



Vitamin D Receptor Expression and its Clinical Significance in Papillary Thyroid Cancer

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Abstract

Objective: This study aimed to evaluate the association between vitamin D receptor (an essential component in the vitamin D signaling pathway) and serum vitamin D as well as its clinical significance in papillary thyroid cancer. **Methods:** This prospective cohort study comprised patients with thyroid tumors who visited our hospital, from 2017 to 2018. The level of vitamin D receptor expression from thyroid tissue was measured in patients with thyroid tumor and evaluated for correlation with serum vitamin D levels and clinicopathologic characteristics of papillary thyroid cancer. Data from 501 patients with papillary thyroid cancer from The Cancer Genome Atlas database were analyzed. **Results:** Increased vitamin D receptor protein and mRNA expression were observed in papillary thyroid cancer compared to those in normal and benign tissues. Lower vitamin D receptor protein expression was associated with high TNM stage papillary thyroid cancer and low p21 protein expression. Lower relative vitamin D receptor mRNA expression in papillary thyroid cancer was associated with low serum 25-hydroxyvitamin D level. The Cancer Genome Atlas database showed a positive correlation among mRNA expression of vitamin D receptor, CYP24A1, and p21. **Conclusions:** An association between decreased vitamin D receptor protein expression and advanced stage papillary thyroid cancer, and a correlation between low vitamin D receptor mRNA expression with low serum 25-hydroxyvitamin D level was observed. Low vitamin D receptor expression in papillary thyroid cancer was shown to positively correlate with low serum vitamin D level and disease aggressiveness.

Keywords

vitamin D, vitamin D receptor, papillary thyroid cancer, surgery

Abbreviations

25(OH)D, 25-hydroxyvitamin D; AJCC, American Joint Committee on Cancer; BRAF, serine/threonine-protein kinase B-Raf; Calcitriol, 1,25-dihydroxyvitamin D₃, 1,25(OH)₂D₃; IHC, immunohistochemistry; miRNA, microRNA; RNA-Seq, mRNA sequencing; PTC, papillary thyroid cancer; qPCR, quantitative polymerase chain reaction; SNAIL, snail family transcriptional repressor; TCGA, The Cancer Genome Atlas; TNM, Tumor, Node, Metastasis; VDR, vitamin D receptor.

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Introduction

Since its discovery in the early 20th century, vitamin D has evolved from being considered as a simple vitamin to a steroid pro-hormone.¹ Previous studies have demonstrated that vitamin D deficiency, which is common worldwide, could be associated with non-skeletal conditions, which include muscle weakness; cardiovascular disorders; and metabolic, autoimmune, and infectious diseases.^{2,3} The associations between low serum 25-hydroxyvitamin D (25(OH)D), a biomarker of vitamin D status, and several malignancies like colorectal cancer, prostate cancer, and breast cancer have been reported, particularly focusing on the potential anticancer effects of vitamin D.⁴⁻⁶ In thyroid cancer, correlation between serum 25(OH)D level and its clinicopathologic characteristics has been controversial.⁷⁻¹⁰

Calcitriol (1,25-dihydroxyvitamin D₃, 1,25(OH)₂D₃), a potent activated form of vitamin D, is tightly regulated through a complex process involving the vitamin D-activating enzymes, 1- α -hydroxylase (also named CYP27B1) and 24-hydroxylase (also named CYP24A1). The enzyme 1- α -hydroxylase is responsible for the final hydroxylation step from 25(OH)D to 1,25(OH)₂D₃, while 24-hydroxylase is the key enzyme in the inactivation of 1,25(OH)₂D₃. The action of calcitriol occurs mainly through its binding to the nuclear vitamin D receptor (VDR), which acts as a hormone-regulated transcription factor.^{11,12}

Several studies have focused on the dynamics of VDR and calcitriol-related enzymes in cancer tissues,^{13,14} suggesting that VDR plays a critical role in the anticancer mechanism of calcitriol. VDR expression in cancer tissue has been shown to have an anticancer effect, and its clinical significance and prognostic value in several cancers has been consistently reported.¹⁵⁻¹⁹ Meanwhile, there have not been many reports on the relationship among serum vitamin D, VDR, and thyroid cancer. A few studies have discussed the overall vitamin D metabolism in normal, benign, and malignant thyroid tissues.²⁰⁻²²

Based on the studies mentioned above, we aimed to investigate VDR expression in papillary thyroid cancer (PTC) and evaluate its clinical significance. We hypothesized that low VDR expression in PTC may be associated with low serum vitamin D levels and aggressive clinicopathologic features. In this study, we first investigated the expression of VDR, CYP27B1, CYP24A1, and markers of cell proliferation (ie, p21, a cell cycle regulator) and metastasis (ie, E-cadherin, a cell-to-cell adhesion marker) in human thyroid tissues, from normal, benign, to PTC. Secondly, we correlated the expression profiles of VDR in PTC with serum vitamin D levels and other clinicopathologic characteristics.

Materials and Methods

A prospective cohort study was conducted on the consecutive patients with thyroid tumors who had visited our hospital

from April 2017 to July 2018. Patients diagnosed with PTC or benign thyroid tumor by fine-needle aspiration biopsy were included in this study. Patients were excluded if they were on medications that might alter vitamin D metabolism; had the following: a disease that could affect serum vitamin D levels, abnormal thyroid function, history of previous neck surgery or irradiation, any prior cancer history; or declined to participate in the study.

Thyroid surgery was determined based on clinical findings and performed by a single surgeon. Blood samples were obtained within one month before surgery. Serum levels of 25(OH)D, 1,25(OH)₂D₃ were measured simultaneously. According to recent criteria of an Endocrine Society clinical practice guideline, serum 25(OH)D levels <20 ng/mL, 20–29.9 ng/mL, and \geq 30 ng/mL are defined as deficient, insufficient, and sufficient, respectively. We categorized the patients into two groups based on their serum vitamin D levels, according to this guideline.²³ A 20 ng/mL cut-off for serum 25(OH)D was used to divide the patients into vitamin D deficient or non-deficient groups,²⁴⁻²⁶ and a 40 pg/mL cut-off was utilized for serum 1,25(OH)₂D₃ as a mean value of the reference range (20-60 pg/mL).²⁷

Normal, benign, and malignant thyroid tissues were obtained from patients for immunohistochemistry (IHC) and real-time quantitative polymerase chain reaction (qPCR). Biopsy samples were collected from a central location in thyroid tumors to obtain a pure tumor sample. Malignant and benign tumors were present simultaneously in four patients, from which both tissues samples were taken. Normal thyroid tissues were taken from the contralateral lobe of the thyroid tumor. Paraffin-embedded tissues from 92 patients and snap-frozen thyroid tissues from 68 patients were collected.

Clinical characteristics and demographic data were reviewed, extracting the following patient data: age at the time of surgery; sex; histologic type of primary tumor; tumor size, multiplicity, and bilaterality; extrathyroidal extension; BRAF (serine/threonine-protein kinase B-Raf) V600E mutations; and the Tumor, Node, Metastasis (TNM) classification. A cut-off value of \geq 1 cm was used for tumor size since a tumor size of <1 cm was used in the definition of papillary thyroid microcarcinoma. The TNM classification system of the American Joint Committee on Cancer (AJCC) (eighth edition) was used for the staging system. The T stage was classified into 1/2 or 3/4, the N stage into N0 versus N1, and the tumor stage into I/II or III/IV. This study was conducted in accordance with the Declaration of Helsinki and was approved by the Institutional Review Board of our hospital (4-2016-0657, approved on September 7, 2016). All patients provided written informed consent. Detailed patient information is not included in the manuscript. The reporting of this study conforms to the STROBE guidelines.

IHC staining for VDR, CYP27B1, CYP24A1, p21, and E-cadherin was performed on paraffin-embedded sections from 73 PTC, 23 benign, and 25 normal thyroid tissues.

Paraffin-embedded tissue specimens were cut into 4- μ m-thick sections and IHC was performed as previously described,²¹ using a Ventana Discovery XT Automated Slide Stainer (Ventana Medical System, Tucson, AZ, USA) with the following primary antibodies: anti-VDR antibodies (ab134826; Abcam, Cambridge, UK; 1:200), anti-CYP27B1 antibodies (sc515903; Santa Cruz Biotechnology, TX, USA; 1:200), anti-CYP24 antibodies (sc365700; Santa Cruz Biotechnology; 1:200), anti-p21 antibodies (ab109520; Abcam; 1:200), and anti-E-cadherin antibodies (ab15148; Abcam; 1:50).

IHC findings were interpreted by a single independent investigator (JSKoo), using an Olympus BX41 microscope (Olympus, Tokyo, Japan). Protein expression of nuclear VDR, cytoplasmic VDR, and nuclear p21 were quantified as the percent of stained nuclei per 100 cells. For cytoplasmic p21, CYP24A1, CYP27B1, and E-cadherin, staining intensity was scored as follows: 0 (no staining), 1 (weak), 2 (modest), or 3 (strong) (Fig. S1). In order to classify samples into high or low expression groups, we used the cut-off levels of 50% for nuclear and cytoplasmic VDR expression,²⁸ and 10% for nuclear p21 expression. For cytoplasmic p21, CYP27B1, CYP24A1, and E-cadherin, a score of 0–1 was categorized as negative, and a score of 2–3 was considered as positive (Figure 1).

Tissue samples obtained during thyroidectomy in 68 patients were immediately snap-frozen in liquid nitrogen and stored at -80°C . Total RNA was extracted using the RNeasy Plus Mini Kit (Qiagen, Hilden, Germany). RNA was quantified using a Spectrophotometer NanoDrop 2000 (Nanodrop Technologies, Wilmington, DE, USA). The acceptable purity as indicated by A260/280 was greater than 1.8, and RNA samples that showed smear or degradation were excluded. The fresh 38 PTC-normal pairs and 10 benign-normal pairs were subjected to qPCR analyses.²⁹

RNA was reversely transcribed to cDNA using an AccuPower RT Premix Kit (Bioneer Inc., Daejeon, South Korea) for experiments. qPCR was performed with an EvaGreen Q Master Mix (LaboPass, Seoul, Korea). Primer sequences for the target genes VDR, CYP27B1, CYP24A1, p21, and E-cadherin are listed in Table S1. Gene expression

was measured by qRT-PCR using a StepOnePlus™ real-time PCR machine (Applied Biosystems, CA, USA). β -Actin was used as an internal control.

In the present study, data is presented as the fold change in target gene expression of the tumors relative to its expression in the counterpart normal tissue. A cut-off level of two (a median value) for the N-fold differential expression of VDR was used to categorize the PTC patients into two groups to evaluate for correlation with clinicopathologic features.

Publicly available mRNA sequencing (RNA-Seq) data of 501 patients with thyroid cancer from The Cancer Genome Atlas (TCGA) database (version 2016_01_28; <https://gdac.broadinstitute.org>) were analyzed.^{30,31} RNA-Seq data of VDR, CYP27B1, CYP24A1, p21, and E-cadherin expressions of PTC were retrieved for further evaluation.

The baseline data is presented as the number and percentage for categorical variables and as the mean \pm standard deviation for continuous variables, unless otherwise specified. Continuous variables were compared using Student's t-test for two group comparisons. Categorical variables were compared using the chi-squared (χ^2) test or Fisher's exact test. The Pearson correlation coefficient was determined in TCGA RNA-Seq data. *P*-values < 0.05 were considered statistically significant. All data were processed and statistically analyzed using IBM SPSS Statistics for Windows, version 25.0 (IBM Corp., Armonk, NY, USA).

Results

Immunohistochemistry Staining

Vitamin D metabolism, cell proliferation, and metastasis were evaluated in 73 PTC, 23 benign, and 25 normal samples by comparing protein expressions of VDR, CYP27B1, CYP24A1, p21, and E-cadherin. A total of 25 pairs of cancer-normal tissues from the same patient were first compared (Table 1). Higher nuclear VDR expression was found in 68.0% of PTC, compared to 20.0% of normal thyroid tissues ($P = 0.001$). Positive CYP27B1 expression was significantly

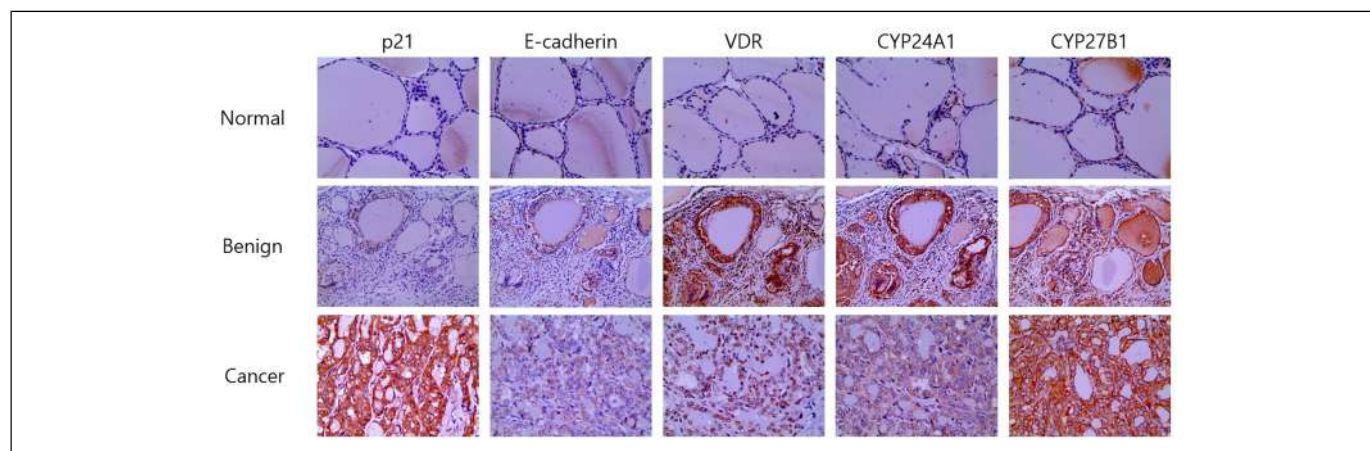


Figure 1. Protein expression profiles of VDR, CYP27B1, CYP24A1, p21, and E-cadherin in normal, benign, and papillary thyroid cancer (400 \times)

Table 1. Protein expression profiles of 25 paired papillary thyroid cancer and normal tissues.

Expression	Normal (n = 25)	Cancer (n = 25)	P value
VDR - Nuclear			
0–50	20 (80.0)	8 (32.0)	0.001
51–100	5 (20.0)	20 (68.0)	
VDR - Cytoplasmic			
0–50	25 (100.0)	25 (100.0)	NA
51–100	0 (0.0)	0 (0.0)	
CYP27B1			
Negative	12 (48.0)	2 (8.0)	0.004
Positive	13 (52.0)	23 (92.0)	
CYP24A1			
Negative	22 (88.0)	18 (72.0)	0.289
Positive	3 (12.0)	7 (28.0)	
P21 - nuclear			
0–10	24 (96.0)	18 (72.0)	0.049
11–100	1 (4.0)	7 (28.0)	
P21 - cytoplasmic			
Negative	22 (88.0)	18 (72.0)	0.289
Positive	3 (12.0)	7 (28.0)	
E-cadherin			
Negative	21 (84.0)	16 (64.0)	0.196
Positive	4 (16.0)	9 (36.0)	

Values are expressed as number (%).

VDR, vitamin D receptor.

higher in PTC compared to that in normal thyroid tissues (92.0% vs 52.0%; $P=0.004$). Nuclear p21 expression was also significantly increased in the PTC group compared to the normal thyroid group (28.0% vs 4.0%; $P=0.049$).

Analyses between 73 PTC and 25 normal tissues revealed a higher protein expression in nuclear VDR (57.5% vs 20.0%; $P=0.001$) and cytoplasmic VDR (23.3% vs 0%; $P=0.005$) in the PTC group (Table 2). There was a significant increase in nuclear p21 (41.0% vs 4.0%; $P<0.001$) and E-cadherin expression (52.1% vs 16.0%; $P=0.002$) in the PTC group compared to the normal tissue group.

We then compared 73 PTC and 23 benign thyroid tissue groups. The PTC group demonstrated higher protein expression in nuclear VDR (57.5% vs 33.3%; $P=0.009$), cytoplasmic VDR (23.3% vs 0%; $P=0.010$), nuclear p21 (41.0% vs 0%; $P=0.001$), and E-cadherin (52.1% vs 16.7%; $P=0.011$). Comparative analyses between 23 benign and 25 normal samples yielded no significant difference in the expression of target proteins.

The association of VDR protein expression with the clinicopathologic characteristics of 73 PTC patients was investigated (Table 3). High nuclear VDR expression was significantly associated with high nuclear p21 expression ($P=0.023$). A significant decrease in nuclear VDR expression was observed in PTC TNM stages 3 and 4 ($P=0.017$). There was no significant association between nuclear VDR and other clinicopathologic features, including serum 25(OH)D and 1,25(OH)₂D₃ levels. Serum vitamin D levels of both 25(OH)D and 1,25(OH)₂D₃ showed no significant correlation with other clinicopathologic features (Tables S2, S3).

Table 2. Protein expression profiles of the thyroid tissue samples, stratified by histology.

Expression	Normal (n = 25)	Benign (n = 23)	Cancer (n = 73)
VDR - Nuclear			
0–50	20 (80.0)	17 (73.9)	31 (42.5) **/#
51–100	5 (20.0)	6 (26.1)	42 (57.5)
VDR - Cytoplasmic			
0–50	25 (100.0)	23 (100.0)	56 (76.7) **/#
51–100	0 (0.0)	0 (0.0)	17 (23.3)
CYP27B1			
Negative	12 (48.0)	11 (47.8)	24 (32.9)
Positive	13 (52.0)	12 (52.2)	49 (67.1)
CYP24A1			
Negative	22 (88.0)	21 (91.3)	62 (84.9)
Positive	3 (12.0)	2 (8.7)	11 (15.1)
P21 - nuclear			
0–10	24 (96.0)	22 (95.7)	43 (58.9) **/###
11–100	1 (4.0)	1 (4.3)	30 (41.1)
P21 - cytoplasmic			
Negative	22 (88.0)	22 (95.7)	64 (87.7)
Positive	3 (12.0)	1 (4.3)	9 (12.3)
E-cadherin			
Negative	21 (84.0)	18 (78.3)	35 (47.9) **/#
Positive	4 (16.0)	5 (21.7)	38 (52.1)

Values are expressed as number (%).

**p < 0.01 versus normal, #p < 0.05 and ###p < 0.01 versus benign.

VDR, vitamin D receptor.

We further investigated the correlation between the protein expression of nuclear VDR and other markers in PTC tissues and found no significant association (Table 3).

Real-Time Quantitative PCR (qPCR)

Comparative analysis was conducted for the relative tumor-normal mRNA expression values in 38 PTC and 10 benign tissues, along with their normal counterparts. The VDR gene showed a higher relative expression in PTC compared to that in benign tumors (20.24 ± 7.63 vs 3.41 ± 1.58 ; $P=0.037$). The other genes yielded no evident difference in relative mRNA expression between PTC and benign tumors (Figure 2).

We examined the correlation of relative cancer-normal mRNA expression of VDR, CYP27B1, CYP24A1, p21, and E-cadherin with VDR protein expression in PTC. Increased mRNA expression of VDR in cancer was observed in patients with higher (> 50) nuclear VDR protein expression (31.21 ± 12.16 vs 3.43 ± 1.04 ; $P=0.033$). The other genes showed no association with nuclear VDR protein expression (Figure 3).

We further investigated the relationship between the relative cancer-normal mRNA expression level of VDR genes and the clinicopathologic characteristics of 38 PTC patients. The group with ≤ 2 -fold cancer-normal VDR mRNA expression ratio had a lower mean serum 25(OH)D level (22.42 ± 9.56 vs 16.01 ± 5.56 ; $P=0.017$). However, serum 1,25(OH)₂D₃ levels were similar, regardless of the relative VDR mRNA expression levels (48.81 ± 25.67 vs 47.50 ± 19.43 , $P=0.860$) (Table 4).

Table 3. Clinicopathological characteristics of patients with papillary thyroid cancer according to vitamin D receptor expression level.

	Nuclear VDR Expression		P-value
	0–50% (N = 31), n (%)	51–100% (N = 42), n (%)	
Age (yr), mean ± SD	46.74 ± 14.21	43.98 ± 13.22	0.395*
Age (yr)			
<55	20 (64.5)	34 (81.0)	0.114 [†]
≥55	11 (35.5)	8 (19.0)	
Sex			
Male	9 (29.0)	13 (31.0)	0.860 [†]
Female	22 (71.0)	29 (69.0)	
25(OH)D (ng/mL)	19.41 ± 8.40	18.00 ± 7.30	0.446*
25(OH)D (ng/mL)			
<20	22 (71.0)	30 (71.4)	0.966 [†]
≥20	9 (39.0)	12 (28.6)	
1,25(OH) ₂ D ₃ (pg/mL)	40.39 ± 18.66	48.56 ± 22.05	0.100*
1,25(OH) ₂ D ₃ (pg/mL)			
<40	16 (51.6)	15 (35.7)	0.174 [†]
≥40	15 (48.4)	27 (64.3)	
Tumor size (cm)	1.29 ± 0.98	1.25 ± 0.82	0.866*
Tumor size			
≤1cm	16 (51.6)	22 (52.4)	0.948 [†]
>1cm	15 (48.4)	20 (47.6)	
Multifocality			
Negative	23 (74.2)	30 (71.4)	0.793 [†]
Positive	8 (25.8)	12 (28.6)	
Bilaterality			
Negative	27 (87.1)	36 (85.7)	1.000 [†]
Positive	4 (12.9)	6 (14.3)	
Extrathyroidal extension			
Negative	10 (32.3)	16 (38.1)	0.607 [†]
Positive	21 (67.7)	26 (61.9)	
T-stage			
T1-T2	11 (35.5)	18 (42.9)	0.525 [†]
T3-T4	20 (64.5)	24 (57.1)	
Regional lymph node			
N0	17 (54.8)	25 (59.5)	0.689 [†]
N1	14 (45.2)	17 (40.5)	
Distant metastasis			
M0	30 (96.8)	39 (92.9)	0.632 [†]
M1	1 (3.2)	3 (7.1)	
TNM stage group			
I-II	18 (58.1)	35 (83.3)	0.017 [†]
III-IV	13 (41.9)	7 (16.7)	
BRAF mutation			
Absent	6 (19.4)	7 (16.7)	0.767 [†]
Present	25 (80.6)	35 (83.3)	
VDR - Cytoplasmic			
Negative to positive 0–50%	21 (67.7)	35 (83.3)	0.119 [†]
Strong Positive 51–100%	10 (32.3)	7 (16.7)	
CYP27B1			
Negative 0–1	10 (32.3)	14 (33.3)	0.923 [†]
Positive 2–3	21 (67.7)	28 (66.7)	

(continued)

Table 3. (continued).

	Nuclear VDR Expression		P-value
	0–50% (N = 31), n (%)	51–100% (N = 42), n (%)	
CYP24A1			
Negative 0–1	27 (87.1)	35 (83.3)	0.657 [†]
Positive 2–3	4 (12.9)	7 (16.7)	
P21 - Nuclear			
Negative 0–10%	23 (74.2)	20 (47.6)	0.023 [†]
Positive 11–100%	8 (25.8)	22 (52.4)	
P21 - Cytoplasmic			
Negative 0–1	30 (96.8)	34 (81.0)	0.069 [†]
Positive 2–3	1 (3.2)	8 (19.0)	
E-cadherin			
Negative 0–1	17 (54.8)	18 (42.9)	0.311 [†]
Positive 2–3	14 (45.2)	24 (57.1)	

*P-values were calculated using Student's t-test. Data are expressed as mean ± SD.

[†]P-values were calculated using χ^2 test. Data are expressed as number (%).25(OH)D, 25-hydroxyvitamin D; 1,25(OH)₂D₃, 1,25-dihydroxyvitamin D₃ (calcitriol).

TCGA Thyroid Cancer Data According to the VDR mRNA Expression status

Owing to restricted mRNA expression profiles in our analyses, we obtained RNA-Seq data of 501 PTC patients from the TCGA database. We investigated the relationship of VDR mRNA expression profiles with CYP27B1, CYP24A1, p21, and E-cadherin. VDR mRNA expression demonstrated a positive association with p21 ($r = 0.109$, $P = 0.015$) and CYP24A1 ($r = 0.215$, $P < 0.001$), and a negative association with E-cadherin ($r = -0.129$, $P = 0.004$). VDR mRNA expression showed no significant relationship with CYP27B1 ($r = -0.05$, $P = 0.269$) (Figure 4).

Discussion

In this study, we identified relevant components of vitamin D metabolism and their effect in thyroid cancer. The presence of VDR in normal thyroid tissue has been previously described.³² PTC samples showed enhanced protein and mRNA expressions of VDR and vitamin D related enzymes when compared to normal and benign human thyroid tissue, suggesting a potential antitumor response.^{20,21} Our study evaluated the protein and mRNA expression profiles of VDR, CYP24A1, and CYP27B1 in normal, benign, and PTC tissues and assessed their anti-proliferation, anti-adhesion, and anti-invasion characteristics in cancer cells.

During IHC analysis, we evaluated VDR protein expression in different cellular compartments (ie, nucleus and the cytoplasm). Only a few studies have evaluated the clinical significance of either nuclear or cytoplasmic VDR expression in cancer.^{33–35} One study reported that IHC staining was generally cytoplasmic in thyroid cancer and was more intense near the tumor capsule.²⁰ Another study revealed that nuclear VDR

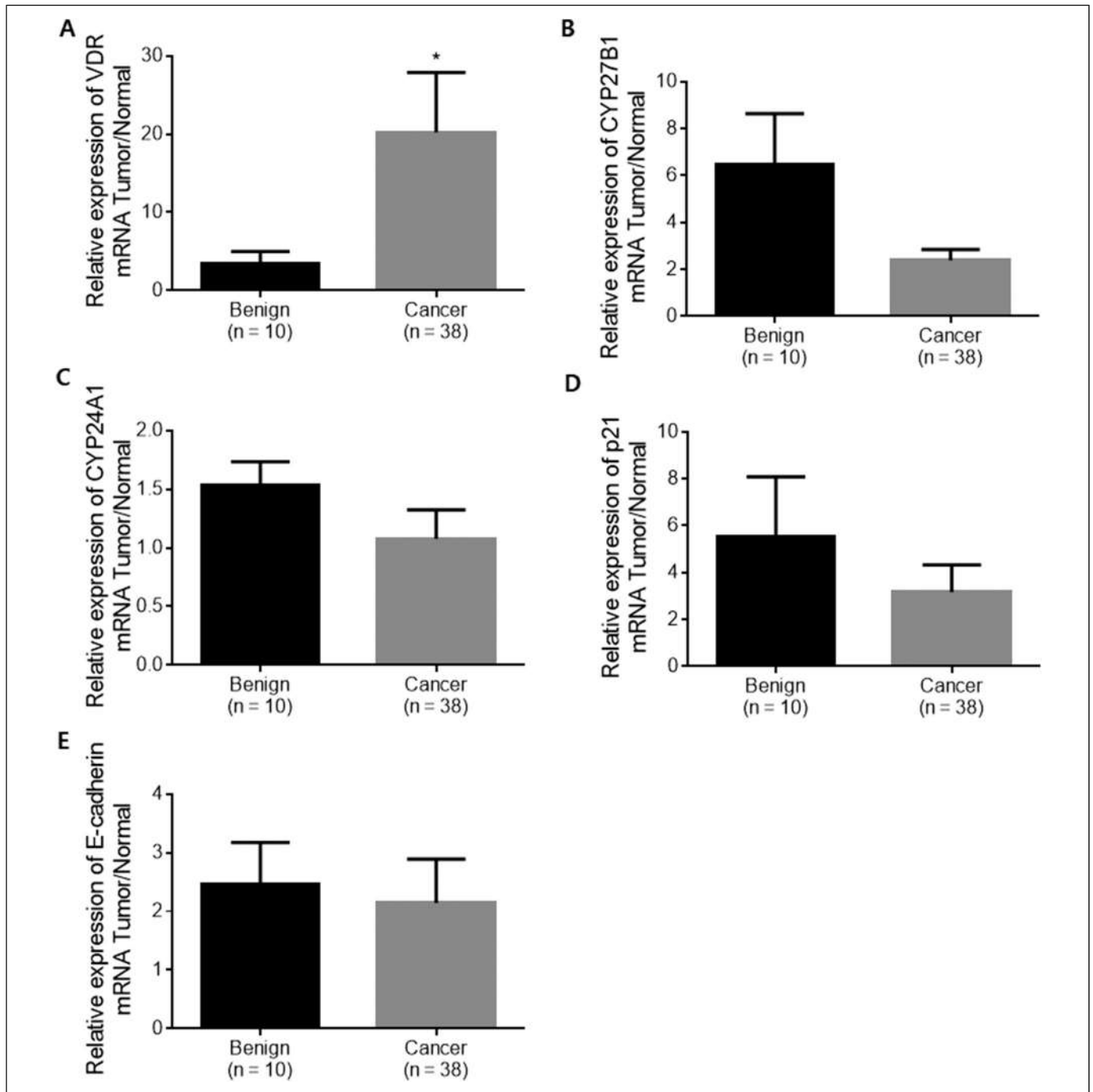


Figure 2. Relative tumor-normal mRNA expression levels of the target genes (A) VDR, (B) CYP27B1, (C) CYP24A1, (D) p21, and (E) E-cadherin in papillary thyroid cancer and benign tumor

Data are expressed as the mean \pm standard error of the mean; * $p < 0.05$; VDR, vitamin D receptor.

expression in PTC samples negatively correlated with STAT3 hyperphosphorylation, which indicates worse clinicopathologic characteristics.³⁵

In our study, nuclear VDR expression was higher than cytoplasmic VDR expression in most PTCs, compared to those in normal and benign samples. Cytoplasmic VDR expression was significantly enhanced in PTC than in normal and benign tissues, and it was detected only in few

patients. The behavior of VDR seems consistent throughout the vitamin D pathway and may contribute to the anticancer activity of calcitriol.^{12,36} Furthermore, nuclear VDR expression was higher than cytoplasmic VDR, implying that VDR activity occurs mainly in the nucleus. Nuclear VDR expression was significantly decreased in TNM stage 3 and 4 PTC, indicating that calcitriol-VDR complex had a reduced anticancer effect.

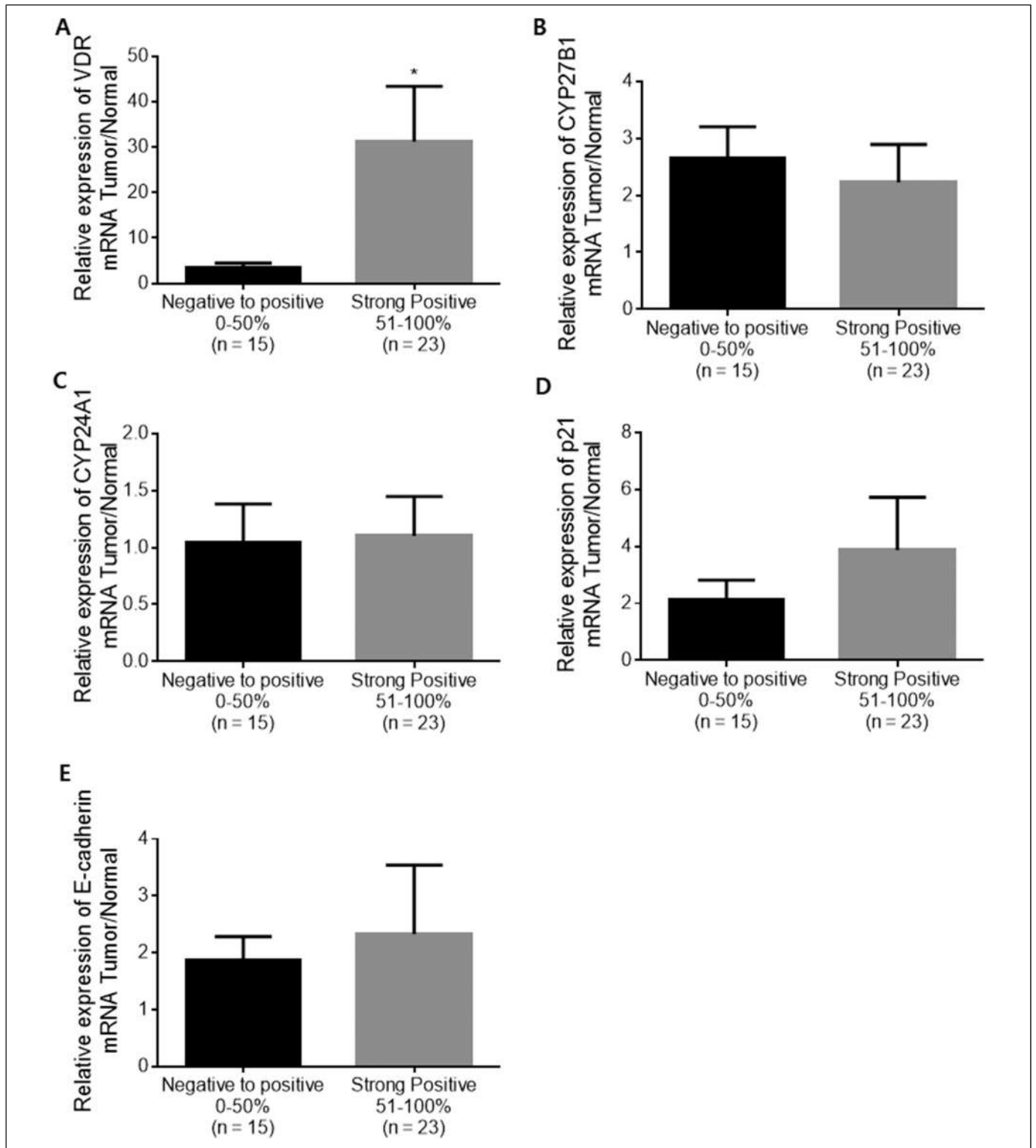


Figure 3. Correlation between the mRNA expression profile of the target genes (A) VDR, (B) CYP27B1, (C) CYP24A1, (D) p21, and (E) E-cadherin with VDR protein expression

Data are expressed as the mean \pm standard error of the mean; *p < 0.05; VDR, vitamin D receptor.

The comparison of 73 PTC with 25 normal samples revealed elevated CYP27B1 and CYP24A1 expression in PTC compared to those in normal tissue, although this was not

statistically significant. Previous studies have demonstrated elevated CYP27B1 and VDR expression can lead to a magnified vitamin D action in thyroid cancer.^{20,21} This study's result of

Table 4. Clinicopathological characteristics of patients with papillary thyroid cancer according to vitamin D receptor mRNA relative expression level.

	VDR mRNA relative expression level		P-value
	T/N < 2 (N = 19), n (%)	T/N ≥ 2 (N = 19), n (%)	
Age (yr), mean ± SD	48.81 ± 12.66	42.53 ± 12.61	0.731*
Age			
<55	16 (84.2)	15 (78.9)	1.000 [†]
≥55	3 (15.8)	4 (21.1)	
Sex			
Male	4 (21.1)	6 (31.6)	0.714 [†]
Female	15 (78.9)	13 (68.4)	
25(OH)D (ng/mL)	16.01 ± 5.56	22.42 ± 9.56	0.017*
25(OH)D (ng/mL)			
<20	16 (84.2)	9 (47.4)	0.038 [†]
≥20	3 (15.8)	10 (52.6)	
1,25(OH) ₂ D ₃ (pg/mL)	40.39 ± 18.66	48.56 ± 22.05	0.100*
1,25(OH) ₂ D ₃ (pg/mL)			
<40	8 (42.1)	7 (36.8)	0.740 [†]
≥40	11 (57.9)	12 (63.2)	
Tumor size (cm)	1.06 ± 0.54	1.57 ± 1.28	0.117*
Tumor size			
≤1cm	10 (52.6)	9 (47.4)	0.746 [†]
>1cm	9 (47.4)	10 (52.6)	
Multifocality			
Negative	12 (63.2)	13 (68.4)	0.732 [†]
Positive	7 (36.8)	3 (31.6)	
Bilaterality			
Negative	13 (68.4)	17 (89.5)	0.232 [†]
Positive	6 (31.6)	2 (10.5)	
Extrathyroidal extension			
Negative	7 (36.8)	7 (36.8)	1.000 [†]
Positive	12 (63.2)	12 (63.2)	
T-stage			
T1-T2	7 (36.8)	8 (42.1)	0.740 [†]
T3-T4	12 (63.2)	11 (57.9)	
Regional lymph node			
N0	13 (68.4)	12 (63.2)	0.732 [†]
N1	6 (31.6)	7 (36.8)	
Distant metastasis			
M0	18 (94.7)	18 (94.7)	1.000 [†]
M1	1 (5.3)	1 (5.3)	
TNM stage group			
I-II	13 (68.4)	13 (68.4)	1.000 [†]
III-IV	6 (31.6)	6 (31.6)	
BRAF mutation			
Absent	5 (26.3)	3 (15.8)	0.693 [†]
Present	14 (73.7)	16 (84.2)	
VDR - Nuclear			
Negative to positive 0–50%	7 (36.8)	8 (42.1)	0.740 [†]
Strong Positive 51–100%	12 (63.2)	11 (57.9)	
VDR - Cytoplasmic			
Negative to positive 0–50%	15 (78.9)	14 (73.7)	1.000 [†]
Strong Positive 51–100%	4 (21.1)	5 (26.3)	
CYP27B1			
Negative 0–1	5 (26.3)	5 (26.3)	1.000 [†]
Positive 2–3	14 (73.7)	14 (73.7)	
CYP24A1			
Negative 0–1	18 (94.7)	15 (78.9)	0.340 [†]
Positive 2–3	1 (5.3)	4 (21.1)	

(continued)

Table 4. (continued).

	VDR mRNA relative expression level		P-value
	T/N < 2 (N = 19), n (%)	T/N ≥ 2 (N = 19), n (%)	
P21 - Nuclear			
Negative 0–10%	9 (47.4)	10 (52.6)	0.746 [†]
Positive 11–100%	10 (52.6)	9 (47.4)	
P21 - Cytoplasmic			
Negative 0–1	17 (89.5)	16 (84.2)	1.000 [†]
Positive 2–3	2 (10.5)	3 (15.8)	
E-cadherin			
Negative 0–1	12 (63.2)	7 (36.8)	0.105 [†]
Positive 2–3	7 (36.8)	12 (63.2)	

* P-values were calculated using Student's t-test. Data are expressed as mean ± SD.

[†]P-values were calculated using χ^2 test. Data are expressed as number (%).

25(OH)D, 25-hydroxyvitamin D; 1,25(OH)₂D₃, 1,25-dihydroxyvitamin D₃ (calcitriol).

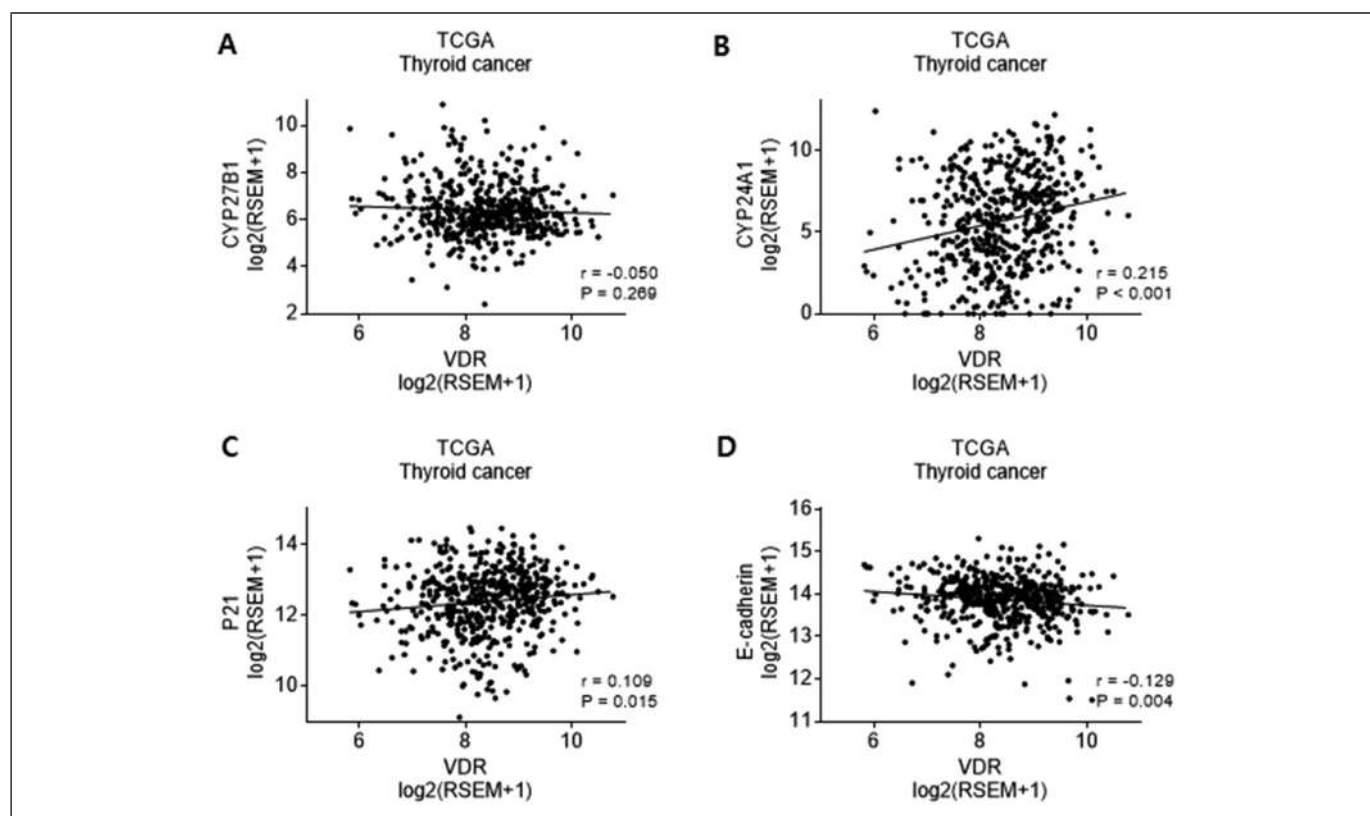


Figure 4. mRNA expression of (A) CYP27B1, (B) CYP24A1, (C) p21, and (D) E-cadherin in the RNA-Seq data of 501 patients with thyroid cancer from The Cancer Genome Atlas database

* $p < 0.05$; VDR, vitamin D receptor.

CYP27B1 elevation in PTC compared to that in paired normal tissue is similar to previous studies.

VDR expression was increased in PTC than in normal and benign tissues, both in protein and mRNA levels. The relative mRNA expression level of VDR in PTC was higher than the protein expression level, which might have resulted from the compartmentalization of VDR protein expression in the nucleus and cytoplasm. This could lead to a greater increase in overall VDR mRNA expression. In addition, higher VDR

protein expression in PTC was evident in patients with high cancer-normal mRNA expression ratio. This supports the augmented expression of VDR in PTC throughout the experiments. These findings are consistent with findings of previous studies that described the potential anticancer effect of VDR in thyroid cancer.^{20,21}

Calcitriol is identified to regulate specific signaling pathways through which it plays roles in anti-proliferation, pro-apoptosis, de-differentiation, anti-inflammation, anti-angiogenesis, and

anti-invasion and anti-metastasis of cancer.^{12,37,38} Recently, Pang *et al* suggested that VDR knockdown attenuates the anti-proliferative, pro-apoptotic, and anti-invasive effect of vitamin D in PTC by activating the Wnt/ β -catenin signaling pathway.³⁹ Zhang *et al* showed that calcitriol enhances doxorubicin-induced apoptosis in PTC cells by regulating VDR/PTPN2/p-STAT3 pathway.³⁵ We hypothesized that increased VDR expression in PTC tissue is caused by similar mechanisms, which may also be impaired in advanced stage PTC.

Previous literature demonstrated that calcitriol inhibits cell proliferation through cell cycle arrest by activating p21 and p27, particularly in the G0/G1 phase.^{40,41} p21 has been recognized for its pro-apoptotic activity in the nucleus and anti-apoptotic activity in the cytoplasm.^{42,43} We evaluated p21 separately according to nuclear and cytoplasmic expression. The human p21 gene contains VDR binding promoter regions and is a transcriptional target of calcitriol-VDR complex.^{37,44} Liu *et al* demonstrated that calcitriol-induced p27 activation in thyroid cancer cells was accomplished by VDR-mediated regulation of p27 phosphorylation and degradation.⁴¹ In our study, p21 mRNA expression did not demonstrate notable findings, but the positive correlation of nuclear p21 and nuclear VDR protein expression may develop into potential anti-proliferative effect in thyroid cancer.

Several studies have reported that E-cadherin is involved in the invasion and metastasis of thyroid cancer, but the results were inconsistent.⁴⁵ Even though the role of E-cadherin in thyroid cancer remains unclear, its elevated protein expression in PTC suggests its anticancer potential according to our study. Previous studies have shown that the VDR activation by calcitriol induces E-cadherin expression by promoting the translocation of β -catenin from the nucleus to the plasma membrane and inhibiting the Wnt/ β -catenin/TCF4 signaling pathway. This also supports its anti-invasion and anti-metastasis roles in thyroid cancer.^{45,46}

Analyses of 501 PTC samples from the TCGA database revealed significant positive correlation between VDR mRNA expression, and CYP24A1 and p21. A previous study reported decreased VDR and CYP24A1 mRNA expression in the PTC N1 stage, accompanied by a decreased p21 expression.²¹ Although the TCGA database was based on mRNA expression level, we observed a similar correlation in the protein level analyses of our study, with decreased nuclear VDR and nuclear p21 protein expression and advanced PTC TNM stage. Although the N stage was not significantly correlated, the findings still explain the loss of anti-proliferative, dedifferentiating functions in aggressive thyroid cancer.

Repressed VDR action may be explained by impaired vitamin D and VDR signaling pathways. Studies have investigated the counteraction of cancer cells via multiple mechanisms to restrain VDR expression, such as in colon and breast cancer. These include the snail family transcriptional repressor (SNAIL) which inhibits transcription, and RAS oncogene mutation which suppresses its transcription.^{47,48} The tumor-suppressor gene, p53, is known to enhance VDR transcription. However, p53 mutant cells can regulate VDR responses by directly binding to VDR and redirecting its transcriptional

program to apoptosis.^{49,50} Others include epigenetic gene silencing, CpG island methylation, and microRNA (miRNA).⁵¹ In thyroid cancer, the mechanism for decreased VDR expression in advanced stages is not clearly defined; hence, further investigation is needed.

In this study, the low VDR mRNA expression in PTC compared to that in normal tissues was associated with a low serum 25(OH)D level. VDR protein expression was not associated with serum vitamin D levels and no clinicopathologic significance in PTC was found with serum vitamin D levels. We assume that this disparity primarily comes from the compartmentalized expression of VDR protein and relative mRNA expression level. Several factors such as unstable environmental factors, unstable co-binding proteins, and patient factor can affect both protein and gene expressions.¹⁴

Nevertheless, the positive correlation between relative VDR mRNA expression and serum 25(OH)D level in PTC implies similar anticancer effect in thyroid cancer. There have been varying results in the correlation between serum vitamin D level and the aggressiveness and prognosis of thyroid cancer.^{7-10,52} There are only a few reports on the relationship of serum vitamin D levels and VDR in different cancers.^{53,54} Since serum 25(OH)D is the best biomarker for vitamin D status, the positive correlation between VDR mRNA expression and serum 25(OH)D level can be a convincing indication of our hypothesis.

This study has several limitations. First, a majority of the patients (71.2%, 51/72) were deficient in serum 25(OH)D. Secondly, for the analysis, serum vitamin D levels were not strictly adjusted for BMI and seasonal variation. Thirdly, in the mRNA expression analyses, we excluded a substantial number of tissues during the qualification control procedure for accurate results, resulting in a smaller number of specimens. A small sample size of 10 benign tumors would especially have led to results with selection bias, since surgery is recommended in large symptomatic benign tumors. Fourthly, assessing the association between serum vitamin D, VDR expression and patient prognosis was not feasible in this study.

Conclusion

In this study, we demonstrated elevated protein and mRNA expression of VDR in PTC compared to normal and benign tissues. However, lower protein expression of nuclear VDR was identified in high TNM stage PTC, which was associated with low nuclear p21 protein expression. This study provides further evidence for the potential anti-proliferative effects of VDR in PTC, which is diminished in aggressive thyroid cancer. Moreover, lower VDR mRNA expression in PTC was associated with low serum 25(OH)D levels. Overall, this is the first report to identify the possibility of a positive correlation between low VDR expression, low serum 25(OH)D level, and aggressiveness of thyroid cancer.

We believe that this study will contribute to a better understanding of vitamin D metabolism and the clinical significance of VDR expression in PTC. Furthermore, large prospective

studies are needed to validate the potential anticancer effect of VDR in thyroid cancer and establish the use of VDR expression and serum vitamin D levels as a prognostic indicator. The therapeutic potential of vitamin D in thyroid cancer should be further investigated in future research.

Ethics Statement

This study was conducted in accordance with the Declaration of Helsinki and was approved by the Institutional Review Board of the Severance Hospital (4-2016-0657). All patients provided written informed consent.

Data Availability Statement

Some or all data generated or analyzed during this study are included in this published article or in the data repositories listed in References.

Supplemental material

Supplemental material for this article is available online.

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