

Correction

Correction: Clinical Correlates of *NRAS* and *BRAF* Mutations in Primary Human Melanoma

In this article (Clin Cancer Res 2011;17:229–35), which was published in the January 15, 2011 issue of *Clinical Cancer Research* (1), the alignment of Table 4 is incorrect. The correct table is shown below.

Table 4. Uncommon genotypes

	<i>NRAS</i>	<i>BRAF</i>	Age, y	Gender	Breslow, mm	Clark	Mitotic index, figures/mm ²	Ulceration
Double mutant	A182G	TG1799AA	58	M	0.85	3	1	No
	C181A	T1799A	58	M	1.46	4	1	No
	A182G	GT1798AA	57	F	1.2	4	3	Yes
Only mutated DNA	C181A	Wild type	49	F	3.9	5	5	No
	A182G	Wild type	51	M	2.2	4	2	No
	Wild type	T1799A	35	F	3.1	3	3	No
	Wild type	GT1798AA	77	M	3.8	4	9	No
	Wild type	T1799A	57	M	5	4	16	Yes

References

1. Ellerhorst JA, Greene VR, Ekmekcioglu S, Warneke CL, Johnson MM, Cooke CP, et al. Clinical correlates of *NRAS* and *BRAF* mutations in primary human melanoma. Clin Cancer Res 2011;17:229–35.

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