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## Prevalence of BRAF<sup>V600E</sup> Mutation in Iranian Patients with Papillary Thyroid Carcinoma: A Single-Center Study

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**Abstract:** The aim of this study was to investigate the frequency of the BRAF<sup>V600E</sup> mutation in Papillary Thyroid Carcinoma (PTC) in Iranian population through PCR-RFLP. Fifty formalin-fixed paraffin-embedded and 26 frozen thyroid tumors including 28 PTCs, two undifferentiated thyroid carcinoma and 46 benign thyroid tumors were evaluated. The BRAF<sup>V600E</sup> mutation was detected in 20 of 28 (71.4%) PTCs, but failed to distinguish the mutant allele in benign thyroid tumors. The age, sex, extrathyroid extension and lymph node metastases distribution did not, significantly, differ between the patients with and without the BRAF<sup>V600E</sup> mutation. In conclusion, these findings might pave our way towards considering the BRAF<sup>V600E</sup> mutation in PTCs in the regions with high prevalence of this alteration as a molecular marker.

**Key words:** Papillary thyroid carcinoma, mutation, BRAF<sup>V600E</sup>, PCR-RFLP

### INTRODUCTION

Thyroid carcinoma is the most common endocrine malignancy. Recently, Kilfoy *et al.* (2009), reported that the world wide incidence rate of thyroid carcinoma varies between 0.7 to 3.4 and 1.8 to 11.8 per 100,000 populations for men and women, respectively in 19 different populations from 1998 to 2002. The thyroid cancer prevalence in Iran has been reported 1.0 and 3.5 per 100,000 population for male and female, respectively and the most frequent morphologic variant is Papillary Thyroid Carcinoma (PTC) with a frequency of 69.9% (Larijani *et al.*, 2005). The BRAF mutations have been found in a variety of human cancers, most notably in melanoma. T1799A transversion (formerly known as T1796A) in exon 15, which is the most common BRAF mutation, accounts for more than 80% of BRAF mutations (Davies *et al.*, 2002). The T1799A mutation results in constitution of Glutamate (E) for Valine (V) (V600E).

This missense mutation alters the conformation of the activation loop in the BRAF kinase domain and activates its kinase activity by simulating phosphorylation. This activation leads to tumorigenesis through the RAS-RAF- MEK-ERK-MAP kinase pathway

(Davies *et al.*, 2002). Although, the BRAF<sup>V600E</sup> mutation is reported to have association with more aggressive tumor phenotype and a higher risk of recurrence and persistent disease in patients with conventional PTC (Kebebew *et al.*, 2007; Xing *et al.*, 2005; Kim *et al.*, 2006; Nikiforova *et al.*, 2003; Xu *et al.*, 2003; Lee *et al.*, 2007), the reports are controversial (Kim *et al.*, 2005; Liu *et al.*, 2005; Ito *et al.*, 2009). Preoperative detection of PTCs by a specific molecular cancer maker, such as BRAF mutation, would specially be helpful for predicting disease aggressiveness while determining the need for and extent of surgery, as well as, the need for adjuvant therapy and follow-up monitoring (Kebebew *et al.*, 2007). As to our knowledge, there is no published data reporting BRAF<sup>V600E</sup> mutation in Iranian population. The aim of this study was to analyze the prevalence of the BRAF<sup>V600E</sup> mutation in Iranian PTC patients and its applicability in the diagnosis of PTC.

### MATERIALS AND METHODS

**Thyroid samples:** This study was conducted in Department of Medical Genetics, Faculty of Medicine, Tehran University of Medical Sciences, Tehran, Iran from

2007 through 2008. A total of 50 Formalin Fixed Paraffin Embedded (FFPE) samples were obtained from patients who underwent thyroidectomy surgery from 2006 to 2007 in referral Shariati Hospital, Tehran University of Medical Sciences, Tehran, Iran. The tissue samples of total of 50 patients, consisting of 25 malignant tumors and 25 non-malignant samples were reviewed by a pathologist. Also, a total of 26 fresh thyroid tissues were obtained from the patients who underwent surgery in 2008 in referral Shariati Hospital, Tehran University of Medical Sciences, Tehran, Iran. Tissue specimens were put in RNAlater (Ambion, Austin, USA) and frozen immediately at  $-80^{\circ}\text{C}$  until use. Final Histologic classification was obtained finally from pathologic reports. The samples included 5 malignant and 21 non-malignant tumors. This study was approved by the Ethics and Research Committee of Tehran University of Medical Sciences and was conducted in accordance with the Declaration of Helsinki principles.

#### Cell line samples, Laser Capture Microdissection (LCM)

**and DNA extraction:** Previously extracted DNA of cell lines, ARO and 8305C (heterozygous for BRAF<sup>V600E</sup> mutation), 8505C (homozygous for BRAF<sup>V600E</sup> mutation) and finally TPC-1 (wild type BRAF gene), were used as positive (heterozygous and homozygous mutation) and negative controls, respectively. The 6  $\mu\text{M}$  FFPE tissue sections were deparafinized and stained with Methyl Green (Sigma-Aldrich, St. Louis, MO) and then manually dissected or using laser capture microdissection system (Leica AS LMD, Wetzlar, Germany). The DNA from dissected FFPE samples was extracted by using QIAamp DNA Micro Kit (QIAGEN, Hilden, Germany) according to the manufacturer's instructions. DNA from frozen tissues was extracted by conventional Phenol-Chloroform method. Quality and quantity of the extracted DNAs were determined by spectrophotometry (NanoDrop ND-1000, Wilmington, Delaware USA).

**PCR-RFLP:** We amplified BRAF exon 15 by PCR using oligonucleotide primer pairs: 5-TCA TGA AGA CCT CAC AGT AAA AAT-3 (forward) and 5-TGG ATC CAG ACA ACT GTT CAA-3 (reverse), as described previously by Takahashi *et al.* (2007). The PCR was carried out with the following conditions: initial denaturation at  $95^{\circ}\text{C}$  for 5 min, followed by 40 cycles of denaturation at  $95^{\circ}\text{C}$  for 30 sec, annealing at  $56^{\circ}\text{C}$  for 30 sec and elongation at  $72^{\circ}\text{C}$  for 30 sec and a final extension at  $72^{\circ}\text{C}$  for 5 min. The PCR products were visualized on 8% polyacrylamid gels with Ethidium Bromide staining.

Then, the amplification products were digested by TspRI (New England Biolabs, Ipswich, MA) restriction

endonuclease that cuts normal allele (ACA GTG AAA) but not the T1799A mutant sequence (ACA GAG AAA). The wild type BRAF fragment produced two 47 bp and 52 bp bands, while the mutant BRAF fragment was not digested and produced a 98 bp band. For confirmation of the results, we sequenced the PCR product in the 5 cell lines previously mentioned and some positive and negative mutant samples.

**Statistical analysis:** All the statistical analyses were performed by the SPSS software. Fisher exact test was used for association of BRAF<sup>V600E</sup> mutation with age and gender. The p-value of less than 0.05 considered to be significant. The True-Positive (TP), True-Negative (TN), False-Positive (FP) and False-Negative (FN) results were defined. Diagnosis sensitivity, specificity, Positive Predictive Value (PPV), Negative Predictive Value (NPV) and accuracy were calculated by using the following formulas:

$$\text{Sensitivity}(\%) = \frac{\text{TP}}{\text{TP} + \text{FN}} \times 100$$

$$\text{Specificity}(\%) = \frac{\text{TN}}{\text{TN} + \text{FP}} \times 100$$

$$\text{PPV}(\%) = \frac{\text{TP}}{\text{TP} + \text{FP}} \times 100$$

$$\text{NPV}(\%) = \frac{\text{TN}}{\text{TN} + \text{FN}} \times 100$$

$$\text{Accuracy}(\%) = \frac{\text{TP} + \text{TN}}{\text{TP} + \text{TN} + \text{FP} + \text{FN}} \times 100$$

## RESULTS

Overall, the prevalence of BRAF<sup>V600E</sup> mutation was 20 out of 30 (66.7%) in malignant thyroid carcinoma and 71.4% in PTCs. The prevalence of BRAF<sup>V600E</sup> mutation regarding to histologic classification of tumours was 72, 100 and 0% in Classic-Variant PTCs (CV-PTC), Tall-Cell variant PTCs (TC-PTC) and Follicular Variant PTC (FV-PTC), respectively (Table 1). It is worthy to note that in one TC-PTC patient the BRAF<sup>V600E</sup> mutation was present as homozygous. No BRAF<sup>V600E</sup> mutation was detected in the normal thyroid tissues surrounding the malignant tissue.

The age, sex, extrathyroid extension and lymph node metastases distribution of the patients with the malignant thyroid samples regarding to BRAF<sup>V600E</sup> mutation status are shown in Table 1. All the 46 benign specimens including 24 goiters, 15 follicular adenomas,

**Table 1: BRAF<sup>V600E</sup> mutation analysis of thyroid histology in 30 malignant and 46 benign thyroid tumors**

Histology type	No. of cases	BRAF <sup>V600E</sup> positive No. of cases (%)	p-value
Papillary thyroid carcinoma	28	20 (71.4)	--
Classic variant	25	18 (72)	
Tall-cell variant	2	2 (100)	
Follicular variant	1	-	
Undifferentiated thyroid carcinoma	2	-	---
Multinodular goiter	24	-	
Hashimoto's thyroiditis	7	-	
Follicular adenoma	15	-	
<b>Gender distribution among 28 PTCs patients</b>			
Male	6	4 (66.7)	NS
Female	22	16 (72.2)	
<b>Age distribution among 28 PTCs patients</b>			
≤40 years	13	9 (69.2)	NS
>40 years	15	11 (73.3)	
<b>Extrathyroidal extension among 25 PTCs</b>			
Present	19	11 (57.9)	NS
Absent	6	6 (100)	
<b>Lymph node metastases among 25 PTCs</b>			
Present	14	9 (64)	NS
Absent	11	8 (72.2)	

NS: Non significant, p>0.05

**Table 2: Correlation between BRAF<sup>V600E</sup> mutation presence and malignancy detection in study results**

Patient groups (n = 76)	Values
TP	20.00
TN	46.00
FP	0.00
FN	10.00
Sn (%)	66.67
Sp (%)	100.00
PPV (%)	100.00
NPV (%)	82.14
Accuracy (%)	86.84

TP: True positive, TN: True negative, FP: False positive, FN: False negative, Sn: Sensitivity, Sp: Specificity, PPV: Positive predicted value, NPV: Negative predicted value

7 Hashimoto's thyroiditis, were BRAF<sup>V600E</sup> mutation negative. Diagnostic sensitivity, specificity, PPV, NPV and accuracy were 66.67, 100, 100, 82.18 and 86.84%, respectively (Table 2).

## DISCUSSION

This study showed BRAF<sup>V600E</sup> mutation in 71.4% of PTC samples and it seems more common in TC-PTC (100%) comparing to CV-PTC (72%). Also there was no BRAF<sup>V600E</sup> mutation in benign thyroid lesions and in normal cells surrounding malignant tissues and this finding is consistent with other reports (Nikiforova *et al.*, 2003; Xu *et al.*, 2003; Xing *et al.*, 2004).

The PTCs are usually curable with standard surgical and adjuvant radioiodine treatment but unfortunately lymph node metastases are found in 30 to 65% of cases at the time of initial diagnosis and 15% of tumors with lymph node metastases also display very aggressive behavior, characterized by local invasion, distant metastasis, treatment resistance and increased mortality (Mazzaferri

and Kloos, 2001). The use of molecular markers for diagnosis and prognosis of thyroid cancers has been explored extensively. The V600E mutation in BRAF kinase appears to be an attractive molecular marker for thyroid cancer diagnosis and prognosis as it has been found to be the most common genetic events and specific for PTCs (Rowe *et al.*, 2006). The aim of present study was to determine the prevalence of the BRAF<sup>V600E</sup> mutation in an Iranian population and its applicability for PTC detection.

This study showed that the BRAF<sup>V600E</sup> mutation was much more common in PTC patients of this study compared to western countries. As meta-analysis by Lee *et al.* (2007) most studies reported that BRAF<sup>V600E</sup> mutation is common in PTCs but the reported frequencies vary from 30 to 83% (mean 49%). Some investigators have suggested that diagnostic technical methods and geographic factors may account for the differences in reported prevalences for BRAF<sup>V600E</sup> mutation in PTCs (Kebebew *et al.*, 2007).

Guan *et al.* (2009) recently reported that there is an association between BRAF<sup>V600E</sup> mutation and Iodine intake. According to their report BRAF<sup>V600E</sup> mutation was found in 69% of PTCs in high iodine content in natural drinking water but it was found only in 53% of PTCs in normal Iodine content in natural drinking water in China (Guan *et al.*, 2009). Present results for BRAF<sup>V600E</sup> mutation prevalence in PTCs is more similar to those published for Asian (Korean and Chinese) populations (Guan *et al.*, 2009; Kim *et al.*, 2004). The higher prevalence of BRAF<sup>V600E</sup> mutation in high Iodine intake in China is in agreement with our results. In agreement to this study reports from Korea, a country with very high iodine intake, indicated that BRAF<sup>V600E</sup> mutation is highly prevalent (52-83%) in PTC (Kim *et al.*, 2004, 2005;

Chung *et al.*, 2006). In contrast, the prevalence of BRAF<sup>V600E</sup> mutation in western countries, Such as Spain and Italy showed low prevalence (37-42%) (Lee *et al.*, 2007). According to the results we can propose that the high prevalence of BRAF<sup>V600E</sup> mutation may be due to high Iodine intake in Iran.

The higher frequency of BRAF<sup>V600E</sup> mutation in the present study might be due to the probable increase in iodine intake after starting a mass programme of iodine supplementation implemented since 1983 in Iran (Larijani *et al.*, 2005). However, this proposal requires definition by further studies. Although, we found that BRAF<sup>V600E</sup> mutation occurred slightly more in females, this was not statistically significant ( $p>0.05$ ). The BRAF<sup>V600E</sup> mutation was observed in 73.3% of PTC patients older than 40 years old and in 69.2% of PTC patients younger than 40 years old. BRAF<sup>V600E</sup> mutation prevalence was slightly more common in older patients than in younger patients, but this was not also a significant difference ( $p>0.05$ ). Present results is in agreement with the results of a meta-analysis study on 1168 PTC patients that showed there were not significant association between BRAF<sup>V600E</sup> mutation and age or gender (Lee *et al.*, 2007).

The results of present research showed that there is not significant association between BRAF<sup>V600E</sup> and extrathyroidal extension or lymph node metastases that is in agreement with some previous reports (Lee *et al.*, 2007; Ito *et al.*, 2009).

In this study, we confirmed that BRAF<sup>V600E</sup> mutation was specific for PTC (100%), because no benign thyroid samples were found to harbor BRAF<sup>V600E</sup> mutation and all the samples that were positive for BRAF<sup>V600E</sup> mutation were malignant. This result is consistent with previous reports (Rowe *et al.*, 2006; Xing, 2005). In the present study, the diagnostic sensitivity and specificity were 66.67 and 100%, respectively. Chung *et al.* (2006) reported the sensitivity of 83.0% and a specificity of 96.0% for direct DNA sequencing method and the sensitivity of 79.6% and a specificity of 80.0% for PCR-RFLP method for detection of PTC in Fine Needle Aspiration Biopsy (FNAB) samples.

High specificity in our study indicates the advantage of BRAF<sup>V600E</sup> mutation for PTCs detection. However, the method based on PCR-RFLP is enough sensitive to detect BRAF<sup>V600E</sup> mutation even in the samples of FFPE tissue and when the DNA yield was low (Kumagai *et al.*, 2007). The lower sensitivity might be explained by the fact that diagnostic sensitivity of BRAF<sup>V600E</sup> mutation strictly depends upon its prevalence in PTC patients.

In conclusion, these findings might pave our way towards considering the BRAF<sup>V600E</sup> mutation in PTCs in the regions with high prevalence of this alteration as a molecular marker.

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