

Table 8. Results from the Taq-Man assay for the *MEIS1* gene for the leukemias; six samples harbored *MLL* rearrangements, and six did not

Sample	Detector	CT	Mean	DCT	DDCT	Ratio sample/ref.
<i>MLL</i>	MEIS 1	26.31	26.255	13.145	-8.53	369.6
	18S	13.1	13.11			
<i>MLL</i>	MEIS 1	22.61	23.505	10.32	-11.355	2619.4
	18S	13.2	13.185			
<i>MLL</i>	MEIS 1	26.59	26.795	14.86	-6.815	112.6
	18S	11.52	11.935			
<i>MLL</i>	MEIS 1	26.38	26.38	11.44	-10.235	1205.2
	18S	14.55	14.94			
<i>MLL</i>	MEIS 1	26.71	26.68	13.39	-8.285	311.9
	18S	12.9	13.29			
<i>MLL</i>	MEIS 1	25.84	25.42	11.445	-10.23	1200.9
	18S	13.81	13.975			
Other	MEIS 1	34.04	34.115	18.76	-2.915	7.5
	18S	15.37	15.355			
Other	MEIS 1	36.47	36.51	21.895	0.22	0.89
	18S	14.8	14.615			
Other	MEIS 1	39.16	39.16	23.885	2.21	0.22
	18S	15.23	15.275			
Other	MEIS 1	36.28	36.16	20.775	-0.9	1.9
	18S	15.14	15.385			
Other	MEIS 1	35.98	36.2	21.27	-0.405	1.3
	18S	14.8	14.93			
Other, reference	MEIS 1	36.66	36.325	21.675	0	1
	18S	14.63	14.65			