Supplementary Table1

mutation pattern	full model	independent model
L	1	3
M	(96)	(6, 4, 4)
$ApCpA \rightarrow ApCpA$	(1)	(1, 1, 1)
$ApCpC \rightarrow ApApC$	(2)	(1, 1, 2)
$ApCpG \rightarrow ApApG$	(3)	(1, 1, 3)
$ApCpT \rightarrow ApApT$	(4)	(1, 1, 4)
$CpCpA \rightarrow CpApA$	(5)	(1, 2, 1)
• • •		• • •
$ApCpA \rightarrow ApGpA$	(17)	(2, 1, 1)
• • •		• • •
$TpTpT \to TpGpT$	(96)	(6, 4, 4)