

S2 Table. Ct values of all target genes in serial dilutions of 5 MB clinical samples.

Target gene	1:10		1:100		1:200		1:1000		1:2000		1:10000	
	Mean	CV ^a	Mean	CV	Mean	CV	Mean	CV	Mean	CV	Mean	CV
RLEP	26.12	0.01	29.58	0.01	30.71	0.02	33.43	0.02	34.48	0.01	36.75	0.02
groEL	29.06	0.01	32.82	0.01	33.65	0.01	36.22	0.01	37.59	0.01	ND ^b	-
pra	30.97	0.01	34.61	0.01	35.40	0.01	38.29	0.01	38.45	0.01	ND	-
esxA	31.27	0.01	34.80	0.01	35.93	0.01	38.25	0.01	38.39	0.01	ND	-
HSP18	31.58	0.01	35.07	0.01	36.23	0.02	38.94	0.01	39.14	0.01	ND	-
85B	32.07	0.01	35.67	0.01	36.86	0.02	38.98	0.01	39.42	0.00	ND	-
rpoT	32.17	0.01	35.76	0.01	36.61	0.02	39.01	0.01	ND	-	ND	-
ML0024	32.29	0.01	35.93	0.01	37.07	0.01	38.66	0.00	ND	-	ND	-
ML1545	32.42	0.01	36.07	0.01	37.08	0.01	39.09	0.01	ND	-	ND	-
ML2179	32.61	0.01	36.23	0.01	37.63	0.01	39.28	0.01	ND	-	ND	-
sodA	33.08	0.01	36.85	0.01	38.00	0.01	39.37	0.01	ND	-	ND	-
16SrRNA	33.09	0.01	36.94	0.01	37.72	0.01	ND	-	ND	-	ND	-
TTC	33.37	0.01	37.41	0.02	38.29	0.01	ND	-	ND	-	ND	-
ML0098	33.42	0.01	37.02	0.01	38.03	0.02	ND	-	ND	-	ND	-
AT	33.49	0.01	37.21	0.01	38.16	0.01	ND	-	ND	-	ND	-
MntH	37.35	0.01	ND	-	ND	-	ND	-	ND	-	ND	-
AGT	ND	-	ND	-	ND	-	ND	-	ND	-	ND	-

The sensitivities of 17 target genes were assessed by qPCR using a series of normalized DNA samples from MB patients.

^aCV = Coefficient of variation.

^bND = Not detected.