

<u>Tumor</u>	<u>MMHCC</u>	<u>Mut freq¹ ($\times 10^{-4}$)</u>	<u>C to T²</u>	<u>C to G</u>	<u>G to A</u>	<u>A to G</u>
K7	BLL	11.5 (21/18,190)	47.6%	9.5%	23.8%	19.0%
K9	DLBCL	14.5 (12/8,285)	33.3%	16.7%	8.3%	41.7%
WT	spleen	12 (10/8,321)	60.0%	0.0%	10.0%	30.0%

¹ In parentheses is the number of mutations per number of total base pairs sequenced.

² Percentage of mutations representing the indicated mutation.