5orf27	0.14	13.07%	0%	0%	0%	0%	Show in Genome Browser
chr1:226250421-226250427	H3F3AP4	0.14	14.28%	0%	0%	0%	0%	Show in Genome Browser
chr20:61733672-61735737	HAR1A	0.14	10.41%	0%	0%	0%	0%	Show in Genome Browser
chr6:26521934-26527610	HCG11	0.14	11.09%	0%	0%	0%	0%	Show in Genome Browser
chr10:70091768-70092684	HNRNP3,PBLD	0.14	11.7%	0%	0%	0%	0%	Show in Genome Browser
chr12:117299028-117319232	HRK	0.14	18.4%	0%	0%	0%	0%	Show in Genome Browser
chr8:1922044-1955108	KBTD11	0.14	20.93%	0%	0%	0%	0%	Show in Genome Browser
chr17:79885705-79888628	MAFG-AS1	0.14	17.37%	0%	0%	0%	0%	Show in Genome Browser
chr16:88729781-88741913	MGC23284	0.14	23.72%	0%	0%	0%	0%	Show in Genome Browser
chr15:25380789-25383200	PAR1	0.14	12.23%	0%	0%	0%	0%	Show in Genome Browser
chr19:5558180-								Show in

5568005	PLAC2	0.14	19.79%	0%	0%	0%	0%	Genome Browser
chr2:62423262-62423490	B3GNT2	0.15	15.28%	0%	0%	0%	0%	Show in Genome Browser
chr17:53038480-53039328	COX11,TOM1L1	0.15	10.88%	0%	0%	0%	0%	Show in Genome Browser
chr9:35860271-35865513	FP588	0.15	13.02%	0%	0%	0%	0%	Show in Genome Browser
chrX:148609134-148621312	LOC100131434	0.15	10.29%	0%	0%	0%	0%	Show in Genome Browser
chr7:104650989-104654588	LOC100216545	0.15	12.22%	0%	0%	0%	0%	Show in Genome Browser
chr15:24409926-24415053	PWRN2	0.15	14.89%	0%	0%	0%	0%	Show in Genome Browser
chr21:46490873-46493126	SSR4P1	0.15	13.66%	0%	0%	0%	0%	Show in Genome Browser
chr5:76372532-76373720	ZBED3	0.15	9.84%	0%	0%	0%	0%	Show in Genome Browser
chr12:56982924-56989379	BAZ2A,RBMS2	0.16	11.61%	0.43%	0%	0%	0%	Show in Genome Browser
chr8:7812535-7813080	FAM66E	0.16	16.48%	0%	0%	0%	0%	Show in Genome Browser
chr10:99625472-99627781	GOLGA7B,CRTAC1	0.16	10.73%	0%	0%	0%	0%	Show in Genome Browser
chr1:59597610-59612479	HSD52	0.16	4.33%	0%	0%	0%	0%	Show in Genome Browser
chr1:168758019-168762120	LINC00626	0.16	25.44%	0%	0%	0%	0%	Show in Genome Browser
chr12:1609657-1613590	LOC100292680	0.16	23.18%	0%	0%	0%	0%	Show in Genome Browser
chr15:89921273-89941718	LOC254559	0.16	11.85%	0%	0%	0%	0%	Show in Genome Browser
chr19:37757716-37759912	LOC284412	0.16	15.47%	0%	0%	0%	0%	Show in Genome Browser
chr3:63088364-63110735	LOC285401	0.16	20.4%	0%	0%	0%	0%	Show in Genome Browser
chr2:6122110-6128364	LOC400940	0.16	16.16%	0%	0%	0%	0%	Show in Genome Browser
chr7:26533415-26534508	LOC441204	0.16	16.45%	0%	0%	0%	0%	Show in Genome Browser
chr18:24267585-24283602	LOC728606	0.16	9.46%	0%	0%	0%	0%	Show in Genome

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chr11:61276272-61278484	LRRC10B	0.16	14.14%	0%	0%	0%	0%	Show in Genome Browser
chr17:72206141-72209460	MGC16275	0.16	15.81%	0%	0%	0%	0%	Show in Genome Browser
chr1:178060643-178062863	RASAL2-AS1	0.16	12.24%	0%	0%	0%	0%	Show in Genome Browser
chr3:156390964-156393502	TIPARP,TIPARP-AS1	0.16	14.1%	0%	0%	0%	0%	Show in Genome Browser
chr1:11865945-11866160	CLCN6,MTHFR	0.17	17.12%	0%	0%	0%	0%	Show in Genome Browser
chrX:49641328-49643350	LOC158572	0.17	17.79%	0%	0%	0%	0%	Show in Genome Browser
chr5:158875564-158886657	LOC285627	0.17	15.36%	0%	0%	0%	0%	Show in Genome Browser
chr2:240115027-240117153	MGC16025	0.17	13.39%	0%	0%	0%	0%	Show in Genome Browser
chr11:65190269-65194001	NEAT1	0.17	15.96%	0%	0%	0%	0%	Show in Genome Browser
chr8:144448794-144450805	RHPN1-AS1	0.17	16.25%	0%	0%	0%	0%	Show in Genome Browser
chrX:47863734-47986684	SPACA5	0.17	17.07%	0%	0%	0%	0%	Show in Genome Browser
chr15:65385344-65398697	UBAP1L	0.17	18.97%	0%	0%	0%	0%	Show in Genome Browser
chr6:31620177-31620477	APOM,BAG6	0.18	18.27%	0%	0%	0%	0%	Show in Genome Browser
chr11:2289729-2292182	ASCL2	0.18	17.31%	0%	0%	0%	0%	Show in Genome Browser
chr16:86588926-86590905	FLJ30679	0.18	16.96%	0%	0%	0%	0%	Show in Genome Browser
chr6:168376604-168377619	HGC6.3	0.18	18.3%	0%	0%	0%	0%	Show in Genome Browser
chr7:27225027-27228912	HOXA11-AS	0.18	17.17%	0%	0%	0%	0%	Show in Genome Browser
chr15:37156644-37178734	LOC145845	0.18	13.15%	0%	0%	0%	0%	Show in Genome Browser
chr11:1330938-1331936	LOC255512	0.18	12.81%	0%	0%	0%	0%	Show in Genome Browser
chr3:106964855-107045810	LOC344595	0.18	18.85%	0%	0%	0%	0%	Show in Genome Browser

chr15:44957931-44968763	PATL2	0.18	17.52%	0%	0%	0%	0%	Show in Genome Browser
chr11:64808376-64809338	SAC3D1	0.18	18.69%	0%	0%	0%	0%	Show in Genome Browser
chrX:118599997-118603083	SLC25A5-AS1	0.18	5.9%	0%	0%	0%	0%	Show in Genome Browser
chr7:45022628-45026259	SNHG15	0.18	8.03%	0%	0%	0%	0%	Show in Genome Browser
chr17:79139307-79156964	AATK-AS1	0.19	22.9%	0%	0%	0%	0%	Show in Genome Browser
chr6:161581174-161583014	AGPAT4-IT1	0.19	16.89%	0%	0%	0%	0%	Show in Genome Browser
chr19:1267471-1268389	CIRBP-AS1	0.19	15.34%	0%	0%	0%	0%	Show in Genome Browser
chr5:92745065-92916953	FLJ42709	0.19	9.83%	0%	0%	0%	0%	Show in Genome Browser
chr9:98568372-98576204	LINC00476	0.19	15.9%	0%	0%	0%	0%	Show in Genome Browser
chr3:45265957-45267814	TMEM158	0.19	17.16%	0%	0%	0%	0%	Show in Genome Browser
chr2:74612845-74621008	DCTN1,DCTN1-AS1	0.2	14.92%	0%	0%	0%	0%	Show in Genome Browser
chr14:24036948-24037243	JPH4,AP1G2	0.2	20.6%	0%	0%	0%	0%	Show in Genome Browser
chr11:2661768-2721228	KCNQ1OT1	0.2	11.25%	0.47%	0.32%	0.13%	0.03%	Show in Genome Browser
chr6:43858765-43905943	LOC100132354	0.2	8.22%	0%	0%	0%	0%	Show in Genome Browser
chr22:46435789-46440748	LOC100271722	0.2	8.09%	0%	0%	0%	0%	Show in Genome Browser
chr2:16080020-16080865	MYCNOS	0.2	11.28%	0%	0%	0%	0%	Show in Genome Browser
chr10:126135999-126138550	NKX1-2	0.2	23.72%	0%	0%	0%	0%	Show in Genome Browser
chr17:1421012-1423832	PITPNA,PITPNA-AS1	0.2	15.01%	0%	0%	0%	0%	Show in Genome Browser
chr1:48567387-48648100	SKINTL	0.2	17.49%	0%	0%	0%	0%	Show in Genome Browser
chr14:105267518-105271046	ZBTB42	0.2	18.33%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr22:50171538-50173958	BRD1,LOC90834	0.21	15.44%	0%	0%	0%	0%	Genome Browser
chr3:129931663-129992648	COL6A4P2	0.21	12.54%	0%	0%	0%	0%	Show in Genome Browser
chr1:175126123-175161949	KIAA0040	0.21	20.6%	0%	0%	0%	0%	Show in Genome Browser
chr14:105287538-105290055	LINC00638	0.21	13.62%	0%	0%	0%	0%	Show in Genome Browser
chr17:8261731-8263859	LOC100128288	0.21	21.41%	0%	0%	0%	0%	Show in Genome Browser
chr1:852953-853555	LOC100130417	0.21	20.85%	0%	0%	0%	0%	Show in Genome Browser
chr5:6582287-6588612	LOC255167	0.21	16.99%	0%	0%	0%	0%	Show in Genome Browser
chr9:107536633-107540044	LOC286367	0.21	20.92%	0%	0%	0%	0%	Show in Genome Browser
chr5:148803653-148808480	MIR143,MIR143HG	0.21	13.27%	0.82%	0%	0%	0%	Show in Genome Browser
chr4:90757894-90758350	SNCA,LOC644248	0.21	21.22%	0%	0%	0%	0%	Show in Genome Browser
chr11:694759-695615	TMEM80,DEAF1	0.21	13.18%	0%	0%	0%	0%	Show in Genome Browser
chr21:43442113-43445060	ZNF295-AS1	0.21	15%	0%	0%	0%	0%	Show in Genome Browser
chr1:111023388-111031256	CYMP	0.22	13.07%	0%	0%	0%	0%	Show in Genome Browser
chr13:50556688-50568097	DLEU2	0.22	19.17%	0%	0%	0%	0%	Show in Genome Browser
chr15:69383255-69388161	LINC00277	0.22	6.64%	0%	0%	0%	0%	Show in Genome Browser
chr15:74418714-74421619	LOC283731	0.22	11.72%	0%	0%	0%	0%	Show in Genome Browser
chr2:121221911-121223925	LOC84931	0.22	20.24%	0%	0%	0%	0%	Show in Genome Browser
chr2:173587920-173600189	RAPGEF4-AS1	0.22	12.4%	0%	0%	0%	0%	Show in Genome Browser
chr6:30296666-30296818	TRIM39	0.22	22.87%	0%	0%	0%	0%	Show in Genome Browser
chr22:28315364-28318166	TTC28-AS1	0.22	23.26%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr10:31605458-31607423	ZEB1-AS1	0.22	10.52%	0%	0%	0%	0%	Genome Browser
chr16:88493879-88507165	ZNF469	0.22	19.89%	0%	0%	0%	0%	Show in Genome Browser
chr1:151694013-151702082	RIAD1	0.23	32.92%	0%	0%	0%	0%	Show in Genome Browser
chr19:33793763-33795962	CEBPA-AS1	0.24	16.9%	0%	0%	0%	0%	Show in Genome Browser
chr9:139216998-139221779	DKFZP434A062	0.24	18.27%	0%	0%	0%	0%	Show in Genome Browser
chr3:197025289-197025447	DLG1-AS1	0.24	24.52%	0%	0%	0%	0%	Show in Genome Browser
chr19:7004964-7006426	FLJ25758	0.24	17.42%	0%	0%	0%	0%	Show in Genome Browser
chr8:145537788-145540372	HSF1,DGAT1	0.24	13.52%	1.1%	0%	0%	0%	Show in Genome Browser
chr6:170188886-170198921	LINC00242,LINC00574	0.24	10.54%	0%	0%	0%	0%	Show in Genome Browser
chr11:67085310-67088804	LOC100130987	0.24	42.16%	0%	0%	0%	0%	Show in Genome Browser
chr22:20193855-20196060	LOC150197	0.24	20.89%	0%	0%	0%	0%	Show in Genome Browser
chr2:105050805-105129214	LOC150568	0.24	8.25%	0%	0%	0%	0%	Show in Genome Browser
chr12:50302668-50304837	LOC283332	0.24	21.29%	0%	0%	0%	0%	Show in Genome Browser
chr1:4472111-4483247	LOC284661	0.24	34.47%	0%	0%	0%	0%	Show in Genome Browser
chrX:130836679-130964671	LOC286467	0.24	18.5%	4.62%	0%	0%	0%	Show in Genome Browser
chr12:46119503-46121704	LOC400027	0.24	14.94%	0%	0%	0%	0%	Show in Genome Browser
chr3:116428635-116435885	LSAMP-AS3	0.24	15.79%	0%	0%	0%	0%	Show in Genome Browser
chr20:61885330-61885748	NKAIN4,FLJ16779	0.24	24.34%	0%	0%	0%	0%	Show in Genome Browser
chr6:32121622-32122181	PPT2-EGFL8,LOC100507547	0.24	24.28%	0%	0%	0%	0%	Show in Genome Browser
chr12:51785071-51785200	SLC4A8,GALNT6	0.24	24.61%	0%	0%	0%	0%	Show in Genome Browser
chr16:2787077-								Show in

2802442	SRRM2-AS1	0.24	27.2%	0%	0%	0%	0%	Genome Browser
chr14:98391947-98444461	C14orf64	0.25	11.97%	0%	0%	0%	0%	Show in Genome Browser
chr22:46449630-46450024	C22orf26,LOC150381	0.25	26.61%	0%	0%	0%	0%	Show in Genome Browser
chr1:47897808-47900313	FOXD2-AS1	0.25	20.63%	0%	0%	0%	0%	Show in Genome Browser
chr5:87836597-87980620	LINC00461	0.25	15.22%	0%	0%	0%	0%	Show in Genome Browser
chr4:1189572-1202750	LOC100130872,SPON2	0.25	25.1%	0%	0%	0%	0%	Show in Genome Browser
chr6:163739826-163745101	PACRG-AS1	0.25	9.36%	0%	0%	0%	0%	Show in Genome Browser
chr1:202976534-202993197	TMEM183B,TMEM183A	0.25	0.09%	0%	0%	0%	0%	Show in Genome Browser
chr6:88032306-88032761	C6orf162,GJB7	0.26	40.94%	0%	0%	0%	0%	Show in Genome Browser
chr16:72127769-72128215	DHX38,TXNL4B	0.26	26.56%	0%	0%	0%	0%	Show in Genome Browser
chr9:131580779-131581464	ENDOG	0.26	26.09%	0%	0%	0%	0%	Show in Genome Browser
chr3:47323985-47324329	KIF9,KLHL18	0.26	26.37%	0%	0%	0%	0%	Show in Genome Browser
chr15:57592563-57599266	LOC283663	0.26	32.17%	0%	0%	0%	0%	Show in Genome Browser
chr1:159824107-159825137	VSIG8,C1orf204	0.26	37.45%	0%	0%	0%	0%	Show in Genome Browser
chr2:53897118-53897449	ASB3,GPR75-ASB3	0.27	27.1%	0%	0%	0%	0%	Show in Genome Browser
chr11:43964106-43965432	C11orf96	0.27	22.6%	0%	0%	0%	0%	Show in Genome Browser
chr1:41326729-41328018	CITED4	0.27	19.06%	0%	0%	0%	0%	Show in Genome Browser
chr17:41857803-41857954	C17orf105	0.28	26.31%	0%	0%	0%	0%	Show in Genome Browser
chr12:54452038-54452348	FLJ12825	0.28	28.93%	0%	0%	0%	0%	Show in Genome Browser
chr8:144063448-144099807	LOC100133669	0.28	20.24%	0%	0%	0%	0%	Show in Genome Browser
chr3:47053032-								Show in

47054957	NRADDP	0.28	35.73%	0%	0%	0%	0%	Genome Browser
chr17:17680000-17682843	SMCR5,RAI1	0.28	19.09%	0%	0%	0%	0%	Show in Genome Browser
chr12:66218240-66220754	HMG2, RPSAP52	0.29	28.96%	0%	0%	0%	0%	Show in Genome Browser
chr7:156333185-156333795	LINC00244	0.29	24.05%	0%	0%	0%	0%	Show in Genome Browser
chr16:1114084-1116526	SSTR5-AS1	0.29	25.01%	0%	0%	0%	0%	Show in Genome Browser
chr4:13547700-13549448	LOC285548	0.3	12.52%	0.28%	0%	0%	0%	Show in Genome Browser
chr12:54519855-54526619	LOC400043	0.3	23.2%	0%	0%	0%	0%	Show in Genome Browser
chr17:33875144-33885110	SLFN14	0.3	23.82%	0%	0%	0%	0%	Show in Genome Browser
chr14:106435819-106438358	ADAM6	0.31	13.56%	0%	0%	0%	0%	Show in Genome Browser
chr12:74931551-74935223	ATXN7L3B	0.31	20.28%	0.4%	0%	0%	0%	Show in Genome Browser
chr13:32420920-32533720	EEF1DP3	0.31	28.34%	0%	0%	0%	0%	Show in Genome Browser
chr10:124639001-124639148	FAM24B, FAM24B-CUZD1, LOC399815	0.31	31.08%	0%	0%	0%	0%	Show in Genome Browser
chr15:70127573-70135305	LINC00593	0.31	20.61%	0%	0%	0%	0%	Show in Genome Browser
chr10:112628648-112630662	LOC282997	0.31	22.18%	0%	0%	0%	0%	Show in Genome Browser
chr10:120789228-120793854	NANOS1	0.31	16.27%	1.81%	0%	0%	0%	Show in Genome Browser
chr5:111310971-111311075	NREP, NREP-AS1	0.31	31.42%	0%	0%	0%	0%	Show in Genome Browser
chr21:42547159-42557166	PLAC4, BACE2	0.31	14.85%	1.5%	0.36%	0%	0%	Show in Genome Browser
chr12:55001971-55003461	GLYCAM1	0.32	32.78%	0%	0%	0%	0%	Show in Genome Browser
chr13:44453420-44453766	LACC1, CCDC122	0.32	32.27%	0%	0%	0%	0%	Show in Genome Browser
chr16:576847-577407	LINC00235	0.32	22.99%	0%	0%	0%	0%	Show in Genome Browser
chr17:61904412-61904809	PSMC5, FTSJ3	0.32	23.11%	0%	0%	0%	0%	Show in Genome Browser

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chr22:19709951-19711065	SEPT5,GP1BB,SEPT5-GP1BB	0.32	12.46%	0.89%	0%	0%	0%	Show in Genome Browser
chr12:120031264-120079363	TMEM233	0.32	23.85%	0%	0%	0%	0%	Show in Genome Browser
chr7:6655527-6663919	ZNF853	0.32	21.98%	3.05%	0%	0%	0%	Show in Genome Browser
chr9:136890561-136891212	LINC00094	0.33	43.1%	0%	0%	0%	0%	Show in Genome Browser
chr2:33153099-33162270	LINC00486,LOC100271832	0.33	17.22%	0%	0%	0%	0%	Show in Genome Browser
chr20:46993632-46999381	LINC00494	0.33	23.3%	0%	0%	0%	0%	Show in Genome Browser
chr19:50431974-50432786	NUP62,IL4I1,ATF5	0.33	20.56%	0%	0%	0%	0%	Show in Genome Browser
chr14:57279901-57397548	OTX2-AS1	0.33	12.23%	0%	0%	0%	0%	Show in Genome Browser
chr1:43424305-43424847	SLC2A1,SLC2A1-AS1	0.33	25.04%	0%	0%	0%	0%	Show in Genome Browser
chr11:61129205-61129472	TMEM138,CYBASC3	0.33	33.58%	0%	0%	0%	0%	Show in Genome Browser
chr7:150020296-150020557	ACTR3C,LRRRC61	0.34	34.35%	0%	0%	0%	0%	Show in Genome Browser
chr4:146019156-146019282	ANAPC10,ABCE1	0.34	34.64%	0%	0%	0%	0%	Show in Genome Browser
chr21:46340736-46348753	ITGB2-AS1	0.34	34.21%	0%	0%	0%	0%	Show in Genome Browser
chr1:22352042-22357713	LINC00339	0.34	30.3%	0%	0%	0%	0%	Show in Genome Browser
chr8:9757574-9760839	LINC00599	0.34	16.37%	0.98%	0%	0%	0%	Show in Genome Browser
chr11:573809-575885	LOC143666	0.34	18.1%	0%	0%	0%	0%	Show in Genome Browser
chr8:142353057-142353314	LOC731779	0.34	34.88%	0%	0%	0%	0%	Show in Genome Browser
chr10:99473481-99477905	MARVELD1	0.34	26.46%	5.52%	0%	0%	0%	Show in Genome Browser
chr3:23958275-23958537	RPL15,NKIRAS1	0.34	34.22%	0%	0%	0%	0%	Show in Genome Browser
chr9:109625378-109625636	ZNF462	0.34	34.74%	0%	0%	0%	0%	Show in Genome Browser

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chr7:123454193-123459484	HYALP1	0.35	6.35%	0%	0%	0%	0%	Show in Genome Browser
chr12:49121218-49159568	LOC255411	0.35	31.45%	0%	0%	0%	0%	Show in Genome Browser
chr1:13516066-13737001	PRAMEF20	0.36	25.26%	0%	0%	0%	0%	Show in Genome Browser
chr1:202137179-202158120	PTPRVP	0.36	21.33%	0%	0%	0%	0%	Show in Genome Browser
chr6:111926979-111927474	TRAF3IP2	0.36	36.29%	0%	0%	0%	0%	Show in Genome Browser
chr15:73075587-73076126	ADPGK-AS1	0.37	34.25%	0%	0%	0%	0%	Show in Genome Browser
chr8:136246374-136311959	LOC286094	0.37	25.88%	0%	0%	0%	0%	Show in Genome Browser
chr1:28975112-28975246	RNU11	0.37	37.77%	0%	0%	0%	0%	Show in Genome Browser
chr1:45272456-45272957	TCTEX1D4	0.37	35.65%	0%	0%	0%	0%	Show in Genome Browser
chr6:31975911-31976684	CYP21A2,TNXA,CYP21A1P,TNXB	0.38	27.21%	0%	0%	0%	0%	Show in Genome Browser
chr3:13729612-13788132	LINC00620	0.38	18.02%	0%	0%	0%	0%	Show in Genome Browser
chr9:94895116-94900911	LOC100128076	0.38	38.13%	0%	0%	0%	0%	Show in Genome Browser
chr9:139979398-139981269	LOC100289341	0.38	31.3%	0%	0%	0%	0%	Show in Genome Browser
chr13:50571143-50571555	TRIM13,DLEU2	0.38	43.07%	0%	0%	0%	0%	Show in Genome Browser
chr1:247273462-247275719	C1orf229	0.39	25.11%	0%	0%	0%	0%	Show in Genome Browser
chr22:19553653-19554362	LOC150185	0.39	35.63%	0%	0%	0%	0%	Show in Genome Browser
chr9:100153121-100158973	LOC286359	0.39	35.82%	0.39%	0%	0%	0%	Show in Genome Browser
chr16:69598997-69600277	NFAT5,MIR1538	0.39	18.34%	0%	0%	0%	0%	Show in Genome Browser
chr14:70234098-70234430	SRSF5,LOC100289511	0.39	39.33%	0%	0%	0%	0%	Show in Genome Browser
chr7:32767562-32769595	ZNRF2P1	0.39	21.28%	0%	0%	0%	0%	Show in Genome Browser

chr3:38495994-38496311	ACVR2B,ACVR2B-AS1	0.4	40.25%	0%	0%	0%	0%	Show in Genome Browser
chr12:113594984-113597080	CCDC42B,DDX54	0.4	38.37%	0%	0%	0%	0%	Show in Genome Browser
chr12:104324189-104324342	HSP90B1,MIR3652	0.4	40.25%	0%	0%	0%	0%	Show in Genome Browser
chr10:4692378-4720262	LOC100216001	0.4	17.36%	0%	0%	0%	0%	Show in Genome Browser
chr4:90759403-90759447	LOC644248,SNCA	0.4	40%	0%	0%	0%	0%	Show in Genome Browser
chr17:46801690-46802119	HOXB-AS5,MIR3185	0.41	34.18%	0%	0%	0%	0%	Show in Genome Browser
chr6:170563422-170571657	LOC154449	0.41	13.09%	0%	0%	0%	0%	Show in Genome Browser
chr3:191046874-191048325	UTS2D,CCDC50	0.41	22.86%	0%	0%	0%	0%	Show in Genome Browser
chr17:42263933-42264082	C17orf65	0.42	42.66%	0%	0%	0%	0%	Show in Genome Browser
chr15:75074425-75075061	CSK	0.42	37.21%	0%	0%	0%	0%	Show in Genome Browser
chr2:172290761-172291183	DCAF17,METTL8	0.42	31.67%	0%	0%	0%	0%	Show in Genome Browser
chr8:12219528-12268510	FAM66A	0.42	22.84%	0%	0%	0%	0%	Show in Genome Browser
chr12:52604714-52615812	LINC00592	0.42	45.29%	0%	0%	0%	0%	Show in Genome Browser
chr11:44995453-44999576	LOC221122	0.42	44.31%	0%	0%	0%	0%	Show in Genome Browser
chr11:1618409-1618477	MOB2,KRTAP5-2,LOC338651	0.42	42.02%	0%	0%	0%	0%	Show in Genome Browser
chr4:69078189-69083798	TMPRSS11BNL	0.42	2.86%	0%	0%	0%	0%	Show in Genome Browser
chr6:112408613-112408673	TUBE1,C6orf225	0.42	42.62%	0%	0%	0%	0%	Show in Genome Browser
chr19:58869855-58874120	ZNF497	0.42	46.34%	0%	0%	0%	0%	Show in Genome Browser
chr19:45653009-45663408	NKPD1	0.43	42.35%	5.59%	0.06%	0%	0%	Show in Genome Browser
chr13:21750372-21750513	SKA3,MRP63	0.43	32.39%	0%	0%	0%	0%	Show in Genome Browser

chr16:1128458-1128731	SSTR5,SSTR5-AS1	0.43	43.06%	0%	0%	0%	0%	Show in Genome Browser
chr2:145277506-145277916	ZEB2-AS1	0.43	36.98%	0%	0%	0%	0%	Show in Genome Browser
chr14:102018560-102022013	DIO3OS	0.44	27.24%	0%	0%	0%	0%	Show in Genome Browser
chr10:102989351-102998615	FLJ41350	0.44	33.11%	0%	0%	0%	0%	Show in Genome Browser
chr14:51955855-52037128	FRMD6-AS2	0.44	15.69%	0%	0%	0%	0%	Show in Genome Browser
chr19:56347702-56347943	NLRP4,NLRP11	0.44	44.62%	0%	0%	0%	0%	Show in Genome Browser
chr2:179315693-179315958	PRKRA	0.44	40.22%	0%	0%	0%	0%	Show in Genome Browser
chr17:61934376-61940490	TCAM1P	0.44	37.01%	0%	0%	0%	0%	Show in Genome Browser
chr9:35732072-35732316	TLN1,CREB3	0.44	44.48%	0%	0%	0%	0%	Show in Genome Browser
chr14:21566970-21571882	ZNF219,C14orf176	0.44	40.55%	0%	0%	0%	0%	Show in Genome Browser
chr16:30200005-30200180	CORO1A,LOC606724	0.45	45.45%	0%	0%	0%	0%	Show in Genome Browser
chr21:43131680-43135935	LINC00479	0.45	33.17%	0%	0%	0%	0%	Show in Genome Browser
chr10:12237769-12237960	NUDT5,CDC123	0.45	45.31%	0%	0%	0%	0%	Show in Genome Browser
chr8:128902835-129108900	PVT1	0.45	27.01%	0%	0%	0%	0%	Show in Genome Browser
chr22:21055402-21058891	TMEM191A	0.45	19.12%	0%	0%	0%	0%	Show in Genome Browser
chr6:31497524-31497997	ATP6V1G2-DDX39B,MCCD1	0.46	32.7%	0%	0%	0%	0%	Show in Genome Browser
chr2:79740060-79740333	CTNNA2	0.46	46.71%	0%	0%	0%	0%	Show in Genome Browser
chr10:91451057-91457685	FLJ37201	0.46	3.96%	0%	0%	0%	0%	Show in Genome Browser
chr11:105948039-105948291	KBTBD3,AASDHPPT	0.46	46.64%	0%	0%	0%	0%	Show in Genome Browser
chr20:62665697-62671314	LINC00176	0.46	23.71%	0.66%	0%	0%	0%	Show in Genome Browser
								Show in

chr18:1268312-1269853	LINC00470	0.46	34.17%	0%	0%	0%	0%	Genome Browser
chr13:101183811-101185997	A2LD1	0.47	39.91%	0%	0%	0%	0%	Show in Genome Browser
chr19:40854332-40854675	C19orf47,PLD3	0.47	25.87%	0%	0%	0%	0%	Show in Genome Browser
chr19:51600864-51611647	CTU1	0.47	41.56%	0%	0%	0%	0%	Show in Genome Browser
chr17:66194801-66196436	LOC440461	0.47	35.39%	0%	0%	0%	0%	Show in Genome Browser
chr4:110354971-110355158	SEC24B,SEC24B-AS1	0.47	47.87%	0%	0%	0%	0%	Show in Genome Browser
chr2:27886196-27886337	SUPT7L,SLC4A1AP	0.47	47.88%	0%	0%	0%	0%	Show in Genome Browser
chr10:45493146-45495864	C10orf25	0.48	26.99%	2.39%	0%	0%	0%	Show in Genome Browser
chr20:61147660-61148731	C20orf166,C20orf166-AS1	0.48	13.91%	0%	0%	0%	0%	Show in Genome Browser
chr6:35694540-35694910	FKBP5,LOC285847	0.48	48.51%	0%	0%	0%	0%	Show in Genome Browser
chr21:43136596-43137334	LINC00112	0.48	49.12%	0%	0%	0%	0%	Show in Genome Browser
chr21:36410235-36411723	RUNX1,RUNX1-IT1	0.48	39.62%	0%	0%	0%	0%	Show in Genome Browser
chr1:3689352-3692545	SMIM1	0.48	53.26%	0%	0%	0%	0%	Show in Genome Browser
chr6:32162621-32163202	GPSM3,NOTCH4	0.49	31.78%	0%	0%	0%	0%	Show in Genome Browser
chr7:65841032-65860943	LINC00174	0.49	40.65%	6.19%	0.96%	0%	0%	Show in Genome Browser
chr15:62929371-62937380	MGC15885	0.49	16.21%	0%	0%	0%	0%	Show in Genome Browser
chr16:4838281-4838397	SEPT12,LOC440335	0.49	49.57%	0%	0%	0%	0%	Show in Genome Browser
chr10:111765726-111768057	ADD3,LOC100505933	0.5	24.06%	0%	0%	0%	0%	Show in Genome Browser
chr15:50647077-50650501	GABPB1-AS1	0.5	3.43%	0%	0%	0%	0%	Show in Genome Browser
chr19:55996594-55998935	NAT14	0.5	54.15%	5.64%	0%	0%	0%	Show in Genome Browser
chr6:32121301-								Show in

32121413	PPT2,LOC100507547	0.5	50.44%	0%	0%	0%	0%	Genome Browser
chr20:35807456-35807974	RPN2,C20orf132	0.5	43.83%	0%	0%	0%	0%	Show in Genome Browser
chr19:35067639-35068596	SCGB1B2P	0.5	44.93%	0%	0%	0%	0%	Show in Genome Browser
chr12:93963598-93963974	SOCS2-AS1	0.5	33.68%	0%	0%	0%	0%	Show in Genome Browser
chr11:65769381-65769549	EIF1AD,BANF1	0.51	51.47%	0%	0%	0%	0%	Show in Genome Browser
chr4:135117491-135122400	PABPC4L	0.51	17.98%	3.68%	0.81%	0%	0%	Show in Genome Browser
chr13:70703916-70713885	ATXN8OS	0.52	47.29%	0%	0%	0%	0%	Show in Genome Browser
chr8:144779285-144780582	BREA2	0.52	44.75%	0%	0%	0%	0%	Show in Genome Browser
chr12:31226779-31227025	DDX11-AS1	0.52	52.22%	0%	0%	0%	0%	Show in Genome Browser
chr3:49190293-49191834	LAMB2P1	0.52	39.81%	0%	0%	0%	0%	Show in Genome Browser
chr9:27245684-27282791	LINC00032	0.52	39.9%	0%	0%	0%	0%	Show in Genome Browser
chr1:1334910-1335683	LOC148413	0.52	39.54%	0%	0%	0%	0%	Show in Genome Browser
chr1:26146397-26146683	LOC646471,FAM54B	0.52	72.28%	0%	0%	0%	0%	Show in Genome Browser
chr8:29996048-30002200	MIR548O2,MBOAT4	0.52	38.88%	0%	0%	0%	0%	Show in Genome Browser
chr9:140172280-140177092	TOR4A	0.52	33.95%	4.23%	0%	0%	0%	Show in Genome Browser
chr3:125899319-125899485	ALDH1L1-AS2	0.53	53.89%	0%	0%	0%	0%	Show in Genome Browser
chr10:99092254-99094458	FRAT2	0.53	22.04%	0.58%	0%	0%	0%	Show in Genome Browser
chr5:134984374-134989624	LOC340074	0.53	20.76%	0%	0%	0%	0%	Show in Genome Browser
chr17:43325303-43339710	MAP3K14-AS1	0.53	38.78%	0%	0%	0%	0%	Show in Genome Browser
chr2:111334679-111334902	RGPD6	0.53	53.12%	0%	0%	0%	0%	Show in Genome Browser
chr10:135043778-								Show in

135045061	UTF1	0.53	30.36%	0%	0%	0%	0%	Genome Browser
chr16:67184768-67184902	B3GNT9	0.54	28.14%	0%	0%	0%	0%	Show in Genome Browser
chr11:66742754-66744478	C11orf86	0.54	30.81%	0%	0%	0%	0%	Show in Genome Browser
chr9:69175960-69178388	LOC440896	0.54	30.42%	0%	0%	0%	0%	Show in Genome Browser
chr7:150102840-150109557	LOC728743	0.54	36.35%	3.09%	0%	0%	0%	Show in Genome Browser
chr18:77915119-77936313	PARD6G-AS1	0.54	14.82%	6.18%	2.67%	0.29%	0%	Show in Genome Browser
chrX:109662285-109662363	AMMECR1, RGAG1	0.55	44.3%	0%	0%	0%	0%	Show in Genome Browser
chr7:106299150-106301591	CCDC71L	0.55	37.05%	0%	0%	0%	0%	Show in Genome Browser
chr5:140071018-140071341	HARS, HARS2	0.55	36.41%	0%	0%	0%	0%	Show in Genome Browser
chr22:51021455-51021752	LOC100144603	0.55	55.7%	0%	0%	0%	0%	Show in Genome Browser
chr10:102849079-102890903	TLX1NB	0.55	31.86%	8.62%	1.91%	0%	0%	Show in Genome Browser
chr1:27320195-27327376	TRNP1	0.55	28.33%	4.36%	0%	0%	0%	Show in Genome Browser
chr19:46714866-46718092	DKFZp434J0226	0.56	54.88%	4.67%	0%	0%	0%	Show in Genome Browser
chr9:140657474-140659222	EHMT1, FLJ40292	0.56	41.79%	0%	0%	0%	0%	Show in Genome Browser
chr19:49122548-49122800	SPHK2, RPL18	0.56	52.96%	0%	0%	0%	0%	Show in Genome Browser
chr19:3750771-3751095	TJP3, APBA3	0.56	46.15%	0%	0%	0%	0%	Show in Genome Browser
chr1:156307945-156308195	TSACC, CCT3	0.56	56.17%	0%	0%	0%	0%	Show in Genome Browser
chr17:43716341-43719669	C17orf69	0.57	24.02%	6.9%	3.33%	0%	0%	Show in Genome Browser
chr5:169758435-169762104	LOC257358	0.57	14.07%	0%	0%	0%	0%	Show in Genome Browser
chr13:28712684-28713311	PAN3, PAN3-AS1	0.57	31.05%	0%	0%	0%	0%	Show in Genome Browser
chr16:2014997-	SNHG9	0.57	34.32%	0%	0%	0%	0%	Show in Genome Browser

2015130								Browser
chr19:56152392-56154835	ZNF580	0.57	36.55%	5.64%	0%	0%	0%	Show in Genome Browser
chr17:36827961-36831187	C17orf96	0.58	22.77%	4.58%	0%	0%	0%	Show in Genome Browser
chr19:405444-409170	C2CD4C	0.58	47.05%	6.85%	1.53%	0%	0%	Show in Genome Browser
chr7:156431062-156433348	C7orf13	0.58	37.34%	0.21%	0%	0%	0%	Show in Genome Browser
chr11:61100380-61100653	DDB1,DAK	0.58	46.35%	0%	0%	0%	0%	Show in Genome Browser
chr9:136891330-136898827	LINC00094,BRD3	0.58	13.82%	3.21%	1.62%	0%	0%	Show in Genome Browser
chr12:54971644-54973023	PDE1B,PPP1R1A	0.58	40.57%	0%	0%	0%	0%	Show in Genome Browser
chrX:48755195-48755349	PQBP1,TIMM17B	0.58	58.06%	0%	0%	0%	0%	Show in Genome Browser
chr11:111383663-111383746	MIR34B	0.59	59.52%	0%	0%	0%	0%	Show in Genome Browser
chr17:79762010-79771889	GCGR	0.6	48.13%	0%	0%	0%	0%	Show in Genome Browser
chr6:29892369-29893428	HCG4B	0.6	27.26%	2.92%	0%	0%	0%	Show in Genome Browser
chr16:90113365-90113504	LOC100130015	0.6	60%	0%	0%	0%	0%	Show in Genome Browser
chr3:48599151-48601201	UCN2	0.6	38.81%	7.88%	0%	0%	0%	Show in Genome Browser
chr2:242254723-242254984	HDLBP,SEPT2	0.61	38.54%	0%	0%	0%	0%	Show in Genome Browser
chr2:47168313-47168864	MCFD2,TTC7A	0.61	38.4%	0%	0%	0%	0%	Show in Genome Browser
chr14:58740697-58764855	FLJ31306	0.62	43.39%	3.75%	0%	0%	0%	Show in Genome Browser
chr19:4501923-4504797	HDGFRP2,PLIN4	0.62	34.03%	6.83%	0.17%	0%	0%	Show in Genome Browser
chr13:52387483-52419286	LINC00282	0.62	27.77%	16.54%	11.62%	3.68%	1.31%	Show in Genome Browser
chr2:64751465-64751746	AFTPH	0.63	46.8%	0%	0%	0%	0%	Show in Genome Browser
chrX:101615317-	NXF2	0.63	27.35%	0%	0%	0%	0%	Show in Genome

101694853								Browser
chr21:34442450-34444727	OLIG1	0.63	11.54%	8.42%	0%	0%	0%	Show in Genome Browser
chr10:135237981-135238121	SPRN	0.63	63.82%	0%	0%	0%	0%	Show in Genome Browser
chr1:185126192-185126446	TRMT1L,SWT1	0.63	63.13%	0%	0%	0%	0%	Show in Genome Browser
chr19:36705567-36705789	ZNF146,ZNF565	0.63	57.39%	0%	0%	0%	0%	Show in Genome Browser
chr17:73642646-73642866	C17orf110	0.64	33.93%	0%	0%	0%	0%	Show in Genome Browser
chr13:24463028-24463559	C1QTNF9B-AS1,MIPEP	0.64	56.26%	0%	0%	0%	0%	Show in Genome Browser
chr15:83654995-83655121	FAM103A1	0.65	65.35%	0%	0%	0%	0%	Show in Genome Browser
chr2:43454350-43455994	LOC100129726	0.65	36.59%	0%	0%	0%	0%	Show in Genome Browser
chr19:50168399-50168887	BCL2L12,IRF3	0.66	51.12%	0.81%	0%	0%	0%	Show in Genome Browser
chr2:55459065-55459449	CLHC1,RPS27A	0.66	68.64%	0%	0%	0%	0%	Show in Genome Browser
chr8:143738875-143751401	JRK	0.66	31.07%	11.65%	3.92%	0%	0%	Show in Genome Browser
chr10:42970939-42972890	LOC84856	0.66	21.46%	0%	0%	0%	0%	Show in Genome Browser
chr22:27053484-27072438	MIAT	0.66	22.23%	5.24%	3.94%	1.99%	1.37%	Show in Genome Browser
chr6:166719168-166721871	PRR18	0.66	20.89%	4.25%	0%	0%	0%	Show in Genome Browser
chr19:54573201-54584634	TARM1	0.66	50.36%	0%	0%	0%	0%	Show in Genome Browser
chr17:75875083-75879620	FLJ45079	0.67	32.96%	3.79%	0%	0%	0%	Show in Genome Browser
chr12:107168399-107168607	LOC100287944,RIC8B	0.67	58.37%	0%	0%	0%	0%	Show in Genome Browser
chr15:62359176-62363109	C2CD4A	0.68	30.93%	9.51%	4.6%	0%	0%	Show in Genome Browser
chr17:36889529-36891858	CISD3,PCGF2	0.68	56.99%	0%	0%	0%	0%	Show in Genome Browser
chr6:146056334-146056991	EPM2A,LOC100507557	0.68	41.94%	0%	0%	0%	0%	Show in Genome Browser

chr8:124213412-124213694	FAM83A,LOC100131726	0.68	43.81%	0%	0%	0%	0%	Show in Genome Browser
chr6:150184544-150185480	LOC100652739,LRP11	0.68	36.6%	0%	0%	0%	0%	Show in Genome Browser
chr11:2004440-2011150	MRPL23-AS1	0.68	27.5%	10.74%	1.34%	0%	0%	Show in Genome Browser
chr14:67936984-67982021	TMEM229B	0.68	29.46%	13.5%	6.54%	2.31%	0%	Show in Genome Browser
chr9:38621085-38623275	FAM201A	0.69	18.7%	1.33%	0%	0%	0%	Show in Genome Browser
chr7:73149399-73150330	ABHD11-AS1	0.7	54.8%	0%	0%	0%	0%	Show in Genome Browser
chr8:143279717-143288236	LINC00051	0.7	30.38%	0%	0%	0%	0%	Show in Genome Browser
chr6:961242-1101567	LOC285768	0.7	24.87%	10.4%	8.26%	4.79%	3.51%	Show in Genome Browser
chr19:36705504-36705523	ZNF565,ZNF146	0.7	70%	0%	0%	0%	0%	Show in Genome Browser
chr11:63527365-63536113	C11orf95	0.71	24.6%	1.63%	0%	0%	0%	Show in Genome Browser
chr13:24463592-24463692	C1QTNF9B-AS1	0.71	71.28%	0%	0%	0%	0%	Show in Genome Browser
chr14:57735340-57735605	EXOC5,AP5M1	0.71	35.71%	0%	0%	0%	0%	Show in Genome Browser
chr15:62455737-62457223	C2CD4B	0.72	35.37%	1.68%	0%	0%	0%	Show in Genome Browser
chr4:166300097-166300680	CPE	0.72	40.58%	0%	0%	0%	0%	Show in Genome Browser
chr6:30523908-30525227	GNL1,PRR3	0.72	32.22%	6.66%	0%	0%	0%	Show in Genome Browser
chr20:30073581-30075377	LINC00028	0.72	63.45%	0%	0%	0%	0%	Show in Genome Browser
chr13:19582399-19586774	LINC00442	0.72	46.3%	1.12%	0%	0%	0%	Show in Genome Browser
chr6:6346699-6569722	LY86-AS1	0.72	43.34%	21%	5.14%	0%	0%	Show in Genome Browser
chr11:60618562-60623444	PTGDR2	0.72	42.59%	9.03%	3.71%	0%	0%	Show in Genome Browser
chr17:8294023-8301144	RNF222	0.72	24.98%	8.4%	7.28%	0.07%	0%	Show in Genome Browser

chr9:137029562-137029686	RNU6ATAC	0.72	72%	0%	0%	0%	0%	Show in Genome Browser
chr3:14526375-14530855	SLC6A6,GRIP2	0.72	13.16%	4.5%	4.06%	0%	0%	Show in Genome Browser
chr19:48442798-48442917	SNAR-C4	0.72	67.5%	0%	0%	0%	0%	Show in Genome Browser
chr1:46685978-46686927	LURAP1	0.73	29.26%	4.52%	1.05%	0%	0%	Show in Genome Browser
chr1:41154752-41157933	NFYC,LOC100130557	0.73	12.64%	0.04%	0%	0%	0%	Show in Genome Browser
chr10:53059333-53059415	PRKG1,MIR605	0.73	39.75%	0%	0%	0%	0%	Show in Genome Browser
chr9:25676394-25678856	TUSC1	0.73	39.26%	3.24%	0%	0%	0%	Show in Genome Browser
chr17:76100566-76104916	TNRC6C,LOC100131096	0.74	16.11%	4.18%	3.3%	0.71%	0%	Show in Genome Browser
chr20:30776949-30778163	TSPY26P	0.74	50.37%	1.48%	0%	0%	0%	Show in Genome Browser
chr17:6918381-6918475	C17orf49,RNASEK-C17ORF49	0.75	75.78%	0%	0%	0%	0%	Show in Genome Browser
chr19:51385354-51398524	KLKP1	0.75	21.86%	0%	0%	0%	0%	Show in Genome Browser
chr8:143808621-143818350	THEM6	0.75	39.43%	0%	0%	0%	0%	Show in Genome Browser
chr9:2621793-2622373	VLDLR,FLJ35024	0.75	76.08%	0%	0%	0%	0%	Show in Genome Browser
chr16:89225628-89227967	LINC00304	0.76	29.72%	9.31%	7.53%	0%	0%	Show in Genome Browser
chr4:68566996-68588220	LOC550112	0.76	9.72%	0%	0%	0%	0%	Show in Genome Browser
chr12:6484671-6484905	SCNN1A,LTBR	0.76	38.72%	0%	0%	0%	0%	Show in Genome Browser
chr20:642241-656823	SCRT2	0.76	37.19%	9.09%	0.83%	0%	0%	Show in Genome Browser
chr6:32008646-32009419	TNXB,CYP21A2	0.76	44.78%	5.44%	0%	0%	0%	Show in Genome Browser
chr5:140098511-140098598	VTRNA1-2	0.76	76.13%	0%	0%	0%	0%	Show in Genome Browser
chr19:7989382-7990443	CTXN1	0.77	54.51%	0%	0%	0%	0%	Show in Genome Browser
								Show in

chr1:16174359-16174602	FLJ37453,SPEN	0.77	47.95%	0%	0%	0%	0%	Genome Browser
chr16:88782606-88782685	MIR4722,PIEZO1	0.77	77.5%	0%	0%	0%	0%	Show in Genome Browser
chr20:48729644-48732494	UBE2V1,TMEM189-UBE2V1	0.77	44.36%	0%	0%	0%	0%	Show in Genome Browser
chr19:45682003-45685057	BLOC1S3	0.78	30.51%	11.47%	6.07%	0%	0%	Show in Genome Browser
chr17:62075711-62078670	C17orf72	0.78	75.05%	0%	0%	0%	0%	Show in Genome Browser
chr12:47610052-47610166	PCED1B,PCED1B-AS1	0.78	78.26%	0%	0%	0%	0%	Show in Genome Browser
chr19:48453553-48453671	SNAR-C3	0.78	78.15%	0%	0%	0%	0%	Show in Genome Browser
chr22:30752627-30752718	CCDC157,SF3A1	0.79	79.34%	0%	0%	0%	0%	Show in Genome Browser
chr3:71803201-71803794	EIF4E3,GPR27	0.79	46.56%	0%	0%	0%	0%	Show in Genome Browser
chr9:131492067-131493673	LOC100506100,ZER1	0.79	25.57%	7.9%	0%	0%	0%	Show in Genome Browser
chr9:26946895-26947309	IFT74,PLAA	0.8	50.46%	0%	0%	0%	0%	Show in Genome Browser
chr1:1353802-1356385	ANKRD65	0.81	18.36%	0%	0%	0%	0%	Show in Genome Browser
chr12:110906232-110906276	FAM216A,GPN3	0.82	82.22%	0%	0%	0%	0%	Show in Genome Browser
chr11:19732481-19736146	LOC100126784,NAV2	0.82	10.06%	2.89%	1.61%	0%	0%	Show in Genome Browser
chr20:34287114-34287378	ROMO1,NFS1	0.82	24.31%	0%	0%	0%	0%	Show in Genome Browser
chr10:72972298-72972821	UNC5B	0.82	37.21%	0%	0%	0%	0%	Show in Genome Browser
chr11:10877691-10879620	ZBED5	0.82	34.57%	0%	0%	0%	0%	Show in Genome Browser
chr6:142958850-142959026	LOC153910	0.83	74.01%	0%	0%	0%	0%	Show in Genome Browser
chr2:28112325-28113123	BRE-AS1,RBKS	0.84	30.41%	5.75%	0%	0%	0%	Show in Genome Browser
chr4:128886461-128886761	C4orf29,MFSD8	0.84	52.49%	0%	0%	0%	0%	Show in Genome Browser
chr7:56563916-								Show in

56564977	DKFZp434L192	0.84	39.73%	2.91%	0%	0%	0%	Genome Browser
chr5:148786440-148801164	MIR143HG	0.84	37.07%	0%	0%	0%	0%	Show in Genome Browser
chr19:812518-813895	MIR3187,LPPR3	0.84	45.5%	0%	0%	0%	0%	Show in Genome Browser
chr2:28113224-28113481	BRE-AS1	0.85	40.31%	5.42%	0%	0%	0%	Show in Genome Browser
chr10:15139086-15139181	C10orf111,RPP38	0.85	85.41%	0%	0%	0%	0%	Show in Genome Browser
chr8:6875817-6875823	DEFA1	0.85	85.71%	0%	0%	0%	0%	Show in Genome Browser
chr12:7072862-7072929	MIR200C	0.85	85.29%	0%	0%	0%	0%	Show in Genome Browser
chr17:76227391-76237067	TMEM235	0.85	21.12%	0%	0%	0%	0%	Show in Genome Browser
chr15:83378346-83378635	AP3B2	0.86	52.41%	0%	0%	0%	0%	Show in Genome Browser
chr16:29753786-29757340	C16orf54	0.86	39.86%	7.59%	2.65%	0%	0%	Show in Genome Browser
chr2:26568954-26569685	EPT1,GPR113	0.86	39.03%	8.36%	0%	0%	0%	Show in Genome Browser
chr1:153232179-153234598	LOR	0.86	47.3%	12.31%	4.98%	0%	0%	Show in Genome Browser
chr17:79780293-79791167	FAM195B	0.87	47.38%	0%	0%	0%	0%	Show in Genome Browser
chr2:5832799-5841516	SOX11	0.87	24.12%	6.48%	2.29%	0%	0%	Show in Genome Browser
chr19:18988719-18989738	CERS1,GDF1	0.88	32.64%	6.37%	0.39%	0%	0%	Show in Genome Browser
chr17:5402748-5404542	LOC728392	0.88	33.86%	0%	0%	0%	0%	Show in Genome Browser
chr17:17717150-17717245	MIR33B,SREBF1	0.88	88.54%	0%	0%	0%	0%	Show in Genome Browser
chr12:56709605-56711427	PAN2,CNPY2	0.88	23.88%	8.55%	5.15%	0.76%	0%	Show in Genome Browser
chr9:70970111-70972757	PGM5-AS1	0.88	42.68%	10.3%	0.1%	0%	0%	Show in Genome Browser
chr19:36486101-36487213	SDHAF1	0.88	39.98%	2.33%	0%	0%	0%	Show in Genome Browser
chr1:1167629-								Show in

1170418	B3GALT6	0.9	24.58%	7.99%	1.07%	0%	0%	Genome Browser
chr11:96074547-96076344	MIR1260B,MAML2	0.9	36.42%	8.34%	0%	0%	0%	Show in Genome Browser
chr1:116379001-116383165	NHLH2	0.9	24.73%	14.08%	5.66%	0%	0%	Show in Genome Browser
chr19:42806284-42806952	PAFAH1B3,PRR19	0.9	21.21%	0%	0%	0%	0%	Show in Genome Browser
chr2:87088946-87089038	CD8B,ANAPC1P1	0.91	91.39%	0%	0%	0%	0%	Show in Genome Browser
chr3:24536143-24536313	LOC644990,THRB	0.91	87.71%	0%	0%	0%	0%	Show in Genome Browser
chr19:50004042-50004125	MIR150	0.92	92.85%	0%	0%	0%	0%	Show in Genome Browser
chr16:22448326-22449036	SMG1P1,RRN3P3	0.92	65.32%	3.55%	0%	0%	0%	Show in Genome Browser
chr7:151106513-151107124	WDR86-AS1	0.93	47.87%	3.1%	0%	0%	0%	Show in Genome Browser
chr11:34937677-34937774	APIP,PDHX	0.94	94.89%	0%	0%	0%	0%	Show in Genome Browser
chr15:40542868-40545110	C15orf56,PAK6	0.94	32.45%	3.91%	0.37%	0%	0%	Show in Genome Browser
chr12:8340328-8353595	FAM66C	0.94	21.44%	9.57%	8.09%	2.32%	1.61%	Show in Genome Browser
chr15:31619083-31670101	KLF13	0.94	11.75%	3.99%	1.38%	0%	0%	Show in Genome Browser
chr6:28058929-28063492	ZSCAN12P1	0.94	19.65%	9.68%	7.35%	3.77%	0.07%	Show in Genome Browser
chr10:89603567-89605365	CFL1P1	0.95	28.84%	6.17%	0%	0%	0%	Show in Genome Browser
chrY:19691333-19691353	FAM224A	0.95	95.23%	0%	0%	0%	0%	Show in Genome Browser
chr20:1373478-1373525	FKBP1A	0.95	91.66%	0%	0%	0%	0%	Show in Genome Browser
chr11:62432779-62432804	C11orf48,METTL12	0.96	96.15%	0%	0%	0%	0%	Show in Genome Browser
chr4:8868773-8870071	HMX1	0.96	22.4%	9.16%	1.15%	0%	0%	Show in Genome Browser
chr11:28078362-28078457	KIF18A,MIR610	0.96	96.87%	0%	0%	0%	0%	Show in Genome Browser
chr3:103946073-103946105	MIR548A3	0.96	96.96%	0%	0%	0%	0%	Show in Genome

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chr4:185719451-185720200	SLED1,ACSL1	0.96	32.93%	9.73%	0%	0%	0%	Show in Genome Browser
chr2:220117965-220118638	TUBA4B,TUBA4A	0.96	25.01%	0%	0%	0%	0%	Show in Genome Browser
chr4:492845-493442	ZNF721,PIGG	0.96	40.92%	0%	0%	0%	0%	Show in Genome Browser
chr20:62374232-62375401	ZBTB46,SLC2A4RG	0.97	20.17%	14.18%	0%	0%	0%	Show in Genome Browser
chr12:6560684-6560883	CD27	0.98	85%	0%	0%	0%	0%	Show in Genome Browser
chr13:21277482-21297236	IL17D	0.98	27.33%	19.26%	13.82%	0.21%	0%	Show in Genome Browser
chr19:1852399-1854759	KLF16	0.98	24.94%	7.32%	4.61%	0%	0%	Show in Genome Browser
chrX:134555868-134560224	LINC00086	0.98	17.07%	3.61%	2.05%	0%	0%	Show in Genome Browser
chr19:518889-519653	TPGS1	0.98	31.5%	4.31%	0%	0%	0%	Show in Genome Browser
chr11:5617347-5617411	TRIM6	0.98	98.46%	0%	0%	0%	0%	Show in Genome Browser
chr12:56223273-56223420	DNAJC14,TMEM198B	1	99.32%	0%	0%	0%	0%	Show in Genome Browser
chrX:134229015-134232733	LINC00087	1	11.49%	4.7%	0.85%	0%	0%	Show in Genome Browser
chr7:98870924-98894540	MYH16	1	47.23%	4.33%	0%	0%	0%	Show in Genome Browser
chr13:27825692-27825722	RPL21,RPL21P28	1	70.96%	0%	0%	0%	0%	Show in Genome Browser
chrY:24442946-24445023	TTY5	1	33.06%	11.85%	0%	0%	0%	Show in Genome Browser
chr16:31446885-31454348	ZNF843	1	53.62%	11.97%	0.05%	0%	0%	Show in Genome Browser
chr3:183977003-183979251	CAMK2N2,ECE2	1.01	48.89%	3.09%	0%	0%	0%	Show in Genome Browser
chr9:89559279-89562104	GAS1	1.01	29.9%	7.67%	2.44%	0%	0%	Show in Genome Browser
chr22:50920012-50924864	ADM2	1.02	31.87%	14.23%	4.48%	0%	0%	Show in Genome Browser
chr12:50222326-50222531	BCDIN3D-AS1	1.02	57.28%	0%	0%	0%	0%	Show in Genome

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chr11:65657875-65659105	CCDC85B	1.02	57.92%	0%	0%	0%	0%	Show in Genome Browser
chr13:112721913-112726020	SOX1	1.02	16.69%	6.79%	5.72%	0%	0%	Show in Genome Browser
chr10:64564516-64568238	ADO	1.03	27.53%	9.83%	1.02%	0%	0%	Show in Genome Browser
chr19:13261282-13265716	IER2	1.03	54.62%	19.23%	2.64%	0%	0%	Show in Genome Browser
chr9:140194083-140196703	NRARP	1.03	24.07%	9.08%	2.74%	0%	0%	Show in Genome Browser
chr6:27325602-27343153	ZNF204P	1.04	42.33%	21.31%	7.52%	0.19%	0%	Show in Genome Browser
chr19:1009172-1009649	GRIN3B,C19orf6	1.05	44.14%	0%	0%	0%	0%	Show in Genome Browser
chr5:148808587-148810208	MIR145	1.05	44.08%	6.96%	1.1%	0%	0%	Show in Genome Browser
chr4:140222676-140222985	NDUFC1,NAA15	1.05	50.64%	9.67%	0%	0%	0%	Show in Genome Browser
chr12:47602203-47610047	PCED1B-AS1	1.05	24.58%	0%	0%	0%	0%	Show in Genome Browser
chr4:139085248-139093173	SLC7A11,SLC7A11-AS1	1.06	6.18%	3.41%	2.81%	2.23%	1.48%	Show in Genome Browser
chr17:655900-658576	DBIL5P	1.07	28.42%	9.93%	0%	0%	0%	Show in Genome Browser
chr5:138609778-138609791	SNHG4,MATR3	1.07	92.85%	0%	0%	0%	0%	Show in Genome Browser
chr7:150076406-150095719	ZNF775	1.07	63.3%	31.32%	5.41%	0%	0%	Show in Genome Browser
chr1:94312388-94312467	BCAR3,MIR760	1.08	57.49%	0%	0%	0%	0%	Show in Genome Browser
chr1:179712298-179785326	FAM163A	1.08	36.99%	19.82%	18.42%	5.63%	1.68%	Show in Genome Browser
chr17:7292290-7293760	C17orf61-PLSCR3,TNK1	1.09	41.11%	6.06%	2.37%	0%	0%	Show in Genome Browser
chr16:30016542-30016836	INO80E,DOC2A	1.09	83.38%	0%	0%	0%	0%	Show in Genome Browser
chr7:39773167-39834221	LINC00265	1.09	68.72%	40.72%	18.71%	2.17%	0.79%	Show in Genome Browser
chr19:4639530-4653952	TNFAIP8L1	1.09	44.35%	10.11%	7.55%	3.47%	0%	Show in Genome Browser

chr19:2250651-2250759	AMH,MIR4321	1.1	96.33%	0%	0%	0%	0%	Show in Genome Browser
chr1:26485511-26489118	FAM110D	1.1	38.38%	12.02%	0%	0%	0%	Show in Genome Browser
chr22:24823530-24823714	ADORA2A	1.11	73.51%	0%	0%	0%	0%	Show in Genome Browser
chr9:140063723-140064491	LRRC26	1.11	72.43%	0%	0%	0%	0%	Show in Genome Browser
chr14:31483852-31483948	MIR624,STRN3	1.11	94.84%	0%	0%	0%	0%	Show in Genome Browser
chr22:46508703-46509565	MIRLET7BHG,MIR4763	1.11	22.01%	10.08%	4.28%	0%	0%	Show in Genome Browser
chr6:32116141-32119720	PRRT1	1.11	38.9%	2.89%	0%	0%	0%	Show in Genome Browser
chr22:19033675-19035888	DGCR11,DGCR2	1.12	11.87%	7.67%	3.79%	1.85%	0.81%	Show in Genome Browser
chr14:24778161-24779854	LTB4R2,CIDEA	1.12	32.29%	6.37%	3.54%	0%	0%	Show in Genome Browser
chr16:69363900-69364498	PDF,COG8	1.12	55.92%	1%	0%	0%	0%	Show in Genome Browser
chr1:84830648-84850494	UOX	1.12	36.23%	0.15%	0%	0%	0%	Show in Genome Browser
chr17:6921230-6921341	MIR497	1.13	67.85%	2.67%	0%	0%	0%	Show in Genome Browser
chr11:117947727-117948020	TMPRSS4-AS1	1.13	36.73%	2.72%	0%	0%	0%	Show in Genome Browser
chr7:29603427-29606911	PRR15	1.14	47.26%	14.19%	0%	0%	0%	Show in Genome Browser
chr20:62584737-62587828	UCKL1-AS1	1.14	37.04%	6.23%	0%	0%	0%	Show in Genome Browser
chr14:102973198-102974279	ANKRD9	1.15	30.03%	13.77%	0.64%	0%	0%	Show in Genome Browser
chr2:6072819-6120350	LOC150622	1.15	15.66%	5.39%	5.03%	4.17%	3.5%	Show in Genome Browser
chr19:49534926-49535044	SNAR-G2	1.15	99.15%	0%	0%	0%	0%	Show in Genome Browser
chr16:30418735-30429915	ZNF771	1.15	43.86%	21.93%	0%	0%	0%	Show in Genome Browser
chr19:47251284-47261832	FKRP	1.16	49.64%	20.34%	5.11%	0%	0%	Show in Genome Browser

chr17:47287589-47288203	GNGT2,ABI3	1.16	60.97%	0%	0%	0%	0%	Show in Genome Browser
chr11:88237745-88242672	GRM5,GRM5-AS1	1.16	18.38%	8.58%	5.31%	0.36%	0%	Show in Genome Browser
chr8:127564687-127570466	FAM84B	1.17	23.76%	20.7%	10.2%	1.23%	0%	Show in Genome Browser
chr20:4702556-4709105	PRND	1.17	30.84%	15.15%	11.59%	2.18%	0%	Show in Genome Browser
chr22:19711103-19711376	GP1BB,SEPT5-GP1BB	1.18	47.44%	0%	0%	0%	0%	Show in Genome Browser
chr6:29942892-29945494	HCG9	1.18	77.01%	7.07%	0%	0%	0%	Show in Genome Browser
chr7:2566780-2568061	LFNG,MIR4648	1.18	40.79%	6.94%	3.12%	0%	0%	Show in Genome Browser
chr22:46505551-46508628	MIRLET7BHG,MIRLET7A3	1.18	34.14%	11.82%	0.48%	0%	0%	Show in Genome Browser
chr16:70284134-70285833	EXOSC6	1.2	38.41%	7.05%	0%	0%	0%	Show in Genome Browser
chr17:38516907-38520945	GJD3	1.2	40.18%	9.23%	3.09%	0%	0%	Show in Genome Browser
chr19:18390563-18392432	JUND	1.2	37.8%	7.86%	2.03%	0%	0%	Show in Genome Browser
chr2:78182033-78182152	SNAR-H	1.2	95%	0%	0%	0%	0%	Show in Genome Browser
chr4:4420785-4421856	STX18,NSG1	1.2	25.37%	11%	1.02%	0%	0%	Show in Genome Browser
chr6:31165537-31171744	HCG27	1.21	36.95%	19.9%	7.51%	0%	0%	Show in Genome Browser
chr15:40824083-40824749	MRPL42P5	1.21	65.36%	0%	0%	0%	0%	Show in Genome Browser
chr13:77454304-77460540	KCTD12	1.22	20.47%	11.2%	5.32%	0%	0%	Show in Genome Browser
chrX:73012040-73048902	XIST,TSIX	1.22	6.07%	0.88%	0.34%	0.14%	0%	Show in Genome Browser
chr11:2155364-2155439	IGF2,INS-IGF2,MIR483	1.23	96.05%	0%	0%	0%	0%	Show in Genome Browser
chr1:236686739-236687361	LGALS8,LGALS8-AS1	1.23	50.71%	0%	0%	0%	0%	Show in Genome Browser
chr6:32812056-32812712	LOC100507463,PSMB8	1.23	29.83%	13.69%	0%	0%	0%	Show in Genome Browser
								Show in

chr7:29685539-29690955	LOC646762	1.23	21%	8.82%	3.96%	1.18%	0.23%	Genome Browser
chr14:101340830-101340920	MIR337	1.23	98.9%	0%	0%	0%	0%	Show in Genome Browser
chr4:69078024-69078079	TMPRSS11BNL,FTLP10	1.23	98.21%	0%	0%	0%	0%	Show in Genome Browser
chr17:19140690-19140844	EPN2	1.25	61.29%	0%	0%	0%	0%	Show in Genome Browser
chr19:1823571-1823784	LOC100288123,REXO1	1.25	81.77%	0%	0%	0%	0%	Show in Genome Browser
chr7:102105390-102105607	LRWD1	1.25	54.12%	0%	0%	0%	0%	Show in Genome Browser
chr17:7517382-7518215	SHBG,FXR2	1.25	61.67%	0%	0%	0%	0%	Show in Genome Browser
chr3:129033615-129043412	H1FX-AS1	1.26	40.53%	13.61%	0%	0%	0%	Show in Genome Browser
chr19:1905373-1913443	SCAMP4,ADAT3	1.27	56.62%	18.9%	10.76%	0%	0%	Show in Genome Browser
chr14:104394817-104395061	TDRD9	1.28	76.32%	0%	0%	0%	0%	Show in Genome Browser
chr10:101491717-101492423	COX15,CUTC	1.29	46.38%	17.06%	0%	0%	0%	Show in Genome Browser
chr21:27107164-27107328	GABPA,ATP5J	1.29	54.54%	0%	0%	0%	0%	Show in Genome Browser
chr10:65224989-65225254	LOC84989,JMJD1C	1.29	84.21%	0%	0%	0%	0%	Show in Genome Browser
chr1:101003728-101007582	GPR88	1.3	74.21%	8.14%	0%	0%	0%	Show in Genome Browser
chr1:1361508-1363166	TMEM88B	1.3	64.16%	3.87%	0%	0%	0%	Show in Genome Browser
chr19:58878991-58892389	ZNF837	1.3	66.47%	14.23%	0.63%	0%	0%	Show in Genome Browser
chr11:2904450-2906696	CDKN1C	1.31	54.65%	0%	0%	0%	0%	Show in Genome Browser
chr19:48259110-48259218	GLTSCR2,SNORD23	1.31	99.08%	0%	0%	0%	0%	Show in Genome Browser
chr9:34568010-34583070	LOC415056,CNTFR	1.31	28.52%	0%	0%	0%	0%	Show in Genome Browser
chr6:5085720-5087455	PPP1R3G	1.31	57.83%	4.78%	0%	0%	0%	Show in Genome Browser
chr22:38239918-								Show in

38240378	ANKRD54,MIR658	1.32	82.3%	0%	0%	0%	0%	Genome Browser
chr17:48638449-48639062	CACNA1G,CACNA1G-AS1	1.32	31.92%	9.28%	0%	0%	0%	Show in Genome Browser
chr16:20817730-20817766	ERL2,LOC81691	1.32	97.29%	0%	0%	0%	0%	Show in Genome Browser
chr19:2511218-2646278	GNG7	1.32	38.74%	18.33%	11.95%	3.93%	2.77%	Show in Genome Browser
chr4:57976043-57976539	LOC255130,IGFBP7	1.32	58.75%	3.42%	0%	0%	0%	Show in Genome Browser
chr7:77976559-77988768	MAGI2,RPL13AP17	1.32	45.44%	28.7%	11.34%	0.06%	0%	Show in Genome Browser
chr17:49242891-49244187	NME2	1.32	36.61%	8.57%	0%	0%	0%	Show in Genome Browser
chr2:63815331-63815851	WDPCP,MDH1	1.32	59.88%	6.14%	0%	0%	0%	Show in Genome Browser
chr22:22049683-22055507	YPEL1,PPIL2	1.32	30.25%	6.2%	4.28%	2.67%	1.06%	Show in Genome Browser
chr3:10326103-10327429	GHRLOS2	1.33	41.07%	6.7%	4.52%	2.03%	0%	Show in Genome Browser
chr9:95381768-95381847	IPPK,LOC100128361	1.33	98.75%	0%	0%	0%	0%	Show in Genome Browser
chr9:94903749-94921890	LINC00475	1.33	33.72%	4.71%	0%	0%	0%	Show in Genome Browser
chr9:127115745-127121461	LOC100129034,PSMB7	1.34	26.44%	8.88%	5.95%	2.16%	0.41%	Show in Genome Browser
chr5:112357797-112363179	MCC,DCP2	1.34	12.11%	4.81%	4.45%	3.15%	1.74%	Show in Genome Browser
chr12:6728001-6745297	LPAR5	1.35	63.1%	23.16%	10.9%	0%	0%	Show in Genome Browser
chr8:30241944-30242674	RBPM5,LOC100128750	1.35	52.53%	2.18%	0%	0%	0%	Show in Genome Browser
chr22:25160468-25161984	PIWIL3,TOP1P2	1.36	30.32%	11.2%	4.61%	0%	0%	Show in Genome Browser
chr4:24797085-24802466	SOD3	1.36	65.84%	17.28%	3.67%	0%	0%	Show in Genome Browser
chr19:5904852-5910261	VMAC	1.36	43.7%	9.18%	7.92%	3.89%	0%	Show in Genome Browser
chr7:100860447-100860984	ZNHIT1,PLOD3	1.36	75.46%	6.13%	0.74%	0%	0%	Show in Genome Browser
chr5:149380218-								Show in

149380742	HMGXB3,TIGD6	1.37	74.66%	0%	0%	0%	0%	Genome Browser
chr17:29858634-29865232	RAB11FIP4,MIR4724	1.37	18.71%	4.15%	3.66%	3.07%	2.31%	Show in Genome Browser
chr21:19161285-19165898	C21orf91,C21orf91-OT1	1.38	12.63%	6.24%	5.87%	2.6%	1.75%	Show in Genome Browser
chr16:75507023-75528926	CHST6	1.38	29.68%	16.9%	13.35%	4.94%	0.01%	Show in Genome Browser
chr7:102086960-102096696	ALKBH4,ORA2	1.39	21.02%	5.1%	4.22%	3.14%	1.45%	Show in Genome Browser
chr20:45947246-45949497	LOC100131496,ZMYND8	1.39	42.8%	8.74%	3.99%	0%	0%	Show in Genome Browser
chr5:36147994-36148090	MIR580,LMBRD2	1.39	74.22%	0%	0%	0%	0%	Show in Genome Browser
chr5:172068276-172118531	NEURL1B	1.39	19.75%	1.94%	0%	0%	0%	Show in Genome Browser
chr19:54976210-54984422	CDC42EP5	1.4	78.63%	9.84%	0%	0%	0%	Show in Genome Browser
chr18:14337422-14342522	CYP4F35P	1.4	44.39%	12.24%	4.59%	3.53%	0.58%	Show in Genome Browser
chr2:219845809-219850379	FEV	1.4	35.57%	10.65%	5.38%	0%	0%	Show in Genome Browser
chr5:87960264-87963409	LINC00461,MIR9-2	1.4	16.49%	9.63%	5.24%	2.03%	0%	Show in Genome Browser
chr3:100119470-100121283	LNP1,TOMM70A	1.4	37.65%	9.47%	0%	0%	0%	Show in Genome Browser
chr7:129414535-129414609	MIR96	1.4	98.66%	0%	0%	0%	0%	Show in Genome Browser
chr17:37760022-37764175	NEUROD2	1.4	37.09%	23.43%	9.96%	1.39%	0%	Show in Genome Browser
chr1:207991724-207995940	LOC148696	1.42	23.16%	8.15%	6.26%	2.32%	0%	Show in Genome Browser
chr14:101346992-101347343	RTL1,MIR431	1.42	90.05%	0%	0%	0%	0%	Show in Genome Browser
chr15:25200070-25213974	SNURF,SNRPN	1.42	65.64%	0%	0%	0%	0%	Show in Genome Browser
chr21:33458517-33458608	LINC00159	1.43	98.91%	0%	0%	0%	0%	Show in Genome Browser
chr1:238643686-238644981	LOC339535	1.43	25.07%	12.8%	6.09%	0%	0%	Show in Genome Browser
chr19:10928102-10928172	MIR199A1	1.43	98.59%	0%	0%	0%	0%	Show in Genome

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chr3:132440807-132593050	NPHP3-AS1	1.43	24.31%	11.94%	6.17%	0%	0%	Show in Genome Browser
chr6:21593972-21598847	SOX4	1.43	31.66%	11.54%	3.07%	1.25%	0%	Show in Genome Browser
chr10:119243805-119303887	EMX2OS	1.44	14.16%	3.07%	2.26%	1.43%	0.93%	Show in Genome Browser
chr11:62453876-62457200	LRRN4CL	1.44	39.42%	19.61%	7.67%	0%	0%	Show in Genome Browser
chr12:6993145-6993766	RPL13P5	1.44	79.42%	0%	0%	0%	0%	Show in Genome Browser
chr22:19956059-19958266	ARVCF,COMT	1.45	39.91%	9.71%	7.97%	0%	0%	Show in Genome Browser
chr19:917342-921014	KISS1R	1.45	82.27%	9.5%	5.55%	0%	0%	Show in Genome Browser
chr6:33175612-33175721	MIR219-1	1.45	95.45%	0%	0%	0%	0%	Show in Genome Browser
chr7:128281295-128301052	FLJ45340	1.46	61.29%	31.85%	13.81%	3.72%	1.12%	Show in Genome Browser
chr17:39202798-39203568	KRTAP2-1	1.46	59.79%	6.22%	0%	0%	0%	Show in Genome Browser
chr17:10725793-10741418	PIRT	1.46	20.29%	14.69%	11.16%	5.74%	2.88%	Show in Genome Browser
chr4:7194374-7194853	SORCS2	1.46	29.37%	18.75%	0%	0%	0%	Show in Genome Browser
chr5:114937794-114952014	TMED7-TICAM2	1.46	14.42%	9.13%	6.86%	5.39%	4.64%	Show in Genome Browser
chr4:16228286-16259810	FLJ39653	1.47	17.46%	13.41%	10.83%	6.01%	0%	Show in Genome Browser
chr22:50964184-50964905	SCO2,TYMP	1.47	75.38%	0%	0%	0%	0%	Show in Genome Browser
chr21:42539728-42540502	BACE2	1.48	48.25%	12.77%	0%	0%	0%	Show in Genome Browser
chr21:26934457-26947473	MIR155HG	1.48	50.1%	16.38%	12.28%	4.77%	0%	Show in Genome Browser
chr15:83735503-83736106	BTBD1,MIR4515	1.49	75%	0%	0%	0%	0%	Show in Genome Browser
chr10:99079022-99081672	FRAT1	1.49	48.09%	9.88%	2.18%	0%	0%	Show in Genome Browser
chr12:121157723-121161442	MIR4700,UNC119B	1.49	17.63%	6.18%	4.59%	3.09%	2.36%	Show in Genome Browser

chrX:12993225-12993287	TMSB4X	1.49	98.41%	0%	0%	0%	0%	Show in Genome Browser
chr1:43735665-43739672	TMEM125	1.5	60.48%	33.38%	17.16%	0%	0%	Show in Genome Browser
chr2:85770793-85772401	GGCX,MAT2A	1.51	32.38%	8.45%	6.33%	0%	0%	Show in Genome Browser
chr2:178148236-178257419	LOC100130691	1.51	24.37%	17.93%	15.61%	12.04%	10.81%	Show in Genome Browser
chr17:43697712-43715327	MGC57346,CRHR1	1.52	23.66%	11.68%	9.31%	7.08%	5.48%	Show in Genome Browser
chr17:74261286-74267379	FAM100B	1.53	54.73%	5.03%	0%	0%	0%	Show in Genome Browser
chr5:180017106-180018487	SCGB3A1	1.53	66.77%	22.2%	0%	0%	0%	Show in Genome Browser
chr14:99976494-99981672	CCDC85C,CCNK	1.54	23.92%	6.65%	0.75%	0%	0%	Show in Genome Browser
chr11:66059373-66064133	TMEM151A	1.54	65.01%	18.12%	6.8%	1.85%	0%	Show in Genome Browser
chr17:15902782-15902832	ZSWIM7,TTC19	1.54	98.03%	0%	0%	0%	0%	Show in Genome Browser
chr17:54961463-54962274	MTVR2	1.55	57.63%	13.54%	0%	0%	0%	Show in Genome Browser
chr19:1753662-1775443	ONECUT3	1.55	27.06%	2.68%	0%	0%	0%	Show in Genome Browser
chr5:92953432-92956835	FAM172A,MIR548AO,MIR2277	1.56	18.18%	8.6%	7.13%	2.64%	0.88%	Show in Genome Browser
chr1:209848670-209849734	G0S2	1.56	69.06%	20.49%	0%	0%	0%	Show in Genome Browser
chr14:37641486-37641865	SLC25A21,LOC100129794	1.56	56.05%	10%	0%	0%	0%	Show in Genome Browser
chr17:39221369-39222131	KRTAP2-4	1.57	78.89%	0%	0%	0%	0%	Show in Genome Browser
chr18:657604-658340	TYMS,C18orf56	1.57	60.4%	19.03%	0%	0%	0%	Show in Genome Browser
chr19:56088892-56092211	ZNF579	1.57	60.62%	26.41%	9.64%	0%	0%	Show in Genome Browser
chr6:31514255-31514627	ATP6V1G2-DDX39B,NFKBIL1	1.58	48.25%	4.55%	0%	0%	0%	Show in Genome Browser
chr1:231950373-231954074	DISC1,DISC2	1.58	19.93%	6.18%	3.45%	2.56%	1.86%	Show in Genome Browser
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chr4:144480625-144482612	GUSBP5	1.58	44.91%	15.84%	0%	0%	0%	Genome Browser
chr19:40129325-40133041	LOC100129935	1.58	33.47%	3.72%	0.74%	0%	0%	Show in Genome Browser
chr19:33166313-33169206	RGS9BP	1.58	40.49%	13.06%	3.35%	0%	0%	Show in Genome Browser
chr1:28835418-28836145	SNHG3,RCC1	1.58	35.85%	8.65%	5.35%	0%	0%	Show in Genome Browser
chr12:53574535-53574621	ZNF740,CSAD	1.58	88.5%	0%	0%	0%	0%	Show in Genome Browser
chr4:89647106-89651254	FAM13A,FAM13A-AS1	1.59	17.63%	5.96%	3.76%	2.49%	1.8%	Show in Genome Browser
chr20:4711929-4721314	PRNT	1.59	40.36%	17.02%	10.5%	1.5%	0%	Show in Genome Browser
chr8:65492814-65496185	BHLHE22	1.6	39.62%	12.78%	4.5%	0%	0%	Show in Genome Browser
chr17:74135979-74137380	RNF157-AS1,FOXJ1	1.6	68.73%	19.47%	9.15%	0%	0%	Show in Genome Browser
chr8:58907113-59062277	FAM110B	1.61	37.78%	32.23%	26.14%	12.96%	7.7%	Show in Genome Browser
chr21:34398239-34401500	OLIG2	1.61	40.94%	18.41%	4.78%	1.48%	0%	Show in Genome Browser
chr20:306239-310865	SOX12	1.61	36.11%	11.99%	5.96%	0.79%	0%	Show in Genome Browser
chr17:79276624-79283048	LINC00482	1.62	55.61%	26.43%	9.6%	0.54%	0%	Show in Genome Browser
chr18:72259012-72265071	LOC400657	1.62	17.94%	8.15%	5.44%	2.3%	0%	Show in Genome Browser
chr2:223162866-223163700	PAX3,CCDC140	1.62	58.57%	20.52%	0%	0%	0%	Show in Genome Browser
chr14:105995059-105996539	TMEM121	1.62	46.18%	16.34%	2.43%	0%	0%	Show in Genome Browser
chr1:240652875-240775462	GREM2	1.63	35.15%	13.23%	12.51%	5.76%	0%	Show in Genome Browser
chr6:31555418-31555743	LST1	1.63	80.07%	0%	0%	0%	0%	Show in Genome Browser
chr6:146205947-146207576	SHPRH,LOC100507557	1.63	27.91%	8.95%	5.76%	2.39%	0%	Show in Genome Browser
chr21:37832920-37948867	CLDN14	1.64	55.4%	29.64%	19.82%	7.93%	0.13%	Show in Genome Browser

chr1:20878932-20881512	FAM43B	1.64	48.39%	15.26%	0.38%	0%	0%	Show in Genome Browser
chr19:59092612-59095763	MGC2752	1.64	34.23%	10.34%	6.59%	0.47%	0%	Show in Genome Browser
chr6:30036881-30038977	RNF39,PPP1R11	1.64	38.54%	15.68%	0%	0%	0%	Show in Genome Browser
chr15:75247443-75249775	RPP25	1.64	41.92%	14.83%	2.74%	0%	0%	Show in Genome Browser
chr16:1559831-1561151	TELO2,IFT140	1.64	74.41%	10.39%	0%	0%	0%	Show in Genome Browser
chr12:111798485-111806925	FAM109A	1.65	60.68%	19.91%	6.61%	2.2%	0%	Show in Genome Browser
chr12:81226312-81226408	LIN7A,MIR617	1.65	98.96%	0%	0%	0%	0%	Show in Genome Browser
chr20:48807376-48809212	CEBPB	1.66	51.6%	11.15%	0%	0%	0%	Show in Genome Browser
chr6:1312675-1314992	FOXQ1	1.66	35.46%	10.31%	8.84%	0%	0%	Show in Genome Browser
chr3:120113062-120114514	MIR198,FSTL1	1.66	19.61%	11.01%	5.5%	3.02%	1.03%	Show in Genome Browser
chr2:239146910-239148681	HES6	1.67	34.99%	23.79%	17.43%	6.44%	0%	Show in Genome Browser
chr11:74165888-74178600	KCNE3	1.67	33.66%	13.89%	12.78%	8.15%	4.66%	Show in Genome Browser
chr11:35639735-35642419	FJX1	1.68	45.84%	15.23%	0.81%	0%	0%	Show in Genome Browser
chr13:20712395-20735183	GJA3	1.68	27.86%	14%	11.39%	7.12%	4.3%	Show in Genome Browser
chrX:1581466-1656037	P2RY8	1.68	43.04%	26.57%	13.53%	3.64%	0%	Show in Genome Browser
chr1:45805788-45806142	TOE1,MUTYH	1.68	80.64%	0%	0%	0%	0%	Show in Genome Browser
chr5:60453536-60458302	C5orf43	1.69	19.55%	14.54%	13.77%	9.9%	8%	Show in Genome Browser
chr2:233734994-233741107	C2orf82	1.7	89.88%	6.67%	0%	0%	0%	Show in Genome Browser
chr5:179105559-179107975	CBY3	1.7	71.43%	28.18%	7.17%	0%	0%	Show in Genome Browser
chr21:44869904-44872385	LINC00319	1.7	37.19%	14.52%	0.22%	0%	0%	Show in Genome Browser
chr17:7816641-								Show in

7819265	LOC284023	1.7	28.35%	7.6%	4.63%	1.78%	0%	Genome Browser
chr6:32723876-32731330	HLA-DQB2	1.71	55.69%	14.37%	5.18%	0%	0%	Show in Genome Browser
chr22:23732793-23744799	ZDHC8P1	1.71	43.29%	27.87%	16.15%	0%	0%	Show in Genome Browser
chr11:45792983-45793908	DKFZp779M0652	1.72	75.3%	12.04%	0%	0%	0%	Show in Genome Browser
chr1:160336861-160342638	NHLH1	1.72	49.72%	18.13%	12.97%	5.9%	0%	Show in Genome Browser
chrY:21238931-21239302	TTY14	1.72	82.52%	1.07%	0%	0%	0%	Show in Genome Browser
chr19:56132107-56135941	ZNF784	1.72	60.7%	7.57%	2.75%	0%	0%	Show in Genome Browser
chr15:41062159-41064645	C15orf62,DNAJC17	1.73	42.01%	18.49%	1.8%	0%	0%	Show in Genome Browser
chr16:1398380-1399242	BAIAP3,TSR3	1.74	45.3%	16.57%	0%	0%	0%	Show in Genome Browser
chr5:172381788-172386371	LOC100268168	1.74	9.68%	3.72%	0%	0%	0%	Show in Genome Browser
chr11:64658609-64658718	MIR192	1.74	99.09%	0%	0%	0%	0%	Show in Genome Browser
chr5:132149033-132152488	SOWAHA	1.74	39.32%	11.92%	5.52%	1.04%	0%	Show in Genome Browser
chr14:34902145-34931468	SPTSSA	1.74	27.25%	6.12%	5.01%	3.63%	0.5%	Show in Genome Browser
chr19:1473244-1479228	C19orf25	1.75	42.33%	21.74%	1.89%	0%	0%	Show in Genome Browser
chr1:2520849-2522080	FAM213B,MME1	1.75	45.12%	9.74%	7.06%	1.7%	0%	Show in Genome Browser
chr10:87359312-87362458	GRID1,GRID1-AS1	1.75	25.92%	13.21%	9.31%	0.31%	0%	Show in Genome Browser
chr21:35818989-35883613	KCNE1	1.75	31.5%	17.05%	15.89%	12.59%	8.75%	Show in Genome Browser
chr17:79876147-79881408	MAFG	1.75	22.06%	3.7%	0%	0%	0%	Show in Genome Browser
chr17:79897521-79900249	MYADML2	1.75	53.04%	21.53%	2.59%	0%	0%	Show in Genome Browser
chr4:13542454-13544152	NKX3-2	1.75	48.79%	18.01%	1.29%	0%	0%	Show in Genome Browser
chr20:2633254-	NOP56	1.75	98.21%	0%	0%	0%	0%	Show in Genome Browser

2633309								Browser
chr12:98906753-98910004	TMPO-AS1	1.75	10.23%	5.64%	1.33%	0%	0%	Show in Genome Browser
chr7:19155093-19157295	TWIST1	1.75	32.94%	18.45%	10.34%	0%	0%	Show in Genome Browser
chr6:4079440-4079914	C6orf201,FAM217A	1.76	48.21%	22.1%	0%	0%	0%	Show in Genome Browser
chr2:233250461-233251754	ECEL1P2	1.76	57.34%	15.91%	4.25%	0%	0%	Show in Genome Browser
chrY:9321930-9322075	FAM197Y2	1.76	99.31%	0%	0%	0%	0%	Show in Genome Browser
chr13:113622757-113623028	MCF2L-AS1	1.76	73.89%	3.3%	0%	0%	0%	Show in Genome Browser
chr15:81293295-81296344	MESDC1	1.77	38.62%	19.73%	2.06%	0%	0%	Show in Genome Browser
chr21:46839631-46844985	COL18A1-AS1	1.78	28.52%	7.24%	5.93%	4.51%	0.78%	Show in Genome Browser
chr2:208627312-208634143	FZD5	1.78	37.41%	27.13%	15.39%	3.93%	0.71%	Show in Genome Browser
chr10:38726472-38741080	LOC399744	1.78	64.72%	41.33%	19.85%	5.31%	0.24%	Show in Genome Browser
chr12:54624732-54635689	CBX5,MIR3198-2	1.8	17.32%	3.53%	3%	2.65%	2.45%	Show in Genome Browser
chr14:91698877-91720224	GPR68	1.8	60.17%	28.45%	11.59%	1.03%	0%	Show in Genome Browser
chr4:7032054-7034844	LOC100129931,TBC1D14	1.8	15.9%	8.77%	7.41%	3.26%	2.29%	Show in Genome Browser
chr4:183060159-183065668	MGC45800	1.8	29.95%	9.84%	7.5%	5.4%	3.51%	Show in Genome Browser
chr9:125026994-125027143	MRRF,RBM18	1.8	85.33%	0%	0%	0%	0%	Show in Genome Browser
chr2:239756726-239757375	TWIST2	1.8	77.07%	10.92%	0%	0%	0%	Show in Genome Browser
chr8:143761875-143764142	PSCA	1.81	71.88%	17.59%	0%	0%	0%	Show in Genome Browser
chr22:19702304-19702314	SEPT5	1.81	90.9%	0%	0%	0%	0%	Show in Genome Browser
chr9:32552432-32552601	TOPORS,LOC100129250	1.81	99.41%	0%	0%	0%	0%	Show in Genome Browser
chr17:27940372-27941773	ANKRD13B,CORO6	1.82	42.08%	17.97%	0%	0%	0%	Show in Genome

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chr9:131937831-131940540	IER5L	1.82	41.18%	14.57%	4.28%	0%	0%	Show in Genome Browser
chr19:1465154-1473200	APC2,C19orf25	1.83	57.01%	13.75%	2.49%	0%	0%	Show in Genome Browser
chr1:11900200-11903200	CLCN6,NPPA-AS1	1.83	23.15%	6.89%	5.66%	4.16%	2.93%	Show in Genome Browser
chr7:99724991-99726119	MBLAC1	1.83	43.75%	16.91%	0%	0%	0%	Show in Genome Browser
chr4:69402905-69403132	UGT2B17	1.83	47.36%	18.42%	2.63%	0%	0%	Show in Genome Browser
chrX:46433192-46457837	CHST7	1.84	51.83%	19.61%	9%	1.68%	0%	Show in Genome Browser
chr18:24486446-24515910	CHST9-AS1	1.85	14.15%	10.42%	9.74%	9.38%	9.15%	Show in Genome Browser
chrX:49216634-49216647	GAGE2A,GAGE8,GAGE2C	1.85	92.85%	0%	0%	0%	0%	Show in Genome Browser
chr1:248100493-248100686	OR2L13	1.85	86.08%	0%	0%	0%	0%	Show in Genome Browser
chr10:5931004-5931860	ANKRD16,FBXO18	1.86	37.57%	24.97%	0%	0%	0%	Show in Genome Browser
chr9:130965663-130966662	DNM1,CIZ1	1.86	79.15%	9.42%	0%	0%	0%	Show in Genome Browser
chr11:3239176-3244361	MIR681-AS1	1.87	19.17%	6.16%	0.5%	0%	0%	Show in Genome Browser
chr5:79950784-79950800	DHFR,MSH3	1.88	94.11%	0%	0%	0%	0%	Show in Genome Browser
chr16:5083274-5083936	NAGPA,NAGPA-AS1	1.88	87.58%	18.49%	0%	0%	0%	Show in Genome Browser
chr19:5823818-5828334	NRTN	1.88	56.84%	10.87%	0%	0%	0%	Show in Genome Browser
chr3:152880029-152886261	RAP2B	1.88	19.15%	10.3%	7.09%	5.23%	0.19%	Show in Genome Browser
chr17:27400528-27402627	TIAF1,MYO18A	1.88	37.74%	18.01%	8.36%	3.82%	0%	Show in Genome Browser
chr1:29446559-29450407	TMEM200B	1.88	62.67%	23.02%	14.17%	4.72%	0%	Show in Genome Browser
chr1:104108057-104108216	AMY2B	1.89	99.37%	0%	0%	0%	0%	Show in Genome Browser
chr4:2061587-2067635	NAT8L	1.89	47.64%	14.89%	4.74%	2.26%	0%	Show in Genome Browser

								Browser
chr8:145618184-145618580	ADCK5,CPSF1	1.92	77.8%	0%	0%	0%	0%	Show in Genome Browser
chr19:14640355-14640381	MIR639,TECR	1.92	96.29%	0%	0%	0%	0%	Show in Genome Browser
chr12:110171914-110211292	FAM222A-AS1	1.93	52.43%	26.78%	12.17%	1.2%	0%	Show in Genome Browser
chr19:44116253-44118649	SRRM5,ZNF428	1.93	38.13%	7.03%	5.7%	3.83%	0.73%	Show in Genome Browser
chr6:96463845-96663487	FUT9	1.94	14.3%	10.57%	10.28%	9.44%	8.57%	Show in Genome Browser
chr22:42348191-42354945	LINC00634	1.94	32.9%	21.61%	15.75%	3.06%	0%	Show in Genome Browser
chr6:32812987-32813562	LOC100507463,TAP1	1.94	68.57%	15.1%	0%	0%	0%	Show in Genome Browser
chr1:148560990-148566880	NBPF15	1.94	80%	24.96%	0%	0%	0%	Show in Genome Browser
chr2:113191108-113191222	RGPD8	1.94	95.65%	0%	0%	0%	0%	Show in Genome Browser
chr20:61637331-61638387	BHLHE23	1.95	61.4%	13.62%	2.74%	0%	0%	Show in Genome Browser
chr9:139006428-139008870	C9orf69	1.95	33.6%	12.81%	10.19%	0.69%	0%	Show in Genome Browser
chr19:2233155-2233904	PLEKHJ1	1.95	44.53%	11.06%	7.06%	0.4%	0%	Show in Genome Browser
chr6:31803040-31803103	C6orf48,SNORD48	1.96	42.18%	17.18%	0%	0%	0%	Show in Genome Browser
chr19:859665-863608	CFD	1.96	69.2%	16.43%	0%	0%	0%	Show in Genome Browser
chr17:39210753-39211463	KRTAP2-2	1.96	75.94%	11.67%	0%	0%	0%	Show in Genome Browser
chr17:79099684-79099749	MIR338,MIR3065	1.96	98.48%	0%	0%	0%	0%	Show in Genome Browser
chr17:55122839-55124155	RNF126P1	1.96	85.42%	11.23%	0%	0%	0%	Show in Genome Browser
chrX:152710179-152711945	TREX2	1.96	56.71%	23.92%	0%	0%	0%	Show in Genome Browser
chr17:79976580-79980773	STRA13	1.97	51.93%	6.57%	0%	0%	0%	Show in Genome Browser
chr7:7679951-7758238	RPA3,LOC729852	1.98	32.81%	19.95%	16.78%	11.82%	8.73%	Show in Genome Browser

chr19:5915232-5917664	CAPS,RANBP3	1.99	32.68%	16.53%	10.65%	0%	0%	Show in Genome Browser
chr3:151488244-151502682	MIR548H2,LOC201651	1.99	55.9%	32.92%	15.51%	0%	0%	Show in Genome Browser
chr20:21376007-21378047	NKX2-4	1.99	52.94%	21.34%	5.69%	0%	0%	Show in Genome Browser
chr5:92919043-92921192	NR2F1	1.99	29.48%	12.27%	8.09%	3.9%	0%	Show in Genome Browser
chr16:69966984-69967083	WWP2,MIR140	1.99	99%	0%	0%	0%	0%	Show in Genome Browser

Highly Covered Genes (Cutoff: 200x)

Nothing found for this cutoff.

BED file used for specification of target regions:
/gpfs/scratch/project/EPIGEN/db/62/TruSeq_exome_hg19.bed