

CpG	Chr	Nt. pos.	P	Rank	Distance to TSS	Genes
cg16351002	10	124600091	8.13E-06	1	208	<i>CUZD1</i>
cg11378686	6	43444256	4.32E-05	2	903	<i>ZNF318</i>
cg05244766	11	67107075	8.03E-05	3	787	<i>GSTP1</i>
cg22445920	5	150663371	1.38E-04	4	149	<i>SLC36A3</i>
cg12026956	5	72147764	2.03E-04	5	490	<i>TNPO1</i>
cg08385610	8	20156963	2.23E-04	6	120	<i>LZTS1</i>
cg10453758	3	133862880	2.50E-04	7	1264	<i>ACAD11</i>
cg19764399	13	47567117	2.81E-04	8	124	<i>MED4</i>
cg00922727	6	109521816	3.18E-04	9	154	<i>SESN1</i>
cg23364287	3	48729780	3.99E-04	10	70	<i>IHPK2</i>
cg04609640	1	45026679	4.21E-04	11	715	<i>VMD2L2</i>
cg23743114	17	31352509	4.83E-04	12	616	<i>CCL15</i>
cg27626899	15	89276929	5.27E-04	13	149	<i>HDDC3</i>
cg23323879	15	63291587	5.85E-04	14	723	<i>CILP</i>
cg16746462	5	96296194	6.36E-04	15	908	<i>LNPEP</i>
cg07246225	2	144994129	6.50E-04	16	257	<i>ZFHx1B</i>
cg05449607	6	111386305	6.78E-04	17	151	<i>C6orf51</i>
cg10167296	1	204747251	7.16E-04	18	251	<i>RASSF5</i>
cg10051054	1	3659032	7.66E-04	19	210	<i>CCDC27</i>
cg23487586	16	83873424	8.29E-04	20	641	<i>MGC22001</i>
cg09450020	7	89679371	8.31E-04	21	435	<i>STEAP2</i>
cg21850254	19	42651693	8.43E-04	22	129	<i>ZNF570</i>
cg08866753	1	39814648	8.66E-04	23	355	<i>PABPC4</i>
cg00600684	10	122600148	8.71E-04	24	537	<i>BRWD2</i>
cg05028306	4	17187761	8.95E-04	25	264	<i>LAP3</i>
cg09180926	15	89338658	9.11E-04	26	150	<i>PRC1</i>
cg04868764	15	38924311	9.17E-04	27	773	<i>SPINT1</i>
cg13726191	4	15549131	9.20E-04	28	62	<i>FGFBP1</i>
cg08303146	11	2423231	9.71E-04	29	NA	<i>KCNQ1</i>
cg18493238	10	37454267	1.00E-03	30	524	<i>ANKRD30A</i>
cg02449978	3	48205171	1.01E-03	31	366	<i>CDC25A</i>
cg03077492	5	43448852	1.02E-03	32	623	<i>CCL28</i>
cg24134767	11	113350848	1.03E-03	33	272	<i>HTR3A</i>
cg23863670	5	43447686	1.04E-03	34	543	<i>CCL28</i>
cg12820481	1	209915383	1.05E-03	35	207	<i>NEK2</i>
cg07123548	19	45588003	1.09E-03	36	69	<i>HIPK4</i>
cg16791508	12	51001958	1.13E-03	37	520	<i>KRTHB3</i>
cg09126273	12	15367034	1.17E-03	38	NA	<i>PTPRO</i>
cg07710481	13	87123385	1.21E-03	39	514	<i>SLITRK5</i>
cg16961218	18	74930131	1.24E-03	40	254	<i>ATP9B</i>