

Original Article

Indicators of multifocality in papillary thyroid carcinoma concurrent with Hashimoto's thyroiditis

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Abstract: Currently, no definitive diagnostic tool is available to distinguish unifocal and multifocal papillary thyroid carcinoma (PTC). This study aims to identify potential diagnostic markers of multifocal PTC. In 471 Hashimoto's thyroiditis (HT) patients, the significant difference was revealed in anti-thyroid peroxidase antibody (TPOAb) concentration, the cytokeratin-19 (CK-19) expression, the occurrence of the B-Raf proto-oncogene serine/threonine kinase (BRAF) mutations and the rearrangement in transformation (RET)/PTC. The patients' samples were assayed for the expression of CK-19, cyclooxygenase-2 (COX-2), galectin-3, and the protein human bone marrow endothelial cell marker-1 (HBME-1) using immunohistochemistry. The BRAF gene mutation was detected using a sequencer. Differences were examined using the Kruskal-Wallis test and the Chi-squared and Fisher's exact tests. The results showed that the elevated CK-19 expression, and the presence of BRAF mutations and RET/PTC rearrangements were indicators of multifocal PTC in HT, suggesting the need for total bilateral thyroidectomy. Among HT patients with TPOAb > 1300 IU/ML, the occurrence of central lymph node metastasis is significantly higher in multi-focal PTC than single-focal PTC. Therefore, these markers may prove useful for discerning between uni- and multifocal PTC, thereby preventing unnecessary surgery in the treatment of unifocal PTC and promoting sufficient treatment of multifocal PTC.

Keywords: Hashimoto's thyroiditis (HT), papillary thyroid carcinoma (PTC), biomarker, thyroidectomy, anti-thyroid peroxidase antibody (TPOAb)

Introduction

Hashimoto's thyroiditis (HT) is an autoimmune condition that leads to the destruction of thyroid cells and is the leading cause of hypothyroidism worldwide. A growing body of evidence indicates that HT patients are at increased risk of developing papillary thyroid carcinoma (PTC) [1]. Although the extent of this relationship is still a matter of debate [2, 3], two large-cohort meta-analyses of recent studies both conclude that HT is clearly associated with PTC [4, 5]. Zhang et al. found that HT and PTC are significantly associated and concluded that long-term exposure to the elevated thyroid stimulating hormone (TSH) levels typical of HT likely predisposes HT patients to PTC [6].

PTC is most often unifocal, but multifocal PTC can arise from the spread of a single primary

tumor or from multiple simultaneous primary tumors. Patients with multifocal PTC are at increased risk for lymph node metastasis, distant metastasis, local recurrence after initial treatment, and regional recurrence [7]. The rate of PTC multifocality is higher in HT patients [8]. The frequency of PTC metastasis to visceral lymph nodes is 4-fold higher in patients with HT than in those without [3]. Wang et al. found that bilateral PTC has greater invasiveness, faster progression, and a shorter 10-year disease-free survival compared to unilateral multifocal PTC. The increased incidence of lymph node metastasis in patients with bilateral PTC contributes to its poor prognosis [9].

The preferred treatment for PTC in HT is surgical resection followed by other treatment modalities. However, opinions vary as to the scope of surgical resection needed. Clearly, multifocal

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PTC in HT should be treated by bilateral thyroidectomy. After unilateral lobectomy, TC recurs in 2-9% of the contralateral lobes. Unilateral resection may leave behind remnant cancer, resulting in contralateral recurrence or even metastases after surgery. Thus, more extensive thyroidectomy is recommended [10]. However, patients with non-total resection have a lower incidence of postoperative complications such as hypothyroidism and hypocalcemia than do those who undergo total resection [11]. In addition, the prognosis of HT patients does not differ significantly between treatment with thyroidectomy and total resection. According to the 2015ATA thyroid cancer diagnosis and treatment guidelines [12], unilateral lobectomy is recommended for low-risk, unifocal PTC.

Unfortunately, no diagnostic tool is available for determining whether HT-associated PTC is uni- or multifocal. Intraoperative frozen section analysis examines only obvious nodules, often missing small cancerous lesions. Unilateral resection only examines the affected side and cannot detect the presence of cancerous lesions in the contralateral side. Pathological examination can only confirm a diagnosis of multifocal cancer after total bilateral resection. Thus, patients with unifocal cancer may be over-treated, exposing them to unnecessary risk of postoperative complications. Therefore, the identification of potential early indicators of multifocal cancer in HT is needed for guiding appropriate clinical treatment plans.

Current research in this area is focused on identifying specific biochemical and genetic markers to distinguish multifocal PTC. Azizi et al. found that elevated serum anti-thyroid peroxidase antibody (TPOAb), the key etiological autoantibody in HT, and TSH ≥ 1 μ U/mL are independent predictors of thyroid cancer in patients with thyroid nodules [13]. The expression of human bone marrow endothelial cell marker-1 (HBME-1), a membrane protein present in the microvilli of mesothelioma and follicular thyroid tumor cells, is significantly higher in PTC patients with malignant than with benign thyroid lesions [14]. Similarly, the expression of Galectin-3, a β -galactoside-binding protein involved in cell migration, adhesion, and apoptosis, is significantly higher in malignant than in benign thyroid lesions [14, 15]. CK-19, a keratin family member responsible for the structural

integrity of epithelial cells, is highly expressed in PTC follicular variants but not in benign lesions such as thyroid adenomas [16]. The expression of cyclooxygenase-2 (COX-2), which promotes cell proliferation and inhibits apoptosis by catalyzing prostaglandin synthesis, is elevated in many precancerous lesions and malignant tumors [17, 18].

Genetic markers under investigation for discerning between metastatic and benign cancers include BRAF gene mutations and rearranged in transformation (RET)/PTC oncogene rearrangements. Both of these genetic alterations are common in thyroid cancer. Mutations in BRAF, a serine/threonine kinase, increase its kinase activity, resulting in the activation of ERK. RET/PTC oncogene rearrangements involve the fusion of the tyrosine kinase domain of RET to the 5' region of unrelated genes, creating dominantly transforming oncogenes [3].

This retrospective study aims to identify markers that distinguish between unifocal and multifocal PTC in HT patients. Blood chemistry parameters, thyroid function assays, and TP-OAb levels were compared between HT patients with unifocal PTC, multifocal PTC, and no PTC. These groups were also compared with respect to immunohistochemical analysis of tissue samples for CK-19, COX-2, Galectin-3, and HBME-1 expression and the presence of BRAF mutations and RET/PTC oncogene rearrangements.

Materials and methods

HT patients

This study is a retrospective analysis of 471 patients with PTC and HT who underwent thyroid surgery from January 2013 to January 2015, including 183 patients in group A (HT with multifocal PTC), 187 in group B (HT with single PTC), and 101 in group C (HT alone). All patients were diagnosed with thyroid nodules due to suspected malignancy or large nodules. Inclusion criteria included thyroid surgery indications and postoperative pathology confirmed as Hashimoto's thyroiditis with papillary thyroid carcinoma or Hashimoto's thyroiditis. Exclusion criteria included hyperthyroidism, history of subacute thyroiditis, history of tuberculosis or other malignant tumors, low immune function, and a history of other major medical conditions.

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Patients who underwent chemotherapy, radiotherapy, or immunotherapy before surgery were excluded. This study was approved by the ethics committee of the First Affiliated Hospital, College of Medicine, Zhejiang University (No. 2016-119).

Clinical data analysis

Blood was collected from all patients before surgery. The following serum indicators were assayed: preoperative T3, T4, TSH, thyroglobulin (TG), calcitonin, and parathyroid hormone; tumor markers, including carcinoembryonic antigen (CEA), carbohydrate antigen CA199, CA-125, alpha-fetoprotein (AFP), ferritin, blood sugar, and blood lipids. The preoperative TPOAb concentration was also determined. Routine pathology results were compared between groups to determine their correlations with unifocal cancer, multifocal cancer, and positive central lymph node metastasis.

Immunohistochemistry

The formalin-fixed, paraffin-embedded samples of all patients were assayed for the expression of cytokeratin-19 (CK-19), cyclooxygenase-2 (COX-2), Galectin-3, and HBME-1 using immunohistochemistry. Paraffin sections were prepared by dewaxing hydration, endogenous peroxidase blocking, non-specific protein blocking, incubation with primary, secondary, and streptavidin-HRP antibodies, DAB color development, hematoxylin counterstaining, and dehydration. The slides were photographed using a microscopic imaging system and carried out quantitative analysis of average optical density using internationally accepted Image Pro Plus software (Media Cybernetics, Rockville, MD, USA). The primary antibodies, all purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA) were as follows: (1) rabbit anti-CK-19 antibody, (2) rabbit anti-COX-2 antibody, (3) rabbit anti-Galectin-3 antibody, and (4) rabbit anti-HBME-1 antibody.

Detection of the BRAF mutation

Genomic DNA was extracted from resected specimens from all patients. PCR was performed to amplify the target gene fragments, and the BRAF gene mutation was detected using a sequencer. PCR primers were designed based on the BRAF genomic DNA sequence

using Primer5 online software, and PCR amplification was performed to amplify exon 15 of the BRAF gene. The primer sequences were as follows:

Forward primer: 5'-TCATAATGCTTGCTCTGAT-AGGA-3'; Reverse primer: 5'-GGCCAAAATTT-AATCAGTGGA-3'. The length of the amplified fragment was 224 bp, and the primer was synthesized by Shenggong Bioengineering Co., Ltd (Shanghai, China). The PCR reaction conditions were as follows: Pre-change at 94°C for 5 min, denaturation at 94°C for 30 s, annealing at 55°C for 30 s, extension at 72°C for 30 s, 35 cycles in total, extension at 72°C for 5 min, ending at 4°C. The PCR products were sequencing using the Sanger Chain Termination method.

RNA extraction

The tissue was placed in a 1-mL Trizol (Invitrogen, Carlsbad, CA, USA) homogenate tube, homogenized in a homogenizer for 20 sec, and incubated for 5 min. After centrifugation, the supernatants were added to 200 µL chloroform, shake well, and stand at room temperature for 2 min. After centrifugation, the supernatants were added to 600 µL isopropanol, mixed well, and left at room temperature for 15 min. After centrifugation, the pellets were washed with 1 ml 75% absolute ethanol and dried at room temperature for 10 min. To the samples was added 40 µL DEPC water to dissolve the RNA, and the solutions were stored at -80°C for later use.

Detection of RET/PTC oncogene rearrangements

Specimens from all patients were selected and confirmed by pathological diagnosis. After total RNA extraction and cDNA synthesis, PCR amplification was carried out using the nested PCR method. The primer sequences were as follows: RET/PTC1 first round: F, 5'-GTCATCTC-GCCGTTCC-3'; R, 5'-CTTTCAGCATCTTCACGG-3'; RET/PTC1 second round: F, 5'-GCTGGAGACCTA-CAAACCTGA-3'; R, 5'-CGTTGCCTTGACCACTTTTC-3'; RET/PTC3 first round: F, 5'-AAGCAAACCTG-CCAGTGG-3'; R, 5'-CGTTGCCTTGACCACTTTTC-3'; RET/PTC3 second round: F, 5'-CAAGCT-CCTTACATACC-3'; R, 5'-CCTTCTCCTAGAGTTTTT-CC-3'; 18S: F, 5'-CGACGACCCATTCGAACGTCT-3'; R, 5'-CTCTCCGGAATCGAACCCCTGA-3'. PCR

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reaction conditions were as follows: pre-change for 5 min at 94°C, denaturation at 94°C for 30 s, annealing at 55°C for 30 s, extension at 72°C for 30 s, 35 cycles in total, extending at 72°C for 5 min, ending at 4°C. The product obtained by PCR amplification was subjected to gelatinization purification, and the base sequence was determined using a DNA sequencer. The sequence was compared to that of the genomic DNA to determine the presence or absence of the mutation and the mutation position.

Statistical analysis

Continuous variables were presented as the median with the interquartile range (IQR), and categorical variables were reported as the number with a percentage. Differences between groups were examined using the Kruskal-Wallis test for continuous variables and the Chi-squared and Fisher's exact tests for categorical variables. Dunn's test was implemented if a significant difference was obtained by the Kruskal-Wallis test. Logistic regression analysis was performed to identify predictors of multifocal PTC in HT, and all variables were evaluated by multivariate analysis with stepwise selection. Receiver operating characteristic (ROC) curves of CK-19 were generated to determine the optimal cut-off point and to distinguish between HT patients with multifocal and unifocal PTC. All *p*-values were two-sided, and *P* < 0.05 was considered statistically significant. All statistical analyses were performed using the statistical software package SAS version 9.4 (SAS Institute Inc., Cary, NC, USA).

Results

Demographic and clinical findings

Of the 471 HT patients enrolled in the study, 183 patients had HT with multifocal PTC (38.9%), 187 patients had HT with single PTC (39.7%), and 101 patients had HT alone (21.4%). The median patient age in these groups was 46, 41, and 46 years, respectively. More than 80% of the patients in all groups were female (86.9%, 87.2%, and 87.1%, respectively). Calcitonin and parathyroid hormone levels were normal in all patients. Only two patients with single PTC had tumor markers (CA-

125 and CA199). HT patients with multifocal (100.0%) or unifocal PTC (98.9%) underwent total thyroidectomy. However, two patients with single PTC underwent a thyroid lobectomy, and all of the patients with HT alone underwent subtotal thyroidectomy (**Table 1**).

Groups differed significantly with respect to the concentration of TPOAb, positive central lymph node metastasis, TSH level, number of tumors, and size of the largest tumor (all *P* < 0.001). More HT patients with multifocal PTC had TPOAb > 1300 IU/mL (71.0%) and were positive for central cervical lymph node metastasis (67.8%) than were patients in other groups (**Table 1**).

In addition, the status of central lymph node metastasis differed with the level of TPOAb between HT patients with multifocal and unifocal PTC (*P* < 0.001). Among HT patients with multifocal PTC and TPOAb concentration > 1300 IU/mL, most were positive for central lymph node metastasis (43.7%); among unifocal patients with TPOAb < 1300 IU/mL, most were negative for central lymph node metastasis (45.5%) (Supplementary Table 1).

Immunohistochemistry results

The expression of CK-19, COX-2, Galectin-3, and HBME-1 differed significantly between tissue samples from different groups (all *P*s ≤ 0.045) (**Figure 2** and **Table 2**). However, Dunn's multiple comparisons test showed that only the median optical density of CK-19 differed significantly between PTC patients with multifocal and unifocal PTC (0.011 vs. 0.006, respectively) (**Table 2**).

We determined the optimal cutoff point in the optical density of CK-19 to determine the relationship between CK-19 expression and multifocal vs. unifocal PTC. ROC analysis indicated that CK-19 optical density had acceptable discrimination for distinguishing between multifocal and unifocal PTC in HT patients (AUC, 0.749; median optimal cutoff point, 0.008) (**Figure 1**).

Genetic findings

The occurrence of BRAF and RET/PTC in HT patients differed significantly between those with multifocal and unifocal PTC (all *P* < 0.001).

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Table 1. Comparison of demographic and clinical data between groups

	HT with multifocal PTC N = 183 n (%)	HT with single PTC N = 187 n (%)	HT alone N = 101 n (%)	P
Demographic				
Age	46.0 (40.0, 51.0)	41.0 (34.0, 54.0)	46.0 (32.0, 51.0)	0.989
Female	159 (86.9)	163 (87.2)	88 (87.1)	0.996
Clinical indicators				
TG (ng/mL)	31.0 (21.0, 44.0)	29.0 (22.0, 44.0)	27.0 (21.0, 38.0)	0.157
Calcitonin = Normal	183 (100.0)	187 (100.0)	101 (100.0)	NA
Tumor markers (Abnormal)				
AFP	0 (0.0)	0 (0.0)	0 (0.0)	NA
CA125	0 (0.0)	1 (0.6)	0 (0.0)	
CA199	0 (0.0)	1 (0.6)	0 (0.0)	
CEA	0 (0.0)	0 (0.0)	0 (0.0)	
Ferritin	0 (0.0)	0 (0.0)	0 (0.0)	
PTH = Normal	183 (100.0)	187 (100.0)	101 (100.0)	NA
Blood sugar = Abnormal	4 (2.2)	4 (2.1)	4 (4.0)	0.651
Blood lipid = Abnormal	11 (6.0)	12 (6.4)	8 (7.9)	0.819
TPOAb > 1300 (IU/mL)	130 (71.0)	22 (11.8)	48 (47.5)	< 0.001
Central compartment of cervical lymph node metastasis = Positive	124 (67.8)	97 (51.9)	0 (0.0)	0.002*
TSH (μ IU/mL)	2.4 (1.6, 3.4) ^f	2.3 (1.6, 2.7) ^c	3.4 (2.6, 4.2) ^{a,b}	< 0.001
Thyroid surgery				
LOBE	0 (0.0)	2 (1.1)	0 (0.0)	
Subtotal	0 (0.0)	0 (0.0)	101 (100.0)	
Total	183 (100.0)	185 (98.9)	0 (0.0)	
Number of tumors	2.0 (2.0, 2.0)	1.0 (1.0, 1.0)	1.0 (1.0, 1.0)	< 0.001
Size of the largest tumor (cm)	0.8 (0.6, 1.2)	0.8 (0.5, 1.2)	3.6 (3.3, 3.9)	< 0.001

HT, Hashimoto's thyroiditis; PTC, papillary thyroid carcinoma; TG, thyroglobulin; AFP, alpha-fetoprotein; CEA, carcinoembryonic antigen; PTH, parathyroid hormone; TPOAb, thyroid peroxidase antibodies; TSH, thyroid-stimulating hormone; LOBE, thyroid lobectomy; Subtotal, subtotal/near-total thyroidectomy; Total, total thyroidectomy. Continuous variables were presented as median and IQR; categorical variables were presented as number and percentage. ^fHT with multifocal PTC vs. HT with single PTC. *Significant difference vs. HT with multifocal PTC, P < 0.05. ^bSignificant difference vs. single PTC, P < 0.05. ^cSignificant different vs. HT alone, P < 0.05.

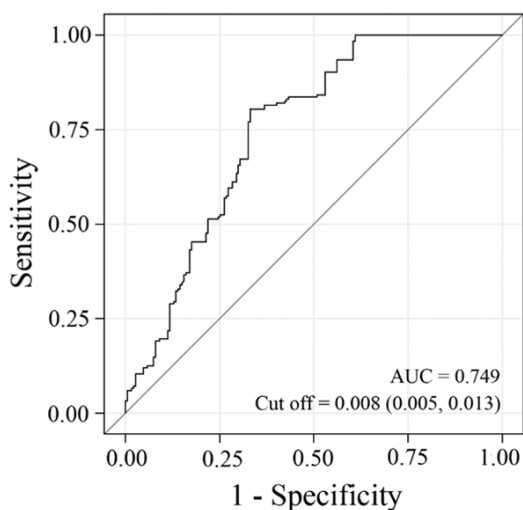


Figure 1. ROC curves to detect the optimal cutoff of CK-19 expression for multifocal PTC in HT.

A greater fraction of HT patients with multifocal PTC had the BRAF mutation (41.5%) or positivity

for RET/PTC (56.8%) than did those with unifocal PTC (Table 3).

Risk factors for multifocal PTC in HT patients

The results of comparisons between HT with multifocal PTC patients and HT single PTC patients in associated factors are shown in Supplementary Table 2. Univariate regression analysis revealed significant differences between HT patients with multifocal and unifocal PTC with respect to TPOAb concentration, central lymph node metastasis, tissue CK-19 optical density, and the presence of BRAF and RET/PTC. After stepwise selection, the concentration of TPOAb and presence of BRAF and RET/PTC were further investigated using multivariate regression analysis. The results show that elevated TPOAb (aOR, 83.11; 95% CI, 33.96-203.44; P < 0.001), BRAF mutation (aOR, 10.92; 95% CI, 4.64-25.72; P < 0.001), and positivity for RET/PTC (aOR, 13.36; 95% CI,

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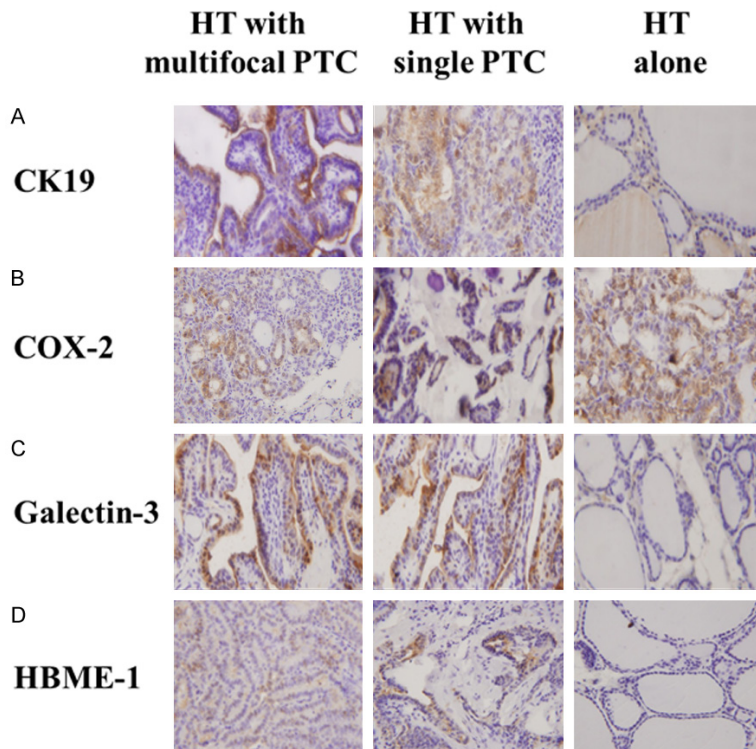


Figure 2. Immunohistochemistry results of CK-19, COX-2, Galectin-3 and HBME-1 in each group. The protein expression of (A) CK-19, (B) COX-2, (C) Galectin-3, and (D) HBME-1 was detected in HT patients with multifocal PTC, with single PTC, and without PTC. Fields at 40 \times magnification were shown, and 1 representative from triplicates was shown for each staining.

5.81-30.73; $P < 0.001$) are associated with multifocal PTC (Supplementary Table 2).

Discussion

This retrospective study of 471 HT patients aims to identify potential diagnostic markers to discern between uni- and multifocal PTC. We observed a significant difference in TPOAb concentration, CK-19 expression, and the occurrence of BRAF mutations and RET/PTC rearrangements between patients with uni- and multifocal PTC. Our results indicate that TPOAb > 1300 IU/mL, elevated TSH only together with TPOAb > 1300 IU/mL, elevated CK-19 expression, and the presence of BRAF mutations and RET/PTC rearrangements are indicators of multifocal PTC in HT. Among HT patients with TPOAb > 1300 IU/MI, the occurrence of central lymph node metastasis is significantly higher in those with multi-focal than single-focus PTC. These markers may prove useful for discerning between uni- and multifocal PTC, thereby preventing unnecessarily extensive surgery in the treat-

ment of unifocal PTC and promoting sufficient treatment of multifocal PTC.

Thyroid peroxidase, a major component of thyroid microsomes, is a key enzyme in the synthesis and secretion of thyroxine. Autoimmune attack on this enzyme by TPOAb, the etiological agent of HT, causes thyroid cell destruction, inhibiting thyroxine production and resulting in a rise in thyroid stimulating hormone (TSH). The chronic inflammation induced by TPOAb is considered to be a risk factor for thyroid cancer [19, 20]. The relationship between elevated TSH and TPOAb is closely related to the development of multifocal cancer. Our results indicate that TPOAb > 1300 IU/mL is a risk factor for HT with multifocal PTC. This finding supports the observation of Dong et al. that a high level of TPOAb (> 1300 IU/mL) is a definitive indicator of multifocal

PTC in HT patients [21]. We also observed that elevated TSH accompanied by TPOAb > 1300 IU/mL is a risk factor for multifocal PTC in HT. In such cases, we recommend bilateral thyroidectomy. In contrast, elevated TSH with mild to moderate elevation of TPOAb (< 1300 IU/mL) did not correlate with multifocal PTC in HT, demonstrating that elevated TSH cannot be used as a sole indicator of multifocal PTC.

Studies suggest that central lymph node dissection should be standard treatment for thyroid cancer patients with central lymph node metastasis [22, 23]. Our results showed that positivity for central lymph node metastasis in HT was significantly higher in multi-focal PTC with TPOAb > 1300 IU/MI than in unifocal PTC. Therefore, we recommend central lymph node dissection for HT patients with high-level expression of TPOAb.

We investigated whether CK-19, COX-2, Galectin-3, and HBME-1 expression can distinguish between uni- and multifocal PTC in HT. These proteins were chosen because of their

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Table 2. Comparison of tissue immunohistochemical data between groups

	HT with multifocal PTC N = 183 n (%)	HT with single PTC N = 187 n (%)	HT alone N = 101 n (%)	P
Immunohistochemistry				
CK19	0.011 (0.007, 0.013) ^{b,c}	0.006 (0.002, 0.011) ^{a,c}	0.003 (0.001, 0.004) ^{a,b}	< 0.001
COX-2	0.005 (0.003, 0.009) ^c	0.004 (0.002, 0.008) ^c	0.002 (0.001, 0.004) ^{a,b}	< 0.001
Galectin-3	0.009 (0.005, 0.021) ^c	0.014 (0.009, 0.021) ^c	0.003 (0.002, 0.004) ^{a,b}	< 0.001
HBME-1	0.006 (0.004, 0.008) ^c	0.006 (0.004, 0.007)	0.006 (0.003, 0.006) ^a	0.045

HT, Hashimoto's thyroiditis; PTC, papillary thyroid carcinoma; CK19, cytokeratin 19; COX-2, cyclooxygenase 2; HBME-1, mouse anti mesothelioma. Continuous variables are presented as the median and IQR. ^aSignificant difference vs. HT with multifocal PTC, P < 0.05. ^bSignificant difference vs. HT with single PTC, P < 0.05. ^cSignificant different vs. patients with HT alone, P < 0.05.

Table 3. Comparison of genetic alterations between groups

	HT with multifocal PTC N = 183 n (%)	HT with unifocal PTC N = 187 n (%)	HT alone N = 10 n (%)	P
Gene detection				
BRAF Mutation	76 (41.5)	18 (9.6)	0 (0.0)	< 0.001*
RET/PTC-positive	104 (56.8)	39 (20.9)	0 (0.0)	< 0.001*

HT, Hashimoto's thyroiditis; PTC, papillary thyroid carcinoma; BRAF, B-type Raf kinase; RET/PTC, rearranged in transformation/papillary thyroid carcinoma. Categorical variables are presented as count and percentage. *Multifocal PTC vs. unifocal PTC.

known patterns of expression and potential specificity as indicated by previous studies. CK-19 and HBME-1 were the two most commonly used immunological markers for PTC diagnosis. These markers could be a useful complement to conventional histological diagnostic methods [14]. Cochand-Priollet et al. found that diagnostic CK-19 and HBME-1 antibody staining had a lower rate of false positive and false negative results in PTC diagnosis compared to morphological methods, thereby improving the accuracy of diagnosis and reducing unnecessary surgical treatment [24]. Liu et al. found that the identification of up-regulated CK-19 (or HBME-1) expression together with down-regulated TPO expression improved the specificity of PTC diagnosis [25]. CK-19 is present in tumors of normal epithelium and epithelial origin and is used to diagnose adenocarcinoma. CK-19 also is expressed in thyroid cancer [18], with high expression in PTC follicular variants but no expression in benign lesions such as thyroid adenomas [16]. Baloch et al. found that analysis of CK-19 expression was very helpful for PTC diagnosis [26]. Guyetant et al. [27] reported that CK-19 was expressed in all PTCs, with 100% sensitivity and 82.15% specificity, and was useful for detecting micrometastases in thyroid tissue. Compared with

these findings, however, we observed significantly higher expression in multifocal than in unifocal PTC for CK-19 but not HBME-1. In addition, ROC analysis indicated that CK-19 optical density had acceptable discrimination for distinguishing between multifocal and unifocal PTC in HT patients. Thus, in intraoperative frozen section analysis of CK-19, a mean optical density > 0.008 indicates multifocal cancer, strongly suggesting that the best treatment choice is total thyroidectomy.

Galectin-3 is a β -galactoside-binding protein involved in cell growth, adhesion, inflammatory responses, immune regulation, and apoptosis. Studies have found that Galectin-3 is expressed in breast cancer, gastric cancer, colon cancer, and other tumors [28], but its expression in thyroid cancer is still uncertain. Galectin-3 is thought to be expressed in malignant tumors, particularly PTC, but not in benign lesions and normal thyroid tissue. Studies have shown that Galectin-3 is valuable for the differential diagnosis of PTC and benign thyroid lesions [29]. Galectin-3 expression in benign lesions may suggest that the lesion has the potential to differentiate into cancer [30]. We observed that Galectin-3 is positively expressed in HT with PTC and weakly positively expressed in HT al-

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one. No significant difference was observed between uni- and multifocal cancer in HT. Therefore, we conclude that Galectin-3 cannot be used as an indicator to distinguish between uni- and multifocal PTC and can be used only as an auxiliary indicator for judging whether PTC is present in HT.

COX-2 catalyzes the conversion of arachidonic acid to prostaglandin, which promotes cell proliferation and inhibits apoptosis. COX-2 expression is induced by growth factors and cytokines and is elevated in many precancerous lesions and malignant tumors. COX-2 expression correlates closely with tumorigenesis, invasion, and metastasis, as it degrades the extracellular matrix and promotes tumor angiogenesis [17, 18]. Studies have shown that tissue COX-2 expression is significantly elevated in patients with PTC compared to those with HT or simple goiter [31]. Similarly, our immunohistochemical analysis revealed significantly higher COX-2 expression in PTC with HT as compared to HT alone. However, we observed no significant difference in COX-2 expression between uni- and multifocal PTC, precluding its use as a marker for multifocal PTC.

The RET/PTC-RAS-BRAF-MEK-ERK pathway (mitogen-activated protein kinase pathway or MAPK), which plays a fundamental role in proliferation, differentiation, apoptosis, and survival, is a key intracellular signaling pathway involved in thyroid carcinogenesis [32]. BRAF gene mutations and RET/PTC oncogene rearrangements are common in thyroid cancer. Studies suggest that thyroid carcinogenesis in HT is associated with the combined elevation in TSH and RET/PTC oncogene rearrangement. Rhoden et al. [33] detected RET/PTC oncogene rearrangements in HT and concluded that HT and PTC are closely related, possibly via HT-induced inflammation. The BRAF V600E mutation results in constitutive activation of BRAF [34, 35] and has been associated with aggressive PTC subtypes [32]. We observed a significant difference in the presence of the RET/PTC oncogene rearrangement between HT patients with uni- and multifocal PTC (20.9% vs. 56.8%, respectively; $P < 0.001$). The presence of the BRAF mutation also differed significantly between these groups (9.6% vs. 41.5%, respectively; $P < 0.001$). These promising results suggested that at the presence of the BRAF mutation or RET/PTC oncogene rearrangement is an indicator of

multifocal PTC. For patients with such test results, we recommend total bilateral thyroidectomy.

In conclusions, TPOAb > 1300 IU/mL, elevated TSH (only together with TPOAb > 1300 IU/mL), elevated CK-19 expression, and the presence of BRAF mutations and RET/PTC rearrangements are markers of multifocal PTC in HT and indicate the need for total bilateral thyroidectomy. Among HT patients with TPOAb > 1300 IU/ml, the occurrence of central lymph node metastasis is significantly higher in those with multi-focal than single-focus PTC, indicating the need for central lymph node dissection. These markers may prove useful for discerning between uni- and multifocal PTC, thereby preventing unnecessarily extensive surgery in the treatment of unifocal PTC and promoting sufficient treatment of multifocal PTC.

If the HT of patients are diagnosed to be accompanied with PTC using the preoperative B-ultrasound and fine needle aspiration (FNA), it suggests that the HT is likely to be accompanied with multi-focal PTC once one of the following conditions is fit: (1) TPOAb > 1300 IU/ml. (2) CK-19 immunohistochemical average optical density > 0.008 (3) BRAF gene mutation. (4) RET/PTC oncogene activation rearrangement is positive. Thus, it is recommended to perform total thyroidectomy combining the intraoperative frozen section at this time.

If the average optical density of CK-19 immunohistochemistry is more than 0.008, it is more likely to be HT multifocal cancer once one of the following conditions is fit: (1) TPOAb > 1300 IU/ml. (2) BRAF gene mutation. (3) RET/PTC oncogene activation rearrangement is positive. It is strongly recommended to perform total thyroidectomy.

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Disclosure of conflict of interest

None.

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Supplementary Table 1. Comparison of central cervical lymph node metastasis according to TPOAb concentration between groups

	HT with multifocal PTC N = 183 n (%)	HT with single PTC N = 187 n (%)	P
Central cervical lymph node metastasis according to TPOAb concentration			< 0.001
Type 1	9 (4.9)	85 (45.5)	
Type 2	50 (27.3)	5 (2.67)	
Type 3	44 (24.0)	80 (42.8)	
Type 4	80 (43.7)	17 (9.1)	

HT, Hashimoto's thyroiditis; PTC, papillary thyroid carcinoma; TPOAb, thyroid peroxidase antibodies. Type 1, negative for central lymph node metastasis with concentration of TPOAb < 1300 IU/mL; type 2, negative for central lymph node metastasis with concentration of TPOAb > 1300 IU/mL; type 3, positive for central lymph node metastasis with concentration of TPOAb < 1300 IU/mL; and type 4, positive for central lymph node metastasis with concentration of TPOAb > 1300 IU/mL.

Supplementary Table 2. Comparison of associated factors between multifocal and unifocal PTC in HT patients

	Univariate		Multivariate	
	OR (95% CI)	P	aOR (95% CI)	P
Clinical indicators				
TPOAb (IU/mL)				
≤ 1300	Reference		Reference	
> 1300	18.40 (10.64, 31.81)	< 0.001	83.11 (33.96, 203.44)	< 0.001
Central cervical lymph node metastasis				
Negative	Reference			
Positive	1.95 (1.28, 2.98)	0.002		
Tissue immunohistochemistry				
CK19 ^a	45.97 (16.65, 126.89)	< 0.001		
Gene detection				
BRAF				
Without mutation	Reference		Reference	
Mutation	6.67 (3.78, 11.77)	< 0.001	10.92 (4.64, 25.72)	< 0.001
RET/PTC				
Negative	Reference		Reference	
Positive	5.00 (3.16, 7.90)	< 0.001	13.36 (5.81, 30.73)	< 0.001

N = 370. HT, Hashimoto's thyroiditis; PTC, papillary thyroid carcinoma; TPOAb, thyroid peroxidase antibodies; CK19, cytokeratin 19; BRAF, B-type Raf kinase; RET/PTC, rearranged in transformation/papillary thyroid carcinoma; OR, odds ratio; aOR, adjusted odds ratio; CI, confidence interval. ^aData was transformed to log₁₀ because of the small values of the data.