

Supplementary Table S3. Summary of MCAM data in mouse models.

ID	Age (months)	Tissue	Hypermethylation		Hypomethylation		Strain
			# of genes	(%)	# of genes	(%)	
<i>NHD13-7</i>	12.0	BM	497	14.43	190	5.52	FVB
<i>NHD13-8</i>	12.0	BM	490	14.65	204	5.92	FVB
<i>NHD13-9</i>	9.0	BM	548	15.91	236	6.85	FVB
<i>RUNX1</i> (D171N)-6	8.2	Spl	300	8.71	125	3.63	B6
<i>RUNX1</i> (D171N)-13	7.8	Spl	270	7.87	134	3.89	B6
<i>RUNX1</i> (S291fs)-58	7.6	Spl	665	19.30	178	5.17	B6