Supplementary Table 1. Beta coefficients and P values after log-linear model of the 140 CpGs present in the $12^{\rm th}$ iteration of the RF algorithm output and their Genetic details

CpGs	CHR ^{\$}	UCSC* chromosomal location	Gene	UCSC* Gene Region	USCS* CpG Island Relation	β-coefficients	P-Value
cg00057114	1	6344362	ACOT7	Body	S_Shelf	1.57	0.032
cg00069771	1	233089275	C1orf57	Body	S_Shelf	-0.11	0.031
cg00349607	1	39546688	MACF1	TSS1500	N_Shore	1.44	0.026
cg00362285	1	236444212	ERO1LB	Body	N_Shore	1.38	0.026
cg00388605	1	89150448	PKN2	Body	Island	-1.55	0.028
cg02641560	1	24828811	RCAN3	TSS1500	N_Shore	2.68	0.011
cg00158434	2	73871782	ALMS1P	TSS1500		1.69	0.017
cg00541803	2	120301835	PCDP1	TSS1500	Island	-2.02	0.031
cg00900242	2	200929967				1.92	0.043
cg01158447	2	190445074	SLC40A1	Body	N_Shore	-1.40	0.016
cg02237186	2	10263043	RRM2;RRM2	Body	Island	1.45	0.039
cg06334988	2	220095083	ANKZF1;ATG9A	Body; TSS1500	S_Shore	-1.12	0.009
cg00071869	3	141599902	ATP1B3	Body	S_Shelf	-1.17	0.023
cg00124095	3	72305082				1.10	0.016
cg00203564	3	176133845				-0.80	0.038
cg00742851	3	3843342	LRRN1	5'UTR	S_Shore	0.80	0.034
cg09306365	3	61547944	PTPRG	1stExon; 5'UTR	Island	-1.17	0.047
cg00050654	4	4576493			N_Shore	-1.64	0.014
cg01214900	4	66784395				0.84	0.104
cg01651499	4	156646379	GUCY1A3	Body		-0.99	0.038
cg02654265	5	137086868			N_Shelf	1.36	0.017
cg11861905	5	176045983			N_Shore	-0.89	0.038
cg17602756	5	179246001	SQSTM1	5'UTR	N_Shore	-1.92	0.028
cg00252472	6	150739173				0.96	0.026
cg02077766	6	42891866	PTCRA	Body	N_Shore	2.26	0.017
cg03473348	6	33160976	COL11A2	TSS1500	Island	-1.84	0.017
cg24303123	6	3076314	RIPK1	TSS1500		1.54	0.016
cg00035220	7	158197980	PTPRN2	Body	N_Shore	0.96	0.034
cg00576402	7	77167212	PTPN12	TSS200; Body	Island	-0.55	0.042
cg00718175	7	158807362	LOC154822	Body		1.53	0.043
cg02433979	7	128824829			N_Shelf	1.07	0.024
cg05345249	7	80291832	CD36	Body		1.30	0.016
cg11570082	7	116899446				1.49	0.014
cg00021978	8	105476917	DPYS	Body	N_Shore	-1.06	0.042
cg00397910	8	80522577	STMN2	TSS1500	N_Shore	1.33	0.016
cg01049274	8	146228340	ZNF252;C8orf77	TSS200; Body	Island	-1.02	0.046
cg01214321	8	26435429	DPYSL2	5'UTR; 1stExon	Island	-1.79	0.038

cg02049232	8	145556129	SCRT1	3'UTR	Island	-1.65	0.024
cg02098905	8	119759090	JOH! I	301R	isiana	-1.02	0.026
cg02919936	8	70982285	PRDM14	5'UTR	Island	-0.57	0.038
cg14839837	8	1834174	ARHGEF10	Body	S_Shelf	1.77	0.025
cg00193668	9	35814283	HINT2	Body	N_Shore	2.85	0.016
cg00306063	9	92289643	LOC100129066	Body	N_Shore	0.95	0.043
cg00077547	10	63170372	TMEM26	Body	11_bhore	-1.55	0.043
cg00198759	10	81444321	LOC650623	Body	Island	1.18	0.047
cg02765031	10	85462980	20 0030023	Body	isiana	1.71	0.009
cg03049303	10	103816154	C10orf76	TSS1500	S_Shore	1.55	0.028
cg04807739	10	88779914	FAM25A	TSS200	S_Shore	-2.03	0.024
cg08353444	10	60506653	BICC1	Body		1.18	0.026
cg20077343	11	1036307	MUC6	Body	N_Shore	-1.62	0.017
cg00224126	12	62861053	MON2	1stExon	S_Shore	-2.18	0.024
cg00369908	12	6772357	ING4	TSS200	Island	1.29	0.028
cg00665405	12	50899548	DIP2B	Body	S_Shore	-1.45	0.022
cg00667315	12	295951		•	N_Shelf	2.03	0.021
cg02583247	12	4555311	FGF6	TSS1500	S_Shore	1.88	0.016
cg02767665	12	129788385	TMEM132D	Body	S_Shore	1.67	0.016
cg04797820	12	129460559	GLT1D1	Body		-1.09	0.016
cg11912049	12	50361401		·	Island	-1.38	0.016
cg00949022	13	73356224	PIBF1;DIS3	TSS200; 1stExon;	Island	0.77	0.038
cg01560119	13	50025780	SETDB2	5'UTR 5'UTR; 1stExon		-0.87	0.026
cg04850479	13	113811569	PROZ	TSS1500	S_Shelf	2.72	0.014
cg05411056	13	112852819			S_Shelf	2.27	0.014
cg22148297	13	99910587	GPR18;UBAC2	1stExon; Body; 5'UTR		0.49	0.037
cg00722180	14	73525139	RBM25	TSS200	Island	1.29	0.038
cg01802073	14	54976343	CGRRF1	TSS1500	Island	1.80	0.034
cg06920136	14	21945858	RAB2B;TOX4	TSS1500; Body	S_Shore	1.54	0.026
cg00229754	15	25018302			Island	-0.55	0.149
cg02203881	15	42386909	PLA2G4D	TSS200		0.73	0.047
cg04980849	15	45671347	LOC145663;GATM	Body; TSS1500	S_Shore	-1.53	0.017
cg00024494	16	73096842			N_Shore	-0.50	0.041
cg00073380	16	5122085	ALG1	Body	Island	-0.78	0.038
cg00161345	16	66866226	NAE1	TSS1500	S_Shore	2.19	0.014
cg00797821	16	49888944			Island	-1.20	0.038
cg00818680	16	68881254	TMCO7	Body	S_Shelf	0.65	0.026
cg00933210	16	3233117			Island	-2.13	0.045
cg01452911	16	86512095				1.72	0.016
cg03910196	16	2256384	MLST8	Body	S_Shore	1.55	0.031
cg05839818	16	85634031				2.56	0.017
cg00354884	17	1057541	ABR	Body; 5'UTR		1.75	0.016

cg00927777	17	1960199	HIC1	Body	Island	-0.92	0.026
cg00320243	19	18314691	RAB3A	5'UTR	Island	-1.91	0.038
cg00337020	19	2241089	SF3A2	5'UTR	S_Shelf	1.04	0.028
cg00201670	21	46221900	UBE2G2	TSS200	Island	-1.95	0.043
cg00015530	8	145052950	PARP10	Body	S_Shore	0.32	0.244
cg00019877	9	115773738	LOC169834	5'UTR	N_Shore	1.82	0.024
cg00546248	1	2412743	PLCH2	Body	S_Shore	-0.45	0.186
cg00778190	1	95068029				0.54	0.101
cg01059116	1	54163070	GLIS1	5'UTR		0.31	0.048
cg00164941	2	73523056			S_Shelf	-0.59	0.11
cg00729049	2	171679402	GAD1	Body	Island	-1.10	0.031
cg00405769	4	187539852	FAT1	Body		-0.57	0.075
cg00866215	4	54956702			N_Shore	0.64	0.123
cg05684300	4	102267366	PPP3CA	Body	N_Shore	0.94	0.061
cg00099094	5	2334514				-0.23	0.189
cg00106739	5	177657311	AGXT2L2	Body	N_Shore	0.97	0.068
cg00580497	5	1153746			N_Shore	0.81	0.122
cg00043444	6	134384706				-0.23	0.633
cg00155619	6	33173278	HSD17B8	Body	S_Shore	0.97	0.08
cg00491418	6	112575482	LAMA4	5'UTR	Island	-1.72	0.066
cg01427769	6	31830600	NEU1	5'UTR; 1stExon	Island	-2.0	0.016
cg12427444	6	31923064	RDBP	Body	N_Shelf	-0.49	0.497
cg00036614	7	47093842			S_Shore	0.33	0.378
cg00064840	7	4784485	FOXK1	Body	N_Shore	1.42	0.068
cg00106345	7	27138396			S_Shore	-0.39	0.213
cg00129269	7	127228222	ARF5	TSS1500	Island	-0.92	0.068
cg00301256	7	82653693	PCLO;PCLO	Body		-0.57	0.264
cg02474076	7	127223255	GCC1	Body	N_Shore	1.47	0.126
cg00161683	8	95563352	KIAA1429	Body	N_Shelf	1.89	0.027
cg00166750	8	102138594			Island	-1.21	0.06
cg00328935	8	142517474	FLJ43860	TSS200		-0.34	0.359
cg00405554	8	65490897	LOC401463	TSS1500	N_Shore	1.34	0.084
cg00449899	8	53626436	RB1CC1	5'UTR	Island	-1.34	0.039
cg00592695	8	134545685	ST3GAL1	5'UTR		-0.92	0.094
cg00598335	8	11627530	NEIL2	5'UTR; 1stExon	Island	0.43	0.489
cg00661018	8	145769086	KIAA1688	Body	N_Shore	1.90	0.053
cg00556719	9	108457484	TMEM38B	Body	Island	1.98	0.075
cg00152946	10	130338781			Island	-1.53	0.068
cg00338996	11	45828080	SLC35C1	Body	S_Shore	0.75	0.06
cg00478064	11	118271813	ATP5L	TSS1500	N_Shore	1.61	0.09
cg12079699	11	131564481	NTM	Body	S_Shelf	-0.77	0.101
cg00104348	12	10772424	STYK1	3'UTR		-0.70	0.149
cg00216961	12	6054731	ANO2	Body		-0.91	0.189

cg00240719	12	132169130			N_Shore	-0.71	0.121
cg00216180	13	114778713	RASA3	Body	Island	0.88	0.213
cg00247571	13	50707065			N_Shore	-1.15	0.027
cg01729020	14	55035889	SAMD4A	Body	S_Shore	0.90	0.25
cg00376910	15	89765061	RLBP1	TSS200		-1.12	0.054
cg00153543	16	84259830	KCNG4	Body	S_Shelf	0.54	0.219
cg00261912	16	89777763	C16orf7	Body	Island	0.47	0.054
cg00489954	16	28855387	TUFM	Body	N_Shore	0.001	0.999
cg00701514	16	1183462			S_Shelf	-0.84	0.075
cg00911376	16	84274417	KCNG4	TSS1500	S_Shelf	0.67	0.162
cg02455706	16	84918851	CRISPLD2	Body		-0.78	0.032
cg00416882	18	44495671	PIAS2	Body	N_Shore	0.88	0.075
cg00417323	18	29307959			S_Shelf	0.76	0.093
cg00486358	18	54309442			S_Shelf	0.54	0.178
cg01080902	19	44039870	ZNF575	3'UTR	S_Shore	1.19	0.06
cg00397859	20	13768987	C20orf7	Body	S_Shelf	0.67	0.065
cg00399938	22	20118646	ZDHHC8	TSS1500	Island	0.44	0.25
cg01133262	22	38863905	KDELR3	TSS200	Island	0.87	0.226

CHR^{\$}= chromosome number UCSC*= University of South Carolina