

# Oncogenic *BRAF* mutation in cutaneous melanoma in Argentina

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## Background

*BRAF* is a critical protein in the mitogen-activated protein kinase (MAPK) pathway, and is mutated in approximately 50% of melanoma patients<sup>1</sup>. A single substitution (V600E) account for 80% of *BRAF* mutations<sup>2-3</sup>. Trunk location, age at diagnosis of primary tumor  $\leq 50$  years and the absence of chronic sun damage predict relationship between primary melanoma and *Braf* mutation.<sup>6,7</sup>

## Objective

The aim of this study was to assess the frequency of oncogenic *BRAF* mutations in cutaneous melanoma and to correlate *BRAF* status with clinic-pathological features in Argentinian melanoma patients.

**Patients and Methods:** Tumor tissue from 92 consecutive patients with primary/metastatic cutaneous melanoma, from 12/2011 to 3/2013 was analyzed in 2 referral centers using real time polymerase chain reaction (RT-PCR) assays for *BRAF* mutation (cobas® 4800 System, v2.0)<sup>4</sup>. Clinical (age, gender, ethnics, primary tumor location) and pathological (Breslow, ulceration, histopathological subtype, mitotic rate, AJCC stage) features were correlated with *BRAF* mutational status

**Inclusion criteria:** > 18 yrs, histology confirmed cutaneous melanoma (any TNM stage), available tissue for mutational analysis (primary, lymph node or metastasis).

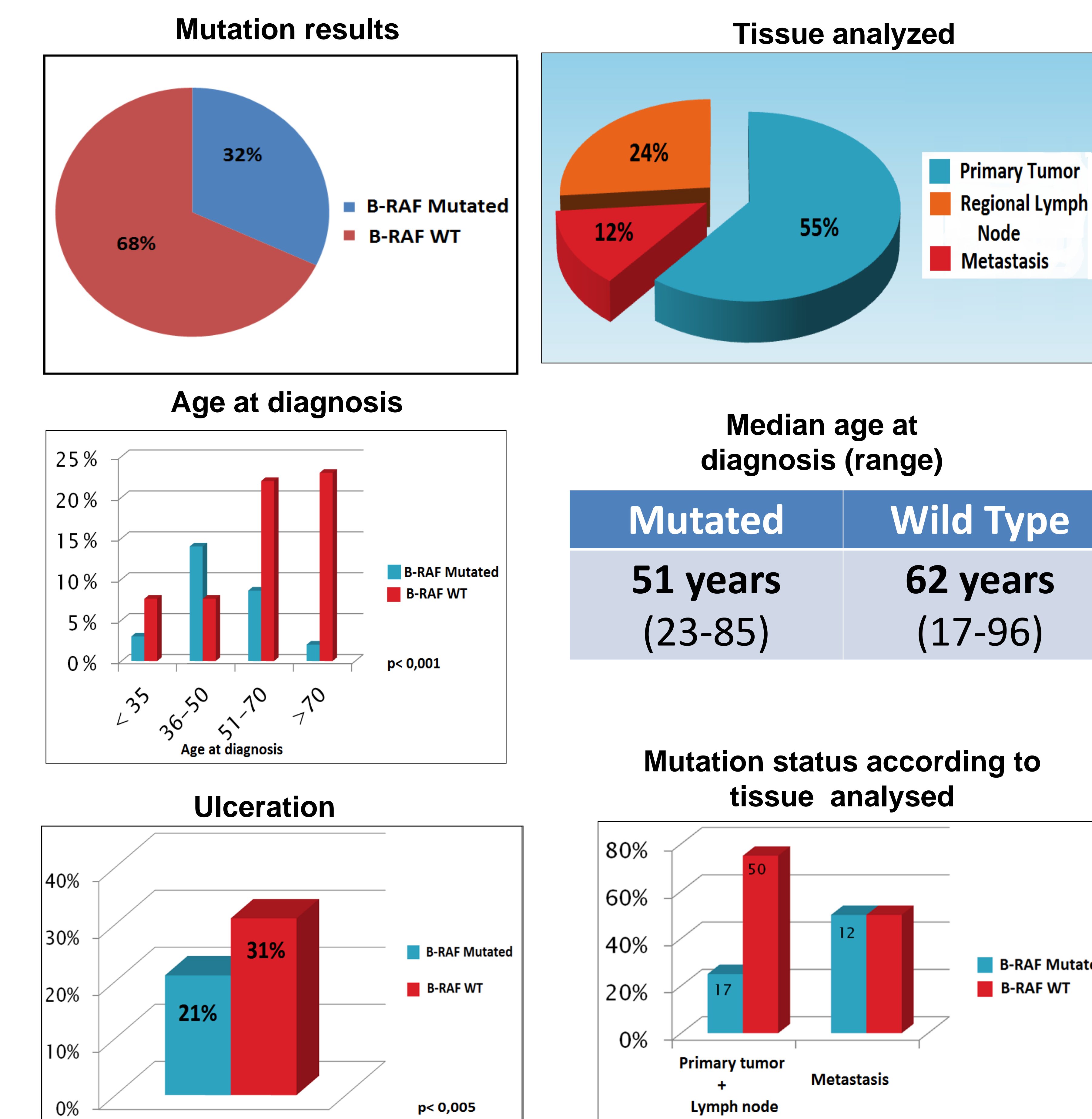
**Exclusion criteria:** Uveal, mucosal or acral melanoma.

## Results

Patient/Tumor Characteristics	N= 92 (%)
<b>Age at diagnosis. Median (range)</b>	58 (17-96) < 35: 17 (18%) 36-50: 20 (31%) 51-70: 34 (36%) >71: 26 (28%)
<b>Sex</b>	47 Males (51%)
<b>Gender</b>	92 caucasian (100%)
<b>Primary tumor location</b>	35 Trunk/Back (38%) 31 Extremities (33%) 14 Other (15%) 10 Head and Neck (10%) 2 Unknown (2%)
<b>Breslow (mm)</b>	$\leq 1$ : 23 (25%) 1,01 a 2: 18 (19%) 2,01 a 3: 10 (10%) 3,01 a 4: 7 (7%) $\geq 4,01$ : 15 (16%) Unknown: 19 (20%)

Patient/Tumor Characteristics	N= 92 (%)
<b>Mitotic Index (per mm<sup>2</sup>)</b>	$\leq 1$ mm <sup>2</sup> : 31 (33%) $\geq 1$ mm <sup>2</sup> : 32 (34%) Unknown: 29 (31%)
<b>Ulceration</b>	Yes: 18 (19%) No: 45 (48%) Unknown: 29 (31%)
<b>Vascular Invasion</b>	Yes: 19 (20%) No: 36 (39%) Unknown: 37 (40%)
<b>Spontaneous regression</b>	Yes: 3 (3%) No: 56 (60%) Unknown: 33 (35%)
<b>Stage</b>	Localized: 52 (56%) Locally advanced: 20 (22%) Metastatic: 20 (22%)

Comparison of 92 patients with *BRAF* WT (32%) and *BRAF* mut (68%) found a statistically significant association with **age at diagnosis** ( $p = .001$ ; Student's Test) and **ulceration** ( $p = .005$ ; Mann-Whitney's Test) only.



## Conclusions

This is the first report of *BRAF* mutation status using RT-PCR COBAS in Argentina. The frequency of *BRAF* mutation is lower than that reported in medical literature, and only the presence of ulceration and age at diagnosis were significantly different in *BRAF*-mutant and *BRAF* wild-type patients. Future studies with more number of patients will clarify our observation.

### References

- Garnett MJ, et al. *Cancer Cell* 2004;6:313-9
- Wan PTC, et al. *Cell* 2004;116:855-67
- Forbes SA, et al. *Nucleic Acids Res* 2010;38:D652-7

- Pritchard C, et al. *Biochem Soc Trans* 2007;35:1329-33
- Flaherty KT, et al. *Cancer* 2010;116:4902-13.
- Viros A, et al. *PLoS Med* 2008;5:e120.
- Long et al. *JCO* 2011



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