

Corrections

In the article entitled, “A Melting Curve Analysis–Based PCR Assay for One-Step Genotyping of β -Thalassemia Mutations: A Multicenter Validation” (Volume 13, pages 427–435 of the July 2011 issue of *The Journal of Molecular Diagnostics*), the authors found several small, but obvious, erroneous calculations in Table 2. The authors have since corrected the values and the corrected Table 2 appears below.

The article entitled “Copy Number Gains in 11q13 and 8q34 Are Highly Linked to Prognosis in Cutaneous Malignant Melanoma” (Volume 13, pages 352–358 of the May 2011 issue), contained an error in the chromosomal band for *MYC* in the title and throughout the text. The correct chromosomal band for *MYC* is 8q24. The authors apologize for the error.

Table 2. Calibrated T_m of 24 Mutations Detected Using the MeltPro HBB Assay

Tube	Color	Genotype	T_m (°C)	ΔT_m (3 SD) (°C)*
1	FAM	Wild-type	65.9	NA
		c.316-197C>T	59.9	6.0 (0.42)
	HEX	Wild-type	65.4	NA
		c.-78A>G	57.0	8.4 (0.39)
		c.-79A>G	58.4	7.0 (0.54)
		c.-80T>C or c.-81A>C [†]	62.1	3.3 (0.42)
		c.-82C>A	54.3	11.1 (0.39)
	ROX	Wild-type	67.7	NA
		c.45-46insG	64.6	3.1 (0.45)
		c.48-49insG	66.0	1.7 (0.35)
		c.52A>T	62.2	5.5 (0.54)
	Cy5	Wild-type	67.1	NA
		c.79G>A	61.7	5.4 (0.31)
		c.92 + 1G>T or c.92 + 5G>C [†]	63.4	3.7 (0.33)
		c.91A>G	59.2	7.9 (0.51)
Wild-type		68.4	NA	
2	FAM	c.79G>A	61.4	7.0 (0.36)
		c.84-85insC	64.0	4.4 (0.45)
		Wild-type	71.0	NA
	HEX	c.124-127delTTCT	60.2	10.8 (0.54)
		c.115delA	61.3	9.7 (0.48)
		c.130G>T or c.113G>A [†]	66.3	4.7 (0.48)
		Wild-type	68.1	NA
	ROX	c.216-217insT	63.8	4.3 (0.54)
		c.216-217insA	62.5	5.6 (0.50)
		c.315 + 2delT	48.1	20.0 (0.45)
		c.315 + 5G>C	51.4	16.7 (0.24)
		Wild-type	64.6	NA
	Cy5	c.-123A>T	58.1	6.5 (0.42)
		c.-140C>T	55.7	8.9 (0.33)

* $\Delta T_m = T_m$ (wild-type) – T_m (mutant).

[†]These mutations are judged to be the same type within pairs using the MeltPro HBB assay. NA, not applicable.