

YOUR EPIGENETIC METHYLATION DATA

A Brief Summary



Can any other DNA company tell you the specific spots they look at to determine your health outcomes? Well, we can!

In fact, we believe that this is our responsibility considering that the data we create for you is owned entirely by you.

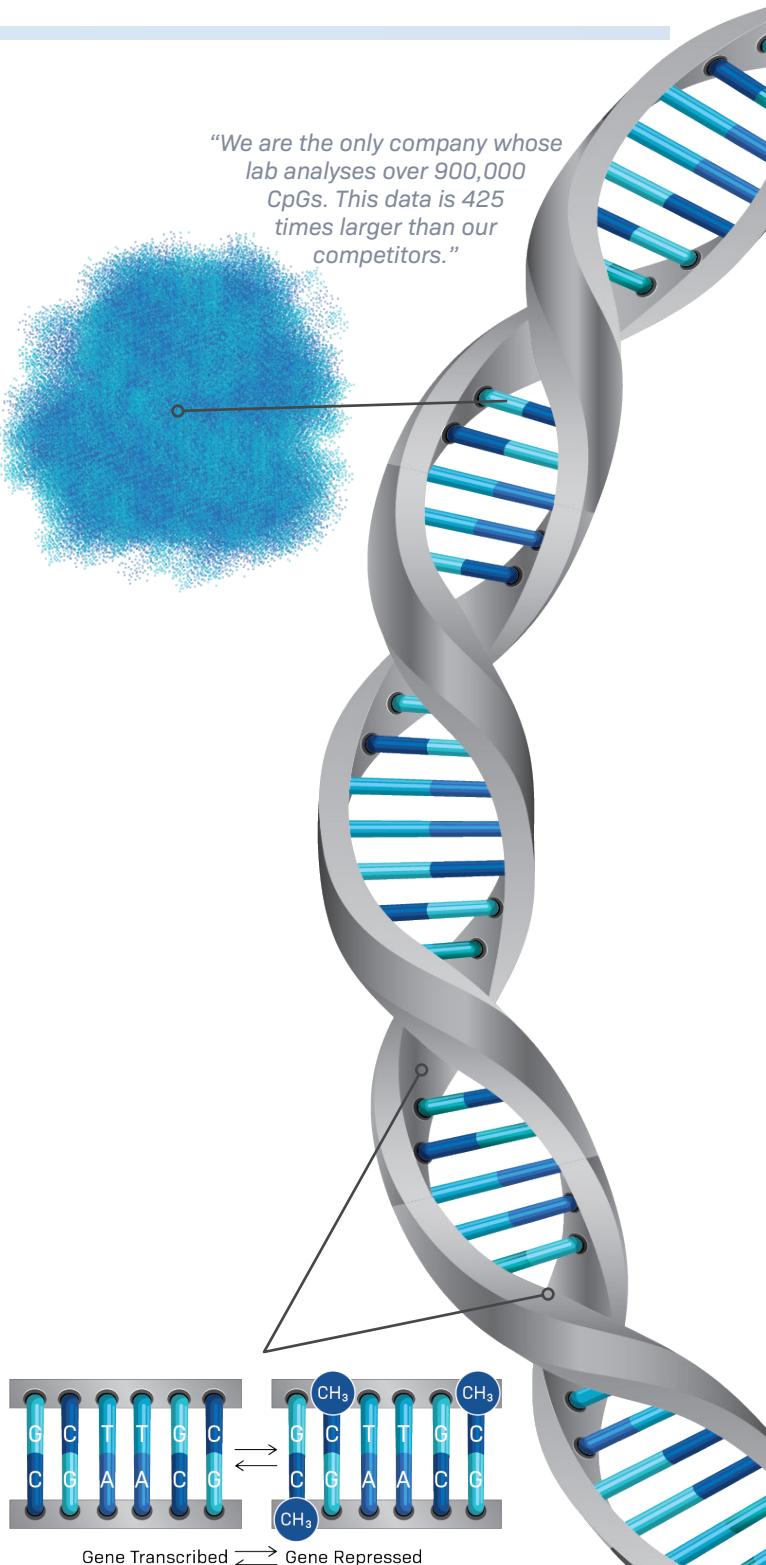
We are committed to total transparency in how we handle and report your data. We also want to deliver your data to you in digestible formats.

Our analysis looks at specific spots in the genome called CpGs. By looking at these locations, and their respective methylation values, we are able to determine the health metrics we report. We use the methylation values of your CpGs to determine your biological age, your pace of aging, and other determinants related to health.

You probably are already familiar with the double-stranded structure of DNA, where each side of the DNA is a strand that is made up of one of four nucleotide bases: cytosine, guanine, adenine, and thymine. Each base has a complementary base that is paired across the double helix, known as base-pairs. These nucleotide base pairings are cytosine with guanine and adenine with thymine.

A CpG is a cytosine that precedes a guanine nucleotide base in a linear order. A CpG site is practically the only site in the genome that will contain the attachment of a methyl group, which is a stable CH_3 molecule. The attachment of a methyl group to the DNA is known as DNA methylation. In mammals, 70 to 80 percent of CpG cytosines are methylated making these sites valuable tools for looking at and analyzing the different characteristics of your health.

"We are the only company whose lab analyses over 900,000 CpGs. This data is 425 times larger than our competitors."



CpG islands are short stretches of palindromic DNA with the sequence "CpG" that code for the same sequence on the complementary strand of DNA (repeated cytosine and guanine nucleotides with the "p" representing the linking phosphate).

The most common way to measure DNA methylation is calculating beta values, which are generated using Illumina's beadchip array and quantifying them from a series of mathematical equations. The beta value is the ratio of the methylated probe intensity. The beta values are a score between 0 and 1 and can be interpreted as the approximation of the percentage of methylation for the population of a given CpG site in the sample.

$$\text{Beta}_i = \frac{\max(y_{i,\text{methy}}, 0)}{\max(y_{i,\text{unmethy}}, 0) + \max(y_{i,\text{methy}}, 0) + \alpha} \quad (1)$$

The intensities are defined as the following mathematical notation shown above.

While we believe that the interpretation of these health metrics is important (such as biological age calculations), we also believe that you should know your methylation status and as a result, we want to provide you the raw data we use in this analysis.

How It Compares Against Other Methylation Clocks

Your data is your property. This means you should have complete access to it. Having access to your beta values for these specific CpG sites gives you the freedom to use your epigenetic information however you would like. You can even use your beta values in published algorithms! This gives you control over how you want to interpret your methylation values beyond what TruDiagnostic™ reports out.

As a reminder, we aren't determining biological age and other outputs based on your survey responses; the survey is meant to gather as much information about you in order to compile the most robust data set based on each person who takes our test. The survey gives us the most accurate depiction of your health history, including phenotypic and genotypic information.

Did you know that most biologic companies don't run your DNA themselves? **At TruDiagnostic™, all testing is done in our private laboratory where data stays encrypted and will never be shared with anyone, except YOU.** Unlike other DNA companies, we will never sell your data to third-party establishments.

The only purpose your health data serves is to grow the worth of our test by sharpening the values assigned to specific sites in your genome. For example, if we have a significant number of people who have gallstones and also have their DNA methylated at the exact same site, we can refine our outputs to include this correlation.

Analyzing CpGs

The technology we use is expensive and unique. Because your DNA was processed in our lab, we want you to take advantage of the outcomes of your DNA methylation levels at these crucial spots. The high-tech array technology we use looks at over 900,000 spots in your DNA.

We work with this vast data set because the more methylation values we utilize, the more robust TruAge™'s findings become. After your DNA's methylation values have been determined with our array scanning software your data is held as an .idat file for long-term storage in our highly secure server.

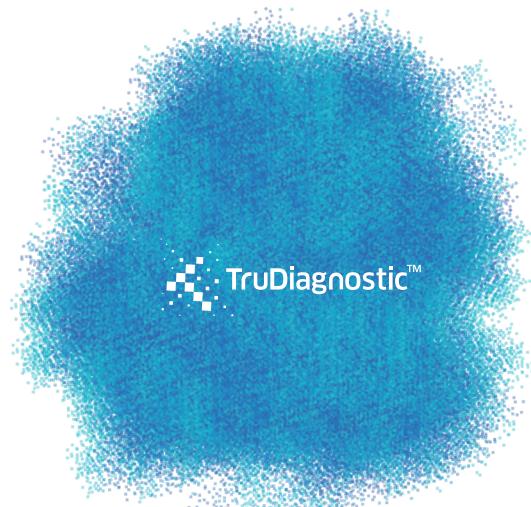
The beauty of storing your data in our repository is that we are able to run future reports based on your methylation values from the sample you have already provided. These future reports are created as soon as relevant epigenetic research has been published, which adds prodigious value to our test at no additional cost to you.

*We are the only company who's lab looks at over **900,000 CpGs**, this data is 4,000 times larger than our competitors.*



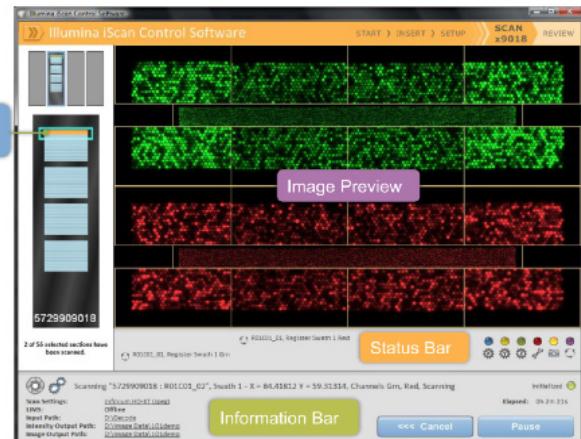
Competitors

vs

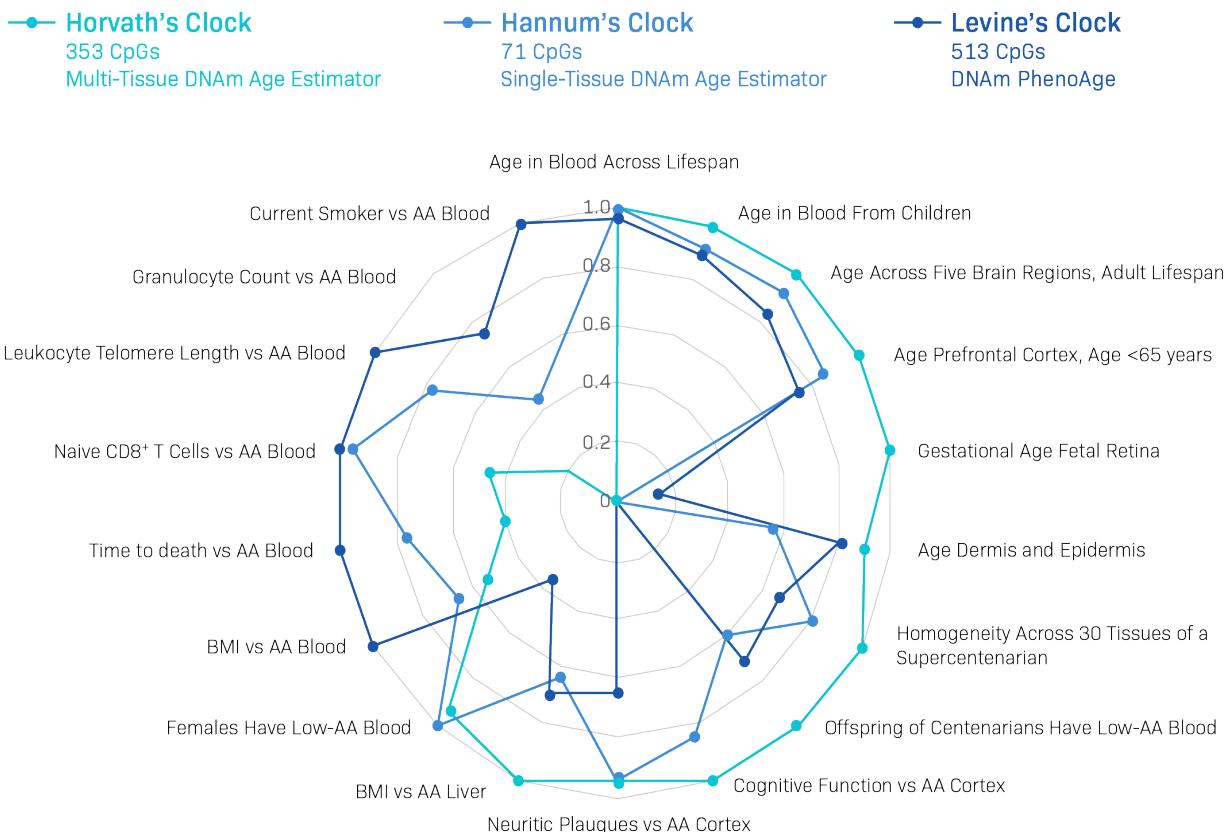


TruDiagnostic™

Monitoring The Scan Progress



Our array scanning software assigns either red or green probes to each CpG to determine the level of methylation at each spot. The iScan control software then converts each spot's methylation data in order to input the data into advanced mathematical algorithms.



A comparison of three biological age estimates based on DNA methylation-based biomarkers. These three clocks' age estimates are resulting from mathematical algorithms for ageing.

Your Specific Beta Values

Using our test will open up many doors for you because now you have autonomous control over whichever algorithm you wish to input your values into. By sharing access to your personal data, you have the power to go above and beyond TruDiagnostic™'s findings by utilizing these results however you please.

On the next page is the complete list of all the CpGs that TruDiagnostic™ and many other popular algorithms use! We are providing your specific beta values at the following CpG locations. The reason we chose this list of 1,300 CpGs versus the total 900,000 is that we believe these are the most useful sites for you.

If you wish to see the complete list of CpGs and your beta values at each spot, we can make that happen per your request. This list is constantly growing and being revised because TruDiagnostic™ does not accept the stagnation of our company's value and neither should you!

Your CpG Values

TruDiagnostic™

CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg00079056	ABHD17B; C9orf85	0.19	cg02480835	NUPL2	0.9
cg00083937	MAPK8IP2	0.34	cg02503970	ST3GAL1	0.81
cg00113951	CABP1	0.74	cg02631957	CSNK1G3	0.06
cg00168942	GJD4	0.86	cg02735486	ANK2	0.15
cg00194146	GFOD1	0.11	cg02802055	USP20	0.22
cg00230271	MGC16169	0.33	cg02976574	FNDC3B	0.19
cg00261781	MATN3	0.37	cg03007010	SNX26	0.34
cg00297600	UNC5D	0.9	cg03112869	FBXW12	0.74
cg00335286	MC2R	0.81	cg03172991	NFIX	0.86
cg00338702	CHFR	0.06	cg03258472	CRB3	0.11
cg00350702	ERGIC3	0.15	cg03340261	HBS1L	0.33
cg00410898	STC1	0.22	cg03387497	BANF2	0.37
cg00412772	C19orf33; YIF1B; oC19orf33	0.19	cg03535648	PMCH	0.9
cg00412805	KBTBD5	0.34	cg03565081	C14orf93	0.81
cg00462994	NAGS	0.74	cg03623878	MCF2L	0.06
cg00503840	DLX5	0.86	cg03703325	HEMK1	0.15
cg00515905	EPS8L3	0.11	cg03724882	CDKN3	0.22
cg00582628	RGS20	0.33	cg03819692	RNA 304"-LINC00304	0.19
cg00687674	EWSAT1	0.37	cg03929796	ALAS1	0.34
cg00744433	CXADR	0.9	cg03977782	SIDT1	0.74
cg00845900	CPA4	0.81	cg03991512	LDHD	0.86
cg00862290	KCNMB3	0.06	cg04007936	CARHSP1	0.11
cg00943950	UBE2B	0.15	cg04014889	MAGEL2	0.33
cg00955230	RFC2	0.22	cg04084157	VGF	0.37
cg01056568	GLP2R	0.19	cg04087608	HSD3B2	0.9
cg01114088	CYTL1	0.34	cg04169469	ODC1	0.81
cg01128603	SF3B2; SNORD13F	0.74	cg04333463	ASPH	0.06
cg01131735	SC65	0.86	cg04359302	MRPL34	0.15
cg01137065	FOXK2	0.11	cg04416752	ST6GALNA4	0.22
cg01311097		0.33	cg04424621	HIST1H2BJ	0.19
cg01221637	C20orf112	0.37	cg04480914	PPP2R5B	0.34
cg01252496	MDS032	0.9	cg04528819	KLF14	0.74
cg01254459	PSMC6	0.81	cg04601137	ADAMTSL5	0.86
cg01261503	POLG2	0.06	cg04616566	THSD4	0.11
cg01335367	C12orf34	0.15	cg04718414	CDC16	0.33
cg01400401	C19orf30	0.22	cg04736140	GPR161	0.37
cg01441777	CSNK1E	0.19	cg04755031	PNPLA5	0.9
cg01450842	STON1-GTF2A1L	0.34	cg04818845	PLEKHG5	0.81
cg01459463		0.74	cg04836038	DOCK9	0.06
cg01511567	SSRP1	0.86	cg05087948	CTSF	0.15
cg01519742	JAKMIP1	0.11	cg05125838	UCN2	0.22
cg01623187	FAM112A	0.33	cg05228408	CLCN6	0.19
cg01626227	TRIM4	0.37	cg05270634	RND2	0.34
cg01651821	MRPL30; MITD1	0.9	cg05294243	KLK13	0.74
cg01918706	UBE2T	0.81	cg05316065	MLZE	0.86
cg01930621	ZNF649	0.06	cg05422352	COPS7B	0.11
cg01946401	RUNX2	0.15	cg05440289	IVL	0.33
cg02016419	TEKT3	0.22	cg05441133	GDF2	0.37
cg02073105		0.19	cg05442902	P2RXL1	0.9
cg02151301	HM13	0.34	cg05473871	ADAMTSL5	0.81
cg02154074	HTRA2	0.74	cg05492270	RNF121	0.06
cg02197293	FBXO27	0.86	cg05501584	ST8SIA2	0.15
cg02228185	ASPA	0.11	cg05532892	PSMA1	0.22
cg02229946	IL1F7	0.33	cg05697249	C11orf52	0.19
cg02309431	BPII3	0.37	cg05759269	CDKN1A	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg05851163	C18orf4	0.19	cg09304040	CDK2	0.9
cg05898102	PCDHB10	0.34	cg09322949	TRFP	0.81
cg06134964	ITIH1	0.74	cg09404633	LMOD1	0.06
cg06144905	PIPOX	0.86	cg09413557	S100A6	0.15
cg06171242	TTRAP	0.11	cg09434995	SRCRB4D	0.22
cg06189653	PRR7	0.33	cg09480837	PLCE1	0.19
cg06295856	CALCA	0.37	cg09548179	AMDHD2	0.34
cg06327515	PCDHB14	0.9	cg09556292	RCOR1	0.74
cg06363129	SOSTDC1	0.81	cg09630437	SLC39A3	0.86
cg06493994	SCGN	0.06	cg09799873	NDUFA7	0.11
cg06533629	KLF13	0.15	cg09809672	EDARADD	0.33
cg06637774	P2RY6	0.22	cg09851465	C1orf87	0.37
cg06638451	FAM107A	0.19	cg09892203	CACNG4	0.9
cg06690548	SLC7A11	0.34	cg10052840	SEMA6B	0.81
cg06908778	SPAG6	0.74	cg10158181	GRK4	0.06
cg06958034	AOF2	0.86	cg10202457	NMNAT2	0.15
cg06975499	EZH1	0.11	cg10225525	FUT8	0.22
cg06994793	PRO1580	0.33	cg10523019	RHBDD1	0.19
cg07038400	PPP2R3A	0.37	cg10570177	C9orf7	0.34
cg07073964	PRSSL1	0.9	cg10591174	MYL2	0.74
cg07180649	WNT4	0.81	cg10636246	AIM2	0.86
cg07211259	PDCD1LG2	0.06	cg10654016	C12orf31	0.11
cg07236943	PEG10	0.15	cg10667970	FBXO6	0.33
cg07265300	PFN4	0.22	cg10669058	CILP2	0.37
cg07484827	CHRNA10	0.19	cg10795646	S100A10	0.9
cg07494518	SNF1LK	0.34	cg10878896	CS	0.81
cg07654934	LXN	0.74	cg10900550	TDRD7	0.06
cg07817698	DHODH	0.86	cg10917602	HSD3B7	0.15
cg07850604	INSM2	0.11	cg10922280	DPEP2	0.22
cg07929310	KIAA0513	0.33	cg11177450	SENP5	0.19
cg08035942	ZADH2	0.37	cg11233384	AKR7A3	0.34
cg08067365	G1P2	0.9	cg11237115	KLHL24	0.74
cg08074477	TNS4	0.81	cg11426590	TRIM62	0.86
cg08169325	EPIM	0.06	cg11459714	KLK10	0.11
cg08212685	ATG10	0.15	cg11487705	OXCT1	0.33
cg08251399	EHD3	0.22	cg11490446	PSEN1	0.37
cg08331960	SLC9A3R2	0.19	cg11600161	TBC1D10C	0.9
cg08424423	CDSN	0.34	cg11618577	KRTCAP3	0.81
cg08475827	RIF1	0.74	cg11631518	KCNC2	0.06
cg08487374	TSC22D4	0.86	cg11833861	TMEM98	0.15
cg08529529	ALOX5AP	0.11	cg11896923	PNMT	0.22
cg08586737	GCC1	0.33	cg11903057	RHOH	0.19
cg08587542	KIAA0141	0.37	cg12145907	PCDHGC4	0.34
cg08654655	IGSF4B	0.9	cg12177001	IFI27	0.74
cg086668790	ZNF154	0.81	cg12188560	CNTFR	0.86
cg08694544	RTBDN	0.06	cg12238343	RLN3R1	0.11
cg08872493	HTR1D	0.15	cg12247247	RPL23	0.33
cg08896629	TYK2	0.22	cg12261786	C10orf116	0.37
cg08899632	ZFP64	0.19	cg12265604	HAS3	0.9
cg08900043	HOXC10	0.34	cg12269343	SEL1L	0.81
cg09045681	TCF19	0.74	cg12289045	SYT3	0.06
cg09096950	UBE4A	0.86	cg12324144	LRRC34	0.15
cg09196959	TRIM40	0.11	cg12373771	CECR6	0.22
cg09254939	KLK10	0.33	cg12402251	CALB1	0.19
cg09294589	CRTAP	0.37	cg12473775	RHOD	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg12743894	C11orf46	0.19	cg15201877	PTGER3	0.9
cg12813792	C20orf85	0.34	cg15344028	ICOS	0.81
cg12864235	CDH9	0.74	cg15381313	NR1H4	0.06
cg12985418	MIB1	0.86	cg15427448	BACE1	0.15
cg12991365	CHDH	0.11	cg15447479	SMO	0.22
cg13042288	ANPEP	0.33	cg15489301	AKR1B10	0.19
cg13119609	APOC2	0.37	cg15498283	TALD01	0.34
cg13120519	ALKBH7	0.9	cg15551881	TRAF1	0.74
cg13218906	PCMTD2	0.81	cg15569512	C14orf153	0.86
cg13258700	C7orf13	0.06	cg15611364	OXSM	0.11
cg13307384	PRPF40B	0.15	cg15642326	MARK4	0.33
cg13323474	TTC18	0.22	cg15811427	FPR1	0.37
cg13351161	SCARA3	0.19	cg15856055	ZNF511	0.9
cg13409216	SUPV3L1	0.34	cg15881088	ARMET	0.81
cg13449372	MRPL43	0.74	cg15887846	CST6	0.06
cg13460409	DSCR6	0.86	cg15903282	EPHB1	0.15
cg13509147	CREB3E3	0.11	cg15963417	PAH	0.22
cg13510262	SP1	0.33	cg15966757	SLC6A13	0.19
cg13514050	GGTL3	0.37	cg16085042	HSP90B1	0.34
cg13550877	C14orf148	0.9	cg16173067	SDCBP2	0.74
cg13564075	PHF12	0.81	cg16295988	C9orf80	0.86
cg13571802	TGM4	0.06	cg16313343	BRF1	0.11
cg13587552	SCNN1D	0.15	cg16319578	HSPA2	0.33
cg13613532	MGC33948	0.22	cg16340918	KIF11	0.37
cg13631913	MGC4399	0.19	cg16354207	UBE2F	0.9
cg13654195	RIOK1	0.34	cg16357381	COL7A1	0.81
cg13656062	CYP4F2	0.74	cg16372520	NRXN3	0.06
cg13656360	MLL	0.86	cg16408970	BAMBI	0.15
cg13700897	RSP02	0.11	cg16466334	MMP3	0.22
cg13718960	RNASE1	0.33	cg16543027	PLCB2	0.19
cg13843773	MGC35043	0.37	cg16612562	RRP22	0.34
cg13854874	CHAF1B	0.9	cg16648841	C8A	0.74
cg13861644	PIWIL1	0.81	cg16713727	SLC25A34	0.86
cg13899108	PDE4C	0.06	cg16718891	ITGA5	0.11
cg13975369	TSGA14	0.15	cg16728114	C1orf66	0.33
cg13994175	KIF9	0.22	cg16743289	CACYBP	0.37
cg14009688	CALD1	0.19	cg16816226	EDEM1	0.9
cg14105047	PDLIM5	0.34	cg16854606	DAND5	0.81
cg14159818	C20orf27	0.74	cg16933388	BSN	0.06
cg14175438	FAM3C	0.86	cg16984944	TBC1D23	0.15
cg14223995	UCP1	0.11	cg17009433	GLDC	0.22
cg14281160	C12orf31	0.33	cg17038116	ASH1L	0.19
cg14350002	GAS1	0.37	cg17129388	NGFR	0.34
cg14423778	MBNL1; LOC40109 3	0.9	cg17133388	C3orf28	0.74
cg14467840	S100A1	0.81	cg17324128	RASSF4	0.86
cg14473016	RASD2	0.06	cg17431739	MSRB2	0.11
cg14550518	ZNF659	0.15	cg17526300	PTER	0.33
cg14689355	ACVR2A	0.22	cg17536848	LMBR1	0.37
cg14747225	POLR3K	0.19	cg17605084	HEM1	0.9
cg14754581	CCRL2	0.34	cg17627559	TSPAN32	0.81
cg14916213	RBM4	0.74	cg17641104	PCTK2	0.06
cg14918082	KCNAB3	0.86	cg17726022	SLC38A1	0.15
cg14972143	EIF4E	0.11	cg17749443	ZFP37	0.22
cg15013019	LYL1	0.33	cg17770886	ZNF511	0.19
cg15171237	CASP3	0.37	cg17861230	PDE4C	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg17896249	E2F4	0.19	cg20570279	ZNF259	0.9
cg17903544	PGAM5	0.34	cg20572838	EME1	0.81
cg17923358	RELN	0.74	cg20652640	ZFPM1	0.06
cg17940013	GPR157	0.86	cg20674577	PHACTR3	0.15
cg17966192	SULT1C2	0.11	cg20761322	CIB2	0.22
cg18001427	C21orf6	0.33	cg20828084	KIAA1199	0.19
cg18003795	CECR1	0.37	cg20891917	IFRD1	0.34
cg18117393	NPL	0.9	cg20967028	ART4	0.74
cg18241647	WFDC12	0.81	cg21006686	NOS1	0.86
cg18267374	NEF3	0.06	cg21053529	CX36	0.11
cg18384097	PTPN7	0.15	cg21081971	TRPC4	0.33
cg18392482	AMDHD1	0.22	cg21099326	TRIM35	0.37
cg18468844	PTAFR	0.19	cg21120249	FLJ36268	0.9
cg18587364	ZNF562	0.34	cg21137706	TMEM5	0.81
cg18691434	GPC2	0.74	cg21184495	ZNF207	0.06
cg18693704	CMTM4	0.86	cg21200703	SLC34A2	0.15
cg18732541	MCEMP1	0.11	cg21201109	ANKRD25	0.22
cg18771300	RHOJ	0.33	cg21207418	ACP5	0.19
cg18809289	ALOX5	0.37	cg21296230	GREM1	0.34
cg18881501	MAP2K3	0.9	cg21363706	NAT1	0.74
cg18996776	KPNA4	0.81	cg21649520	PMP2	0.86
cg19008809	SFMBT1	0.06	cg21712685	UNQ3045	0.11
cg19028160	HDAC4	0.15	cg21762589	BNIP3	0.33
cg19104072	B3GNT6	0.22	cg21801378	BRUNOL6	0.37
cg19149785	KLK8	0.19	cg21835643	RBPSUHL	0.9
cg19287114	SLC44A1	0.34	cg21907579	TBX5	0.81
cg19297232	SMPD3	0.74	cg21926612	PARK2	0.06
cg19345165	NDUFS3	0.86	cg21993406	CENPH	0.15
cg19356189	KLK10	0.11	cg22090592	C14orf173	0.22
cg19371795	INSR	0.33	cg22179082	GPR12	0.19
cg19378133	A2BP1	0.37	cg22194129	CLEC4C	0.34
cg19398783	TLR10	0.9	cg22197830	TXNDC15	0.74
cg19439331	CXXC6	0.81	cg22282672	FLJ32921	0.86
cg19514469	ELMO3	0.06	cg22395019	GALNT14	0.11
cg19556572	AKNA	0.15	cg22396353	GTSE1	0.33
cg19560210	PGBD5	0.22	cg22407458	TCP11	0.37
cg19566405	FLJ10260	0.19	cg22473095	RBBP9	0.9
cg19573166	SLC22A17	0.34	cg22484793	TLR9	0.81
cg19586576	GJA7	0.74	cg22495124	CACNG5	0.06
cg19615059	TRPV4	0.86	cg22511262	WT1	0.15
cg19632206	FAM5C	0.11	cg22512531	CRTAM	0.22
cg19663795		0.33	cg22580353	PVR	0.19
cg19685066	ACTL8	0.37	cg22582569	PPP2R1B	0.34
cg19686152	TMOD3	0.9	cg22594309	SYT2	0.74
cg19722847	IPO8	0.81	cg22736354	NHLRC1	0.86
cg19724470	CD274	0.06	cg22809047	RPL31	0.11
cg19731122	MYO1C	0.15	cg22947000	BCMO1	0.33
cg19883905	CIDEA	0.22	cg22971191	SLC10A2	0.37
cg20066677	CHFR	0.19	cg22983092	KRT25A	0.9
cg20090497	TAS2R9	0.34	cg22991148	C21orf29	0.81
cg20162159	MON1A	0.74	cg23124451	CBX7	0.06
cg20173259	LHB	0.86	cg23127998	FLRT3	0.15
cg20234170	CAT	0.11	cg23152772	FIBCD1	0.22
cg20492933	MRPL22	0.33	cg23159337	ATP13A4	0.19
cg20550118	CRABP1	0.37	cg23173910	ACTG2	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg23191950	ALDH1A3	0.19	cg26746469	KIAA0406	0.9
cg23213217	DEGS1	0.34	cg26815229	CYP2J2	0.81
cg23234999	MKRN3	0.74	cg26824091	GL01	0.06
cg23239039	D4S234E	0.86	cg26842024	KLF2	0.15
cg23338195	SLC30A8	0.11	cg26866325	PPBPL2	0.22
cg23376526	SLC27A3	0.33	cg26898166	EMX1	0.19
cg23568913	PES1	0.37	cg26932976	KIAA0664	0.34
cg23668631	CAMKK1	0.9	cg27015931	C16orf65	0.74
cg23710218	MSC	0.81	cg27187881	NAGA	0.86
cg23818978	AMICA1	0.06	cg27244482	CAMK2A	0.11
cg23832061	PPIL6	0.15	cg27367952	MGC50811	0.33
cg24110063	COX6A2	0.22	cg27440834	SNX4	0.37
cg24125648	HYPK	0.19	cg27493997	CRABP2	0.9
cg24208206	FYCO1	0.34	cg27514224	NUP188	0.81
cg24304712	C1orf101	0.74	cg27626102	STEAP2	0.06
cg24332433	SLC39A11	0.86	cg27655905	C11orf24	0.15
cg24407308	DGKZ	0.11	cg00075967	STRA6	0.22
cg24493940	MMP17	0.33	cg00374717	ARSG	0.19
cg24505122	WNT5B	0.37	cg00864867	PAWR	0.34
cg24505341	CADPS	0.9	cg00945507	SEC61G	0.74
cg24556026	GNE	0.81	cg01027739	DOLPP1	0.86
cg24651706	CHCHD5	0.06	cg01353448	C7orf16	0.11
cg24674703	CD5	0.15	cg01584473	MUC17	0.33
cg24921089	AMPD3	0.22	cg01644850	ZNF551	0.37
cg25022327	P53AIP1	0.19	cg01656216	ZNF438	0.9
cg25092328	PXMP4	0.34	cg01873645	FAM108B 1; C9orf85	0.81
cg25136687	REV3L	0.74	cg01968178	REEP1	0.06
cg25229964	CNKS1R	0.86	cg02085507	TRIP10	0.15
cg25251635	POP7	0.11	cg02217159	KHDRBS2	0.22
cg25256723	F5	0.33	cg02331561	ABCA17P; ABCA3	0.19
cg25428451	LOC339524	0.37	cg02332492	C8G	0.34
cg25459323		0.9	cg02364642	GEFT	0.74
cg25536676	DHCR24	0.81	cg02388150	SFRP1	0.86
cg25713185	SLC19A3	0.06	cg02479575	MIR7-3; C19orf30	0.11
cg25769980	TLR6	0.15	cg02489552	CCDC105	0.33
cg25881193	BTN2A2	0.22	cg02580606	KRT33B	0.37
cg25898500	HAGH	0.19	cg02654291	C9orf64	0.9
cg26022315	SNF8	0.34	cg02827112	SMARCA D1	0.81
cg26091688	ZNF200	0.74	cg02972551	KDM3A	0.06
cg26096837	FGF19	0.86	cg03103192	SPATA18	0.15
cg26104204	MAP3K10	0.11	cg03167275	CXADR	0.22
cg26109803	WNT8B	0.33	cg03270204	DDR1	0.19
cg26201213	MGMT	0.37	cg03565323	ZNF287	0.34
cg26212924	CYC1	0.9	cg03588357	GPR68	0.74
cg26219051	SCUBE1	0.81	cg03760483	ALOX12	0.86
cg26312920	NEUROD6	0.06	cg04126866	C10orf99	0.11
cg26350286	RBM35A	0.15	cg05250458	ZNF177	0.33
cg26357744	RASSF1	0.22	cg05365729	LOXL2	0.37
cg26382071	TXNL5	0.19	cg05675373	KCNC4	0.9
cg26394737	ABHD4	0.34	cg05755779	COLEC10	0.81
cg26394940	C22orf26; LOC15038 1	0.74	cg05921699	CD79A	0.06
cg26581729	NPDC1	0.86	cg05960024	CLOCK	0.15
cg26614073	SCAP	0.11	cg06121469	SPG11	0.22
cg26665419	C6orf203	0.33	cg06361108	CCNF	0.19
cg26711820	MYF6	0.37	cg06462291	NTSDC3	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg06557358	TMEM132 E; C17orf102	0.19	cg15974053	HSD17B14	0.9
cg06738602	PTGER2	0.34	cg15988232	CSPG5	0.81
cg06810647	CRAMP1L	0.74	cg16150435	C6orf15	0.06
cg06952310	NCAN	0.86	cg16241714	CEBPD	0.15
cg06993413	DPP8	0.11	cg16494477	FGF18	0.22
cg07285276	RAPGEF1	0.33	cg16547529	KLHL35	0.19
cg07291563	GRWD1	0.37	cg16579101	NOP2	0.34
cg07337598	ANXA9	0.9	cg17063929	NOX4	0.74
cg07455279	NDUFA3	0.81	cg17099569		0.86
cg07595943	ADAD2	0.06	cg17285325	TYMP	0.11
cg08030082	POMC	0.15	cg17408647	C7orf44	0.33
cg08090772	ADHFE1	0.22	cg17655614	CDH1	0.37
cg08124722	CCL7	0.19	cg17729667	NINL	0.9
cg08251036		0.34	cg17853587	NDST3	0.81
cg08370996	NR2F2	0.74	cg17960516	DOK7	0.06
cg08413469	DEPDC1	0.86	cg18055007	DDAH2	0.15
cg08434234	DGKI	0.11	cg18180783	MYOZ1	0.22
cg08771731	LOC28569 6; BASP1	0.33	cg18440048	ZNF70	0.19
cg08965235	LTBP3	0.37	cg18573383	KCNC2	0.34
cg09019938	PRKG1	0.9	cg18983672	FOXE3	0.74
cg09118625	DIRAS3	0.81	cg18984151	C3orf75	0.86
cg09191327	PRDM12	0.06	cg19167673	PDGFB	0.11
cg09418283	PAWR	0.15	cg19273182	PAPOLG	0.33
cg09509673	CCR10; CNTNAP1	0.22	cg19305227	SLC28A2	0.37
cg09785172	WFS1	0.19	cg19346193	BCCIP; UROS	0.9
cg09869858	P11	0.34	cg19478743	ZMYND15; CXCL16	0.81
cg09885951	CENPF	0.74	cg19514928	TMEM56	0.06
cg10281002	TBX5	0.86	cg19692710	DNAJB13	0.15
cg10376763	TNP1	0.11	cg19945840	SDF4; B3GALT6	0.22
cg10377274	PATE1	0.33	cg20295671	YPEL1	0.19
cg10486998	GALR1	0.37	cg20305610	PDLIM5	0.34
cg10920957	JPH3	0.9	cg20524216	C3orf75	0.74
cg11932564	TNFRSF13 C	0.81	cg20692569	FZD9	0.86
cg12351433	LHCGR	0.06	cg20795863	NEU2	0.11
cg12768605	LYPD5	0.15	cg20914508	GAP43	0.33
cg12830694	PPP1R14A	0.22	cg20947775	SCD5	0.37
cg12946225	HMG20B	0.19	cg20999813	USP10	0.9
cg13038560	C2orf60; C2orf47	0.34	cg21096399	MCAM	0.81
cg13216057	DKK3	0.74	cg21378206	IL1F5	0.06
cg13319175	CAPZB	0.86	cg21460081	HOXB4	0.15
cg13682722	C14orf102	0.11	cg21870884	GPR25	0.22
cg13836627	TJP1	0.33	cg22006386	CATSPER G	0.19
cg14258236	OR5V1	0.37	cg22289837	CA3	0.34
cg14308452	PRR22	0.9	cg22432269	CYFIP1	0.74
cg14329157	WDR69	0.81	cg22449114	TCF15	0.86
cg14424579	AGBL5	0.06	cg22679120	SNX8	0.11
cg14501253	C8orf79	0.15	cg22901840	DIRAS3	0.33
cg14658362	RBPM5	0.22	cg22920873	C7orf55	0.37
cg14723032	PITPNM3	0.19	cg23517605	TUBB2B	0.9
cg14894144	LAMA3	0.34	cg23662675	ZMYND8	0.81
cg14992253	EIF3I; C1orf91	0.74	cg23941599	FEM1C	0.06
cg15341340	DNASE2	0.86	cg24116886	DEFB127	0.15
cg15381769	PTPRK	0.11	cg24126851	DCHS1	0.22
cg15547534	C7orf47	0.33	cg24254120	RFC3	0.19
cg15661409	C14orf105	0.37	cg24262469	TIPARP; LOC10028 7227	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg24450312	RASSF5	0.19	cg05590257	PLD6	0.9
cg24580001	CCDC88B	0.34	cg05847778	BBS5	0.81
cg24834740	PPP1R16B	0.74	cg05903609	PRPF8	0.06
cg25070637	SDC2	0.86	cg06044899	TMSL3; FAM190A	0.15
cg25148589	GRIA2	0.11	cg06117855	CLEC3B	0.22
cg25505610	EIF3M	0.33	cg06513075	NAT10	0.19
cg25552492	LGI3	0.37	cg06688848	RSPRY1; FAM192A	0.34
cg25683012	BAZ2A	0.9	cg06836772	PRKAA2	0.74
cg25771195	C16orf80	0.81	cg06926735	UBE2V1; TMEM198-UBE2V1	0.86
cg25781123	THUMPD3	0.06	cg07158339	FXN	0.11
cg26003813	PLK1	0.15	cg07388493	NDUFS5	0.33
cg26005082	MIR7-3; C19orf30	0.22	cg07403456	PGLYRP2	0.37
cg26045434	HR; HR	0.19	cg07498421	CRADD	0.9
cg26297688	C12orf23	0.34	cg07663789	NPR3	0.81
cg26372517	TFAP2E	0.74	cg07730301	ALDH3B1	0.06
cg26453588	BIK	0.86	cg07770222	C8orf31	0.15
cg26620959	SYNE1	0.11	cg07849904	MN1	0.22
cg26845300	SNX9	0.33	cg08186124	LZTFL1	0.19
cg27092035	ARL10	0.37	cg09133026	RPS6K1	0.34
cg27169020	BNC1	0.9	cg09441152	PQLC1	0.74
cg27319898	ZNF804B	0.81	cg09646392	TNFSF13B	0.86
cg27377450		0.06	cg09722397	GRIN2C	0.11
cg27413543	SEC31A	0.15	cg09722555	CCL27	0.33
cg27494383	LTK	0.22	cg10045881	CHI3L2	0.37
cg00091693	KRT20	0.19	cg10266490	ACOT11	0.9
cg00431549	MGP	0.34	cg10345936	SLC36A2	0.81
cg00436603	CYP2E1	0.74	cg10865119	C6orf122; C6orf208	0.06
cg01027805	ZNF219; C14orf176	0.86	cg10940099	CD164	0.15
cg01234063	ST3GAL4	0.11	cg11025793	IER2; STX10	0.22
cg01262913	DSCR9	0.33	cg11299964	MAPKAP1	0.19
cg01407797	CCDC117	0.37	cg11314684	AKT3	0.34
cg01459453	SELP	0.9	cg11388238	KCTD18	0.74
cg01485645	MLLT6	0.81	cg11653266	MRPL38	0.86
cg01560871	C10orf27	0.06	cg12413566	XIRP1	0.11
cg01570885	FAM50B	0.15	cg12616277	ESYT3	0.33
cg01820374	LAG3	0.22	cg12941369	PDCD6IP	0.37
cg02047577	UCKL1AS; UCKL1	0.19	cg13129046	C10orf35	0.9
cg02071305	VPS18	0.34	cg13269407	C22orf26; LOC15038 1	0.81
cg02275294	SOAT1	0.74	cg13302154	MGP	0.06
cg02335441	NEK11; ASTE1	0.86	cg13547237	C11orf68; DRAP1	0.15
cg03019000	TEX264	0.11	cg13828047	MPI	0.22
cg03286783	CASC4	0.33	cg13931228	MPP6	0.19
cg03330058	ABTB1	0.37	cg14060828	PTH2 (TIP39)	0.34
cg03578041	LARP6	0.9	cg14163776	ACAP2	0.74
cg03682823	SGCE; PEG10	0.81	cg14408969	C8orf40; SLC20A2	0.86
cg03891319	ACY1	0.06	cg14409958	ENPP2	0.11
cg03947362	C2orf60; C2orf47	0.15	cg14597908	GNASAS; GNAS	0.33
cg04005032	OSBPL10; ZNF860	0.22	cg14654875	NAT15; ZNF597	0.37
cg04094160	ZBTB5	0.19	cg14727952	BIRC2	0.9
cg04121983	CASKIN2	0.34	cg15185286	AIG1	0.81
cg04268405	CHST3	0.74	cg15262928	TIMM17A	0.06
cg04431054	PRRC1	0.86	cg15703512	C16orf65	0.15
cg04452713	DST	0.11	cg15804973	MAP3K5	0.22
cg04474832	ABHD14B	0.33	cg16034652	BTBD7; KIAA1409	0.19
cg04999691	C7orf29; LRRC61	0.37	cg16168311	APOA1BP	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg16358826	GABRA4		cg00073460	ZC3H12D	
cg16403394	RXRA		cg00147638	RUNX3	
cg16419345	ACOX1; C17orf106		cg00481951	SST	
cg16744741	PRKGG		cg00486113	PSORS1C1	
cg16899442	CCDC78; HAGHL		cg00495443	IL17RA	
cg17274064	ERG; ERG		cg00748589		
cg17338403	SLC03A1		cg00871371		
cg17589341	SLC14A1		cg01124420	EDAR	
cg17686885	TOM1L1		cg01345395		
cg18031008	MRPS21		cg01372366	PTPRJ	
cg18139769	SGCE; PEG10		cg01528542		
cg18328933	ABHD14B		cg02033323	FOXP3	
cg18956095	ZHX1		cg02046143	1GSF9B	
cg19044674	LEPRE1; C1orf50		cg02085953	ARID5A	
cg19046959	COL8A2		cg02650266		
cg19420968	HCRTR1		cg02867102		
cg19569684	MGC295_6		cg02988775	TNFRSF25	
cg19706682	LRRK50; HSDL1		cg02989940	AHSP	
cg19761273	CSNK1D		cg03032497		
cg19853760	LGALS1		cg03132824	RGS12	
cg20100381	NAE1		cg03370106	VGLL4	
cg20240860	ACCS		cg03399905	ANKRD34C	
cg21211748	E2F2		cg03467087	SYNJ2	
cg21305265	KCTD9; CDCA2		cg03473532	MKLN1	
cg21370143	MYBPC3		cg03607117	SFMBT1	
cg21395782	NDUFA13; TSSK6		cg04400972	TRIM45	
cg21950518	IL6ST		cg04416734	ALDOA	
cg22171829	PDK4		cg04683740	LBH	
cg22190114	NLRP8		cg04875128	OTUD7A	
cg22568540	NCRNA00_181; A1BG		cg04940570	TEAD1	
cg22613010	CLCN2		cg05217983	RUNX2	
cg22637507	ALKBH3		cg05392293	FRMD4A	
cg23092072	AFF1		cg05442902	MGC16703	
cg23180365	GLB1; TMPPE		cg05913271	CUL9	
cg23786576	ATPAF1		cg06245711	FAM100B	
cg24058132	GALC		cg06419846	CD248	
cg24081819	EPHX2		cg06445016		
cg24471894	KIAA0020		cg06639320	FHL2	
cg24888049	FES; FURIN		cg06685111	HCG18	
cg24899750	SNRPB2		cg06874016	NK1RAS2	
cg25101936	ZBTB16		cg06952412	LTBP4	
cg25159610	PLK2		cg07082267		
cg25166896	C22orf25		cg07094298	TN1PN2	
cg25411725	SLC22A13		cg07107916	PPP2R5A	
cg25564800	KPNA1		cg07547549	SLC12A5	
cg25657834	NTSR2		cg07553761	TRIM59	
cg25809905	ITGA2B		cg07583137	CHMP4C	
cg25928579	HOXB8		cg07899551		
cg26043391	FBXO28		cg07927379	C70rf13	
cg26162695	ELAC2		cg07955474	1RF8	
cg26456957	PPP1R12C		cg07955995	KLF14	
cg26723847	VPS26B; NCAPD3		cg08097417	KLF14	
cg27016307	HRC		cg08234504		
cg27202708	C1orf65		cg08415592	APOL1	
cg27544190	C21orf63		cg08454507	RPTOR	

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg08482359	PRKAR1B	0.19	cg225 12670	RPS6KA1	0.9
cg08540945		0.34	cg22513455	B4GALT4	0.81
cg08688907		0.74	cg22796704	ARHGAP2	0.06
cg08945443	ZMYND17	0.86	cg23001918		0.15
cg09025210	ZBTB38	0.11	cg23091758	NR1P3	0.22
cg09197075	SNN	0.33	cg23500537		0.19
cg09651136	PKM2	0.37	cg23606718	FAM123C	0.34
cg10274029		0.9	cg23731272	SMAD3	0.74
cg10493055	TSPAN5	0.81	cg23744638		0.86
cg10501210		0.06	cg23876292	LMAN1	0.11
cg10688297	IQCE;IQCE	0.15	cg24079702	FLH2	0.33
cg10909506	ORMDL3	0.22	cg24376214	RASGRP4	0.37
cg11067179	CD248	0.19	cg24735235	FLG37453	0.9
cg11685391	KCN1P2	0.34	cg25020550	TG	0.81
cg12966876	ACOXL	0.74	cg25130381	SLC9A1	0.06
cg13001142	STXBP5	0.86	cg25289028	LRFN3	0.15
cg13553498	CLEC2D	0.11	cg254 10668	RPA2	0.22
cg13608166	NAFTc1	0.33	cg25428494	HPSE	0.19
cg14361627	KLF14	0.37	cg25478614	SST	0.34
cg14518178		0.9	cg25521400	B3GNT2	0.74
cg14556683	EPHX3	0.81	cg25639084		0.86
cg14692377	SLC6A4	0.06	cg26091609	CTLA4	0.11
cg14969094		0.15	cg26164712	CXCR5	0.33
cg15535471	H1PK2	0.22	cg26485825	LAMA3	0.37
cg15867698	ACTN1	0.19	cg26655856	CBFA2T3	0.9
cg15989436		0.34	cg26720010	TBRG4	0.81
cg16054275	F5	0.74	cg18328933	ABHD14A	0.06
cg16419235	PENK	0.86	cg03440846	ACSS2	0.15
cg16549957		0.11	cg10235817	ADRA2C	0.22
cg16662477	RFFL	0.33	cg14826456	ADRB1	0.19
cg16867657	EL0VL2	0.37	cg12111714	ATP8A2	0.34
cg17478979	ZC3H12D	0.9	cg18236477	ATP8A2	0.74
cg17820878	SLC9A1	0.81	cg19945840	B3GALT6	0.86
cg17850367	AB11	0.06	cg17241310	BARHL2	0.11
cg18346531	GAS7	0.15	cg13547237	Bles03	0.33
cg18442362	OGDH	0.22	cg07621046	C10orf82	0.37
cg18473521	H0XC4	0.19	cg08909157	C9orf66	0.9
cg19283806	CCDC102B	0.34	cg08872742	CDH5	0.81
cg19722847	1P08	0.74	cg11136562	CENTD3	0.06
cg19935065	DNTT	0.86	cg19885761	CPLX2	0.15
cg20052760		0.11	cg00399483	DCC	0.22
cg20244489		0.33	cg05822532	ELN	0.19
cg20386303		0.37	cg13921352	FAM19A4	0.34
cg20426994	KLF14	0.9	cg00201234	FBLN2	0.74
cg20723792	FAM53B	0.81	cg00107187	FLJ42486	0.86
cg20822990	ATP13A2	0.06	cg03734874	FLJ42486	0.11
cg21097090	TNFA1P8	0.15	cg25044651	FLJ90650	0.33
cg21139312	MS12	0.22	cg11981599	GATA4	0.37
cg21593149	MAEA	0.19	cg13434842	GATA4	0.9
cg21912203	CACHD1	0.34	cg24646414	GATA4	0.81
cg22016779	DNER	0.74	cg02844545	GCM2	0.06
cg22158769	LOC375196	0.86	cg00059225	GLRA1	0.15
cg22213242	CD248	0.11	cg15425280	GRIA2 (GLUR2)	0.22
cg22285878	KLF14	0.33	cg25148589	GRIA2 (GLUR2)	0.19
cg22454769	FHL2	0.37	cg16464322	HNRP	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg06760035	HOXB4	0.19	cg26933021		0.9
cg06291867	HTR7	0.34	cg20822990	ATP13A2	0.81
cg24826867	IRF8	0.74	cg07312601	MRT04	0.06
cg14614211	IRXL1	0.86	cg09993145	RUNX3	0.15
cg20792062	KCNA5	0.11	cg23605843		0.22
cg27409364	KCNC1	0.33	cg25410668	RPA2	0.19
cg06572160	KCNC3	0.37	cg17879376	BAI2	0.34
cg27553955	KCNG3	0.9	cg14962509	TFAP2E	0.74
cg27320127	KCNK12 (THIK-2)	0.81	cg24375409	EPHA10	0.86
cg12782180	LEP	0.06	cg22851420	HPCAL4	0.11
cg19594666	LEP	0.15	cg24107728	LRP8;LRP8;LRP8;LRP8	0.33
cg08468689	LGP1	0.22	cg14614643		0.37
cg19831077	LOC349136	0.19	cg00257455		0.9
cg10031651	LRRC2	0.34	cg23045908	PDE4B	0.81
cg15784615	LTBR	0.74	cg15201877	PTGER3	0.06
cg20134215	MCHR2	0.86	cg18933331		0.15
cg13603171	MOXD1	0.11	cg05675373	KCNC4;KCNC4;KCNC4	0.22
cg23290344	NEF3	0.33	cg19269039	DENND2D;DENND2D	0.19
cg02994956	NEFH	0.37	cg16008966		0.34
cg12799895	NPTX2	0.9	cg14565725	TBX15;TBX15	0.74
cg25511429	NRN1	0.81	cg05940231	TBX15	0.86
cg20366906	PCDH8	0.06	cg03984502	RORC	0.11
cg00911351	PCDHGB4	0.15	cg16599143	FMO3;FMO3	0.33
cg23563234	PCDHGB7	0.22	cg02901139	NENF;NENF	0.37
cg07408456	PGLYRP2	0.19	cg11298786	WNT3A	0.9
cg02154186	PNMA2	0.34	cg09809672	EDARADD	0.81
cg24199834	POU4F2	0.74	cg05940691	WDR64	0.06
cg23282949	RENBP	0.86	cg12869659	ZNF238	0.15
cg13614181	RGC32	0.11	cg23028740	C1orf100	0.22
cg06156376	SHOX2	0.33	cg10959651	RSAD2	0.19
cg06092815	SKIP	0.37	cg00522231	ITGB1BP1;ITGB1BP1	0.34
cg21992250	SLC15A3	0.9	cg01752203		0.74
cg16232126	SLC5A7	0.81	cg23643435	GALM;GALM	0.86
cg15201635	SMPD3	0.06	cg01243072		0.11
cg02008154	TBX20	0.15	cg07589899		0.33
cg01293143	TCEA2	0.22	cg10855531		0.37
cg13282837	TCL1A	0.19	cg22943590		0.9
cg27210390	TOM1L1	0.34	cg23398076	MEIS1	0.81
cg07533148	TRIM58 (BIA2)	0.74	cg12082609	MEIS1	0.06
cg15747595	TSPYL5	0.86	cg01447660		0.15
cg12457773	VMP	0.11	cg02085953	ARID5A	0.22
cg20616414	WNK2	0.33	cg15149655	VWA3B;VWA3B	0.19
cg14456683	ZIC1	0.37	cg22809047	RPL31;RPL31;RPL31	0.34
cg03975694	ZNF540	0.9	cg06639320	FHL2;FHL2;FHL2;FHL2	0.74
cg27389185	ZNF540	0.81	cg22454769	FHL2;FHL2;FHL2;FHL2	0.86
cg05508084	ZNF667	0.06	cg00017842		0.11
cg19246110	ZNF671	0.15	cg23606718	FAM123C	0.33
cg03734874	FLG42486	0.22	cg22061831	ARHGEF4;ARHGEF4	0.37
cg27320127	KCNK12	0.19	cg12757011	TBR1	0.9
cg07533148	TRIM58	0.34	cg01620164	FIGN	0.81
cg19885761	CLPX2	0.74	cg12105450	CASP10;CASP10;CASP10	0.06
cg00911351	PCDHGB4	0.86	cg00760938		0.15
cg15425280	GRIA2	0.11	cg23077820	PAX3	0.22
cg13547237	Bles03	0.33	cg08166272	IRS1	0.19
cg12140144	FLJ42875;PRDM16;FLJ42875;PRDM16	0.37	cg10523019	RHBDD1;RHBDD1	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg23462687	HDAC4	0.19	cg16281600	GRIA1;GRIA1	0.9
cg20669012		0.34	cg03555227	RANBP17	0.81
cg03183882	SH3BP5;SH3BP5	0.74	cg14345676	HRH2;HRH2;HRH2	0.06
cg15910502	KAT2B	0.86	cg14314729		0.15
cg12941369	PDCD6IP;PDCD6IP;PDCD6IP	0.11	cg23375552		0.22
cg02244028	SCN11A	0.33	cg16867657	ELOVL2	0.19
cg24888989	KIF15;KIF15;KIAA1143	0.37	cg21572722	ELOVL2	0.34
cg00702638	KIF15;KIF15;KIAA1143	0.9	cg01527307		0.74
cg07303143	KIAA1143;KIF15	0.81	cg10699171		0.86
cg19381811	UBA7	0.06	cg06493994	SCGN;SCGN	0.11
cg03019000	TEX264;TEX264;TEX264	0.15	cg02281167	TRIM15	0.33
cg01844642	GPR62	0.22	cg03771840	TRIM15	0.37
cg04474832	ABHD14B	0.19	cg06685111	HCG18;TRIM39;TRIM39;HCG18;TRIM39	0.9
cg03607117	SFMBT1;SFMBT1;SFMBT1	0.34	cg06462220	KIAA1949;KIAA1949	0.81
cg22264409	PROK2;PROK2	0.74	cg08420066	KIAA1949;KIAA1949	0.06
cg11205552	BOC	0.86	cg21467614	TNF	0.15
cg17321954	STXBP5L	0.11	cg12753631		0.22
cg06796779	GATA2;GATA2;GATA2	0.33	cg18501647	PRRT1	0.19
cg18303397	IFT122	0.37	cg04576021	HLA-DOB	0.34
cg14423778	MBNL1	0.9	cg10192196	PSMB8;PSMB8;PSMB8	0.74
cg15277914	DHX36;DHX36	0.81	cg18468088		0.86
cg07553761	TRIM59	0.06	cg03894990	SPATS1	0.11
cg06737494	GHSR;GHSR	0.15	cg01740766		0.33
cg01059398	TNFSF10	0.22	cg16255583		0.37
cg26824216	RTP4	0.19	cg04642300	ARMC2	0.9
cg07110949	MAEA;MAEA	0.34	cg17266282		0.81
cg23239150	EVC	0.74	cg07095347		0.06
cg21254939	SORCS2	0.86	cg16333846		0.15
cg23995914	ZNF518B	0.11	cg13221458	TIAM2;TIAM2;TIAM2	0.22
cg23836737		0.33	cg05468948	SOD2;SOD2;SOD2	0.19
cg10699857		0.37	cg00795927		0.34
cg05024939		0.9	cg08911208	PRKAR1B	0.74
cg05106770		0.81	cg22372849	CYP2W1	0.86
cg01511232		0.06	cg16012294		0.11
cg25148589	GRIA2;GRIA2;GRIA2;GRIA2;GRIA2	0.15	cg27009703	MAD1L1;MAD1L1;MAD1L1	0.33
cg24843443	DDX60L	0.22	cg11671968	HOXA9	0.37
cg03364683	DDX60L	0.19	cg26312920	FKBP14	0.9
cg21815258	CLPTM1L	0.34	cg19663246	NEUROD6;NEUROD6	0.81
cg12608692	MYO10	0.74	cg14396995		0.06
cg20755989		0.86	cg18442362	GLI3	0.15
cg12238343	RXFP3	0.11	cg09748749	OGDH;OGDH;OGDH	0.22
cg02328239	GDNF	0.33	cg23857078	ASL;ASL;ASL;ASL	0.19
cg21878650	ADAMTS6	0.37	cg21743182		0.34
cg26921969		0.9	cg09436502	CDK14	0.74
cg10837404	DCP2	0.81	cg14175438	PDK4	0.86
cg01883408	CCNI2	0.06	cg20665157	FAM3C;FAM3C	0.11
cg06448705	TRPC7;TRPC7;TRPC7;TRPC7;TRPC7;TRPC7	0.15	cg21184711	CADPS2;CADPS2;CADPS2	0.33
cg16983159	TMEM173	0.22	cg02383785	CADPS2;CADPS2;CADPS2	0.37
cg11006267		0.19	cg02821342	MKLN1;FLJ43663	0.9
cg21874213	NRG2;NRG2;NRG2;NRG2	0.34	cg20397034	MKLN1;FLJ43663	0.81
cg26843711	PCDHGA4;PCDHGA6;PCDHGA1;	0.74	cg08280936	ZNF783	0.06
	PCDHGA5;PCDHGB1;PCDHGA6;	0.86	cg18769120		0.15
	PCDHGA3;PCDHGA2;PCDHGB2;	0.11	cg26101086	ARHGEF10	0.22
	PCDHGB3	0.33	cg19859445		0.19
cg08587542	KIAA0141;KIAA0141	0.37	cg07502389	NEFM;NEFM	0.34

Your CpG Values

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CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg18267374	NEFM;NEFM;NEFM	0.19	cg21907579	TBX5;TBX5;TBX5;TBX5	0.9
cg00582628	RGS20	0.34	cg10281002	TBX5;TBX5	0.81
cg23710218	MSC;MSC	0.74	cg07172885	ZNF268;ZNF268;ZNF268;ZNF268;ZNF268;	0.06
cg19497517	PLEC1;PLEC1;PLEC1	0.86		ZNF268;ZNF268;ZNF268;ZNF268	0.15
cg13586038	SMARCA2;SMARCA2	0.11	cg20404336	ZNF268;ZNF268;ZNF268;ZNF268;ZNF268;	0.22
cg07158339	FXN;FXN;FXN	0.33		ZNF268;ZNF268;ZNF268;ZNF268	0.19
cg24046474	RPL12;LRSAM1;LRSAM1	0.37	cg18582260	PARP4	0.34
cg14059835	C9orf78;USP20;USP20;USP20	0.9	cg06648759		0.74
cg10570177	C9orf7;C9orf7	0.81	cg23357533	C13orf15	0.86
cg13649056		0.06	cg20102280	HTR2A;HTR2A	0.11
cg13734401		0.15	cg13767001	MIR759	0.33
cg06231995		0.22	cg23389651		0.37
cg14411282	ZMYND19	0.19	cg09646392	TNFSF13B;TNFSF13B	0.9
cg12530994	AKR1C3	0.34	cg00593462		0.81
cg23754392	BMI1	0.74	cg07115626		0.06
cg06908778	SPAG6;SPAG6	0.86	cg24058132	GALC;GALC;GALC;GALC	0.15
cg22796704	ARHGAP22	0.11	cg13027206	CCDC88C	0.22
cg18738190	CHST3	0.33	cg15480367	CHGA;CHGA	0.19
cg25427880		0.37	cg12177001	IFI27;IFI27	0.34
cg09671951	C10orf26	0.9	cg23709172	SLC25A29	0.74
cg06888746	SH3PXD2A	0.81	cg14334310		0.86
cg24838825		0.06	cg04875128	OTUD7A	0.11
cg13848598	ADRB1	0.15	cg03361973	SEMA6D;SEMA6D;SEMA6D;SEMA6D;	0.33
cg07906193		0.22		SEMA6D;SEMA6D	0.37
cg12776156	TSPAN4	0.19	cg16717122	SCG3;SCG3;SCG3;SCG3	0.9
cg05928581	LSP1	0.34	cg04858164	TCF12;TCF12;TCF12;TCF12	0.81
cg17627559	TSPAN32;C11orf21;C11orf21	0.74	cg25005357	TCF12;TCF12;TCF12;TCF12	0.06
cg23091758	NRIP3	0.86	cg08454546	FAM81A	0.15
cg10825530	SOX6;SOX6;SOX6;SOX6	0.11	cg15188939	ARIH1	0.22
cg20654468	LPXN;LPXN	0.33	cg20540209	ISL2	0.19
cg26552743		0.37	cg05542681	FBXL16	0.34
cg21992250	SLC15A3;SLC15A3	0.9	cg08949164	BAIAP3	0.74
cg15015340	DAGLA	0.81	cg09183146	UNKL	0.86
cg22843803		0.06	cg00454305	UNKL	0.11
cg02532488	EHBP1L1	0.15	cg02871659	SNHG9	0.33
cg13547237	C11orf68;DRAP1;C11orf68	0.22	cg00991848	SNHG9	0.37
cg05496363	CD248	0.19	cg02331561	ABCA17P;ABCA3	0.9
cg20063906		0.34	cg26974444	SNX29	0.81
cg12328429	ME3;ME3;ME3	0.74	cg06112560	PLK1	0.06
cg25969122		0.86	cg10917602	HSD3B7	0.15
cg18633600	LRTM2;LRTM2;CACNA2D4;LRTM2	0.11	cg09155044	VKORC1;VKORC1	0.22
cg08622677	PRMT8	0.33	cg04031656	CHD9	0.19
cg25719851	PTPN6	0.37	cg03746976	C16orf57	0.34
cg13828440	KLRD1;KLRD1;KLRD1	0.9	cg18693704	CMTM4;CMTM4	0.74
cg26986871		0.81	cg00658652		0.86
cg21747310		0.06	cg03991512	LDHD;LDHD	0.11
cg13909661	SSPN;SSPN	0.15	cg03486383	SLC7A5	0.33
cg26311454		0.22	cg02228185	ASPA;ASPA	0.37
cg00753885		0.19	cg23668631	CAMKK1;CAMKK1	0.9
cg18573383	KCN2;KCN2;KCN2;KCN2;KCN2;KCN2	0.34	cg14522800	ZMYND15;ZMYND15	0.81
cg15405572	NAV3	0.74	cg25135555	GAS7;GAS7;GAS7;GAS7	0.06
cg08993878		0.86	cg13029847	SEZ6;SEZ6	0.15
cg22827210	CHST11	0.11	cg06874016	NKIRAS2	0.22
cg03670162		0.33	cg04267101	EPN3	0.19
cg04596060	RBM19;RBM19;RBM19	0.37	cg22507023	CUEDC1	0.34

Your CpG Values



CpG	Gene	Your Beta Value	CpG	Gene	Your Beta Value
cg13093111	ARSG	0.19	cg25436157	C18orf45	0.9
cg13683374	GPR142	0.34	cg24217948	SETBP1;SETBP1	0.81
cg10644544	TRIM47	0.74	cg17589341	SLC14A1	0.06
cg03643998	C1QTNF1	0.86	cg17243289	SMAD2;SMAD2;SMAD2	0.15
cg11620135	BAIAP2	0.11	cg12459502	BCL2	0.22
cg25436157	C18orf45	0.33	cg26005082	MIR7-3;C19orf30	0.19
cg24217948	SETBP1;SETBP1	0.37	cg11766468	EVI5L;EVI5L	0.34
cg17589341	SLC14A1	0.9	cg10586358	SYCE2	0.74
cg17243289	SMAD2;SMAD2;SMAD2	0.81	cg14556683	EPHX3;EPHX3	0.86
cg12459502	BCL2	0.06	cg18335931	MAST3	0.11
cg26005082	MIR7-3;C19orf30	0.15	cg10498798	FFAR1	0.33
cg11766468	EVI5L;EVI5L	0.22	cg27212234	ATP4A	0.37
cg10586358	SYCE2	0.19	cg17110586		0.9
cg14556683	EPHX3;EPHX3	0.34	cg21944491	LTBP4;LTBP4;LTBP4	0.81
cg18335931	MAST3	0.74	cg21940708	PRMT1;C19orf76	0.06
cg10498798	FFAR1	0.86	cg06458239	ZNF549	0.15
cg27212234	ATP4A	0.11	cg10729426	ZNF549	0.22
cg17110586		0.33	cg21911021	ZIK1	0.19
cg21944491	LTBP4;LTBP4;LTBP4	0.37	cg19702785	KCNS1	0.34
cg21940708	PRMT1;C19orf76	0.9	cg07547549	SLC12A5;SLC12A5	0.74
cg06458239	ZNF549	0.81	cg12303084	ZMYND8;ZMYND8;ZMYND8	0.86
cg10729426	ZNF549	0.06	cg17274064	ERG;ERG	0.11
cg21911021	ZIK1	0.15	cg09428349	C2CD2;C2CD2	0.33
cg19702785	KCNS1	0.22	cg10636297	AGPAT3;AGPAT3	0.37
cg07547549	SLC12A5;SLC12A5	0.19	cg12373771	CECR6;CECR6	0.9
cg12303084	ZMYND8;ZMYND8;ZMYND8	0.34	cg05442902	MGC16703;P2RX6;P2RX6	0.81
cg17274064	ERG;ERG	0.74	cg16612562	RASL10A;RASL10A	0.06
cg09428349	C2CD2;C2CD2	0.86	cg19015086	SEC14L2	0.15
cg10636297	AGPAT3;AGPAT3	0.11	cg21205978	APOL6	0.22
cg12373771	CECR6;CECR6	0.33	cg01949403	APOL3;APOL3	0.19
cg05442902	MGC16703;P2RX6;P2RX6	0.37	cg08415592	APOL1	0.34
cg16612562	RASL10A;RASL10A	0.9	cg19853760	LGALS1;LGALS1	0.74
cg19015086	SEC14L2	0.81	cg25459323	SEPT3;SEPT3	0.86
cg21205978	APOL6	0.06	cg27187881	NAGA;NAGA	0.11
cg01949403	APOL3;APOL3	0.15	cg00343092	TSPO;TSPO	0.33
cg08415592	APOL1	0.22	cg00347775	MPPED1	0.37
cg04031656	CHD9	0.19	cg27131176	C22orf26;LOC150381	0.9
cg03746976	C16orf57	0.34	cg22982767	LOC150381	0.81
cg18693704	CMTM4;CMTM4	0.74	cg07016730	SBF1	0.06
cg00658652		0.86	cg01892695		0.15
cg03991512	LDHD;LDHD	0.11			0.22
cg03486383	SLC7A5	0.33			0.19
cg02228185	ASPA;ASPA	0.37			0.34
cg23668631	CAMKK1;CAMKK1	0.9			0.74
cg14522800	ZMYND15;ZMYND15	0.81			0.86
cg25135555	GAS7;GAS7;GAS7;GAS7	0.06			0.11
cg13029847	SEZ6;SEZ6	0.15			0.33
cg06874016	NKIRAS2	0.22			0.37
cg04267101	EPN3	0.19			0.9
cg22507023	CUEDC1	0.34			0.81
cg13093111	ARSG	0.74			0.06
cg13683374	GPR142	0.86			0.15
cg10644544	TRIM47	0.11			0.22
cg03643998	C1QTNF1	0.33			0.19
cg11620135	BAIAP2	0.37			0.34