

Supplementary Table S7. Summary of MCAM data in human MDS patients.

ID	FAB	Age (years)	Tissue	Hypermethylation		Hypomethylation		Race	CIMP
				# of genes	(%)	# of genes	(%)		
MDS-07	CMML	70	BM	639	6.6	193	2.0	White	+
MDS-08	RAEB	62	BM	942	9.8	338	3.5	Black	+
MDS-14	RAEB	72	BM	1677	17.4	397	4.1	White	+
MDS-12	RAEB-T	61	BM	674	7.0	373	3.9	White	+
MDS-18	RAEB-T	61	BM	773	8.0	447	4.6	White	+
MDS-20	RAEB-T	67	BM	889	9.2	377	3.9	White	+
MDS-21	RAEB-T	80	BM	1109	11.5	304	3.2	White	+
MDS-15	CMML	61	BM	409	4.2	414	4.3	White	-
MDS-09	CMML	81	BM	383	4.0	361	3.7	White	-
MDS-16	RA	70	BM	205	2.1	627	6.5	White	-
MDS-19	RAEB	56	BM	468	4.9	326	3.4	White	-
MDS-10	RAEB	71	BM	527	5.5	464	4.8	White	-
MDS-13	RAEB	72	BM	332	3.4	392	4.1	White	-
MDS-17	RAEB	73	BM	510	5.3	676	7.0	White	-
MDS-22	RAEB	73	BM	285	3.0	249	2.6	White	-
MDS-24	RAEB	73	BM	315	3.3	418	4.3	White	-
MDS-06	RAEB	75	BM	486	5.0	355	3.7	White	-
MDS-05	RAEB	79	BM	564	5.8	333	3.5	White	-
MDS-11	RAEB	80	BM	351	3.6	442	4.6	White	-