Case report: poorly differentiated thyroid carcinoma in a young female patient

Shane Kaysen[1]^, Peter Anderson[2], Jeffrey Mohlman[3]

[1]Rocky Vista University College of Osteopathic Medicine, Parker, CO, USA; [2]South Valley ENT and Steward Healthcare, Salt Lake City, UT, USA; [3]Professional Diagnostics Inc. and Steward Healthcare, Salt Lake City, UT, USA

Correspondence to: Jeffrey Mohlman, MD, MPH. Jordan Valley Medical Center, 3580 W 9000 S West Jordan, UT 84088, USA. Email: Jeffrey.mohlman@steward.org.

Background: Poorly differentiated thyroid carcinoma (PDTC) is a rare subtype of thyroid cancer that is often overlooked on differentials for thyroid neoplasms, and its etiology is poorly understood. This report provides an additional clinical and genetic profile to the literature of this rare and often complex entity as well as discusses a multitude of genetic findings seen in PDTC.

Case Description: A young female patient (<21) presented to her primary care provider with a history significant for a large, asymptomatic, right sided, immobile neck mass. A CT scan revealed a mass measuring 4.0×4.0×2.6 cm^3 lying adjacent to the thyroid. The mass was subsequently removed via a right hemithyroidectomy. The histology demonstrated a diffuse trabecular, and solid growth pattern consisting of poorly differentiated follicular cells with mitotic figures, capsular invasion, lymphovascular invasion, nerve involvement and focal areas of necrosis. Immunohistochemical stains were performed and were useful in determining the tumor's thyroid origin. Multiple pathology consults were obtained in the workup for this case along with the application of the Turin criteria. The patient was recommended to undergo a complete thyroid resection and seek genetic counseling due to a known association between this disease and a rare syndrome known as DICER syndrome. Genetic studies did not identify a *DICER1* mutation, and the patient was lost to follow-up after referral to a tertiary care center.

Conclusions: This tumor type is rare in this age group and has few studies explaining its etiology. The genetic analysis in this case had a unique genetic profile showing *MYC* amplification and mutations of undetermined significance without the previously reported finding of a *DICER1* mutation.

Keywords: Case report; poorly differentiated thyroid carcinoma (PDTC); DICER syndrome; Turin criteria; *MYC* amplification

Introduction

Poorly differentiated thyroid carcinoma (PDTC) is a rare form of thyroid cancer that represents 4–7% of all thyroid carcinomas and, rarer still, is the occurrence in those under the age of 21 (1). The etiology of the disease is largely unknown but thought to arise from well differentiated thyroid carcinoma (WDTC) with the potential to advance to anaplastic thyroid carcinoma (ATC). This case report seeks to present and discuss PDTC in the form of patient presentation, workup, microscopic findings, the Turin criteria, and genetic analysis to further the understanding of PDTC, specifically in patients <21. It will largely focus on genetic findings which include, a lack of a *DICER1* mutation, a *MYC* amplification, and a variety of mutations

^ ORCID: 0000-0003-4418-8589.

Figure 1 Transverse (A) and sagittal (B) CT images of the mass located in the right thyroid lobe. Gross specimen post-resection measuring 4.0×3.5×2.5 cm^3 (C).

with unknown clinical significance. We present the following case in accordance with the CARE reporting checklist (available at https://aot.amegroups.com/article/view/10.21037/aot-21-20/rc).

Case presentation

A young previously healthy female (<21) patient presented with a large firm neck mass right of midline and adjacent to the thyroid gland without other signs or symptoms. Family history was negative for pertinent findings. A CT scan showed the mass appeared potentially contiguous with the right thyroid lobe with mild tracheal compression (*Figure 1*). Fine needle aspiration and cytology reported mixed solid, acinar, and focal microfollicular architecture. It was thus signed out as being most consistent with a benign adenomatous nodule with recommended correlation with imaging. The mass was resected with a right hemithyroidectomy and sent for further histopathological evaluation which included frozen sections, standard H&E

staining, and immunohistochemistry. The patient was subsequently referred to a tertiary cancer treatment center and follow up information was unable to be obtained. Despite multiple attempts, informed consent was unable to be obtained from the patient as follow-up was lost. The patient's contact information was unreliable at the time this case report was written. All procedures performed in this study were in accordance with the ethical standards of the institutional and/or national research committee(s) and with the Helsinki Declaration (as revised in 2013).

Timeline of events

Late 2020:

❖ Patient presents complaining of a solid neck mass, confirmed with palpation and visual inspection by PCP. Referred for ultrasound evaluation.

❖ Ultrasound identifies a homogenous hypoechoic neck mass with prominent vasculature, without definitive involvement of the right thyroid lobe.

FNA is ordered.

❖ FNA and subsequent cytopathologic findings give the preliminary diagnosis favoring a benign follicular lesion. Patient is referred to a surgeon for mass resection.

Early 2021:

❖ Surgeon orders a CT scan which confirms size and location of the mass in addition to mild tracheal compression. The patient remained asymptomatic during this time.

❖ A right hemithyroidectomy is performed and the mass is removed without complications.

❖ Initial frozen sections were deferred to permanent due to the complexity of the lesion.

❖ Permanent sections were then sent from the community hospital setting to a university hospital. An expert on thyroid pathology is also consulted during this time.

❖ The three entities agree on the final diagnosis of PDTC, likely arising from a prior papillary thyroid carcinoma (PTC), follicular type.

❖ Genetic analysis is recommended due to the rarity of this tumor in this age group, looking for a *DICER1* mutation.

❖ Patient is also referred to a tertiary cancer treatment center and follow-up is lost.

Mid 2021:

❖ Genetic testing is completed and no *DICER1* mutation was found. Instead, a *MYC* amplification and multiple other mutations are identified. These findings are reported out and an addendum to the case is made.

Macroscopic description

A 22 g red 4.0×3.5×2.5 cm³ nodule. Sectioning the specimen showed a fleshy, red-tan cut surface.

Microscopic description

Frozen section slides showed a follicular lesion, but definitive diagnosis was deferred to permanents. Permanent histological examination reveals an insular, solid, and trabecular growth pattern (*Figure 2*) comprised of small, poorly formed follicles with loss of PTC nuclear features in a vast majority of lesions with several lesser areas showing PTC-like nuclear features. Within the trabecular and solid architecture there are multiple foci of necrosis and

dispersed mitotic figures. There is a background of normal looking thyroid follicles scattered sparsely throughout with colloid remnants. Notable is the tumor's invasion of both the lymphovasculature and nervous tissue in the periphery (*Figure 3*). Although partially encapsulated, the tumor is freely infiltrative. Individual cells have normal to enlarged nuclei, prominent nucleoli, and an abundance of eosinophilic cytoplasm. The nuclei are relatively round with and are lacking nuclear grooves, Pseudonuclear inclusion bodies, and nuclear clearing (Orphan-Annie eye).

Immunohistochemistry staining showed TTF1 positive follicular cells in the lesion, helping rule out ATC. PAX8 was also positive confirming the lesion is of thyroid origin. Staining was negative for synaptophysin and chromogranin (*Figure 4*). A Congo red stain was performed, and polarization was negative for amyloid.

Diagnosis and therapeutic intervention

The frozen section diagnosis given was: Follicular lesion, defer to permanents. The final diagnosis was PDTC likely arising from a PTC, follicular variant.

An extensive workup was used to diagnose the mass including ultrasound, a core needle biopsy, CT scan with contrast, and ultimately surgical excision followed by multiple pathology consults. The patient continued to be asymptomatic and recovered well from the right hemithyroidectomy. No extrathyroidal extension was identified clinically. It was recommended the patient follow-up to completely resect the remaining thyroid and seek genetic testing for a *DICER1* mutation. Neoadjuvant chemotherapy is also commonly used to treat the disease, although it is unclear if this patient pursued this option.

Genetic testing on the formalin fixed paraffin embedded tissue (NeoTYPE Discovery Profile for Solid Tumors from Neogenomics Laboratories) was negative for suspected *DICER1* mutation that can be seen in young patients with PDTC that carries an elevated risk of developing additional cancers (2). The panel used tests for well over 100 biomarkers and resulted in pertinent negatives such as *BRAF*, *BRCA1*, *BRCA2*, *EGFR*, *KIT*, *KRAS*, *NRAS*, *PDGFRA*, *PIK3CA*, *ALK* FISH, *MET* FISH, *PDGFRe* FISH, *PTEN* FISH, *RET* FISH, and *ROS1* FISH. Other pertinent findings from the genetic analysis included a *MYC* amplification by fluorescent in situ hybridization (FISH) with 4 or more *MYC* signal per nucleus in at least 40% of tumor cells (48%) consistent with gene amplification (*Figure 5*). Additional copies of
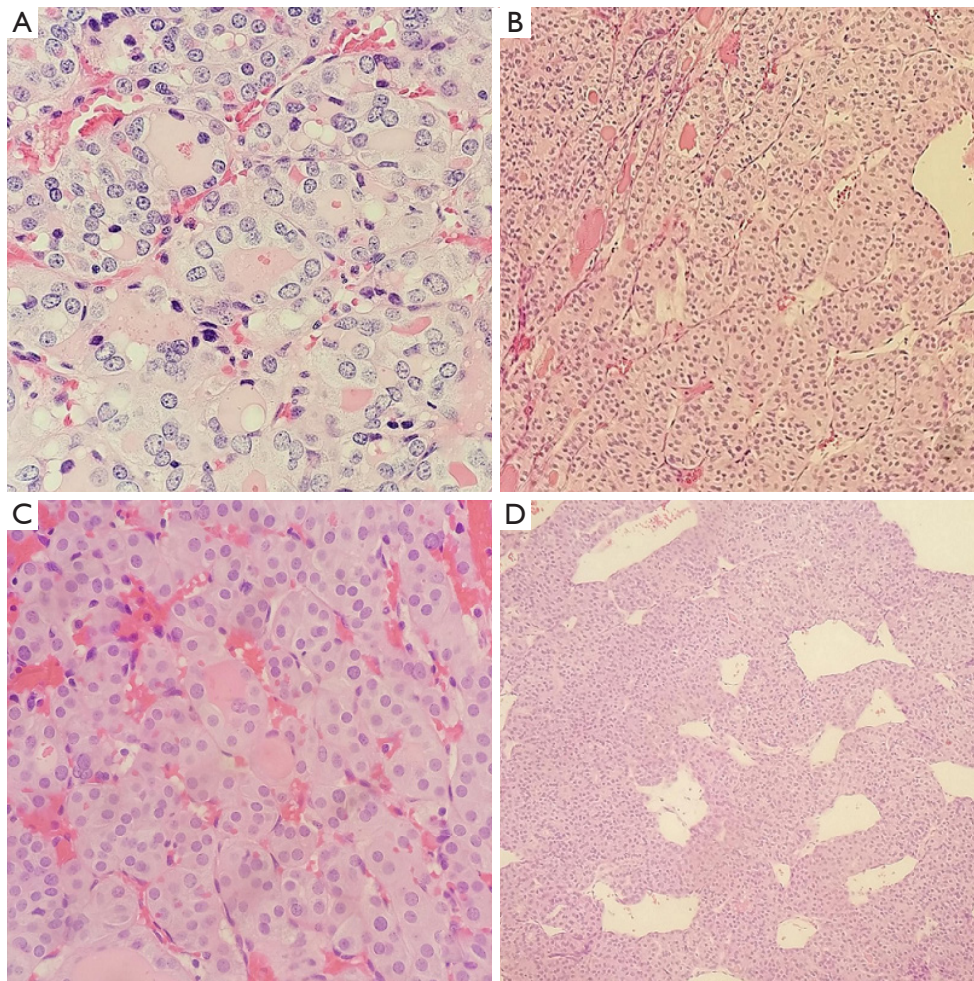
Figure 2 Solid (A, 400× magnification) trabecular (B, 200× magnification and C, 400× magnification) and insular (D, 100× magnification) patterns from various tumor sections. All images are stained with hematoxylin and eosin (H&E).

BRAF (>2F, 80.0%, negative <25.1%) [Fusion cutoff, % of cells counted, Negative cutoff], *RET* (>2F, 24.0%, negative <16.2%), and *PTEN* (>2R>2G, 32.0%, negative <21.8%) were observed which is an abnormal result suggestive of gains of *BRAF* (7q34) or chromosome 7, *RET* (10q11), *PTEN* (10q23) or chromosome 10. No additional known mutations of clinical significance were identified; although 5 variants of unknown clinical significance were present in the *BLM* (D88E NM_000057.4:c.264C>A), *KMT2A* (E495K NM_005933.4:c.1483G>A), *MPL* (T469I NM_005373.3:c.1406C>T), *NF2* (R338C NM_000268.3:c.1012C>T) and *PRKDC* (A2345V NM_006904.7:c.7034C>T) gene loci. Prognosis in this case of a young patient without a *DICER1* mutation is still largely unknown (3).

Discussion

PDTC is a relatively rare and aggressive form of thyroid cancer making up approximately 4–7% of all thyroid carcinomas. PDTC occurs most commonly in older populations with the mean age of clinical detection being 57 years with a predominately female predilection (2). It typically presents as a painless thyroid mass averaging 4.0 cm in diameter at initial presentation and may affect one or both thyroid lobes. It may arise from a precursor lesion or without a prior lesion. Precursor lesions include WDTC such as follicular or PTC as in this case. The diagnosis can often be missed, lending itself to being the highest cause of fatalities from non-anaplastic thyroid entities if not considered in the differential (4-6). PDTC looks and behaves as an intermediate between
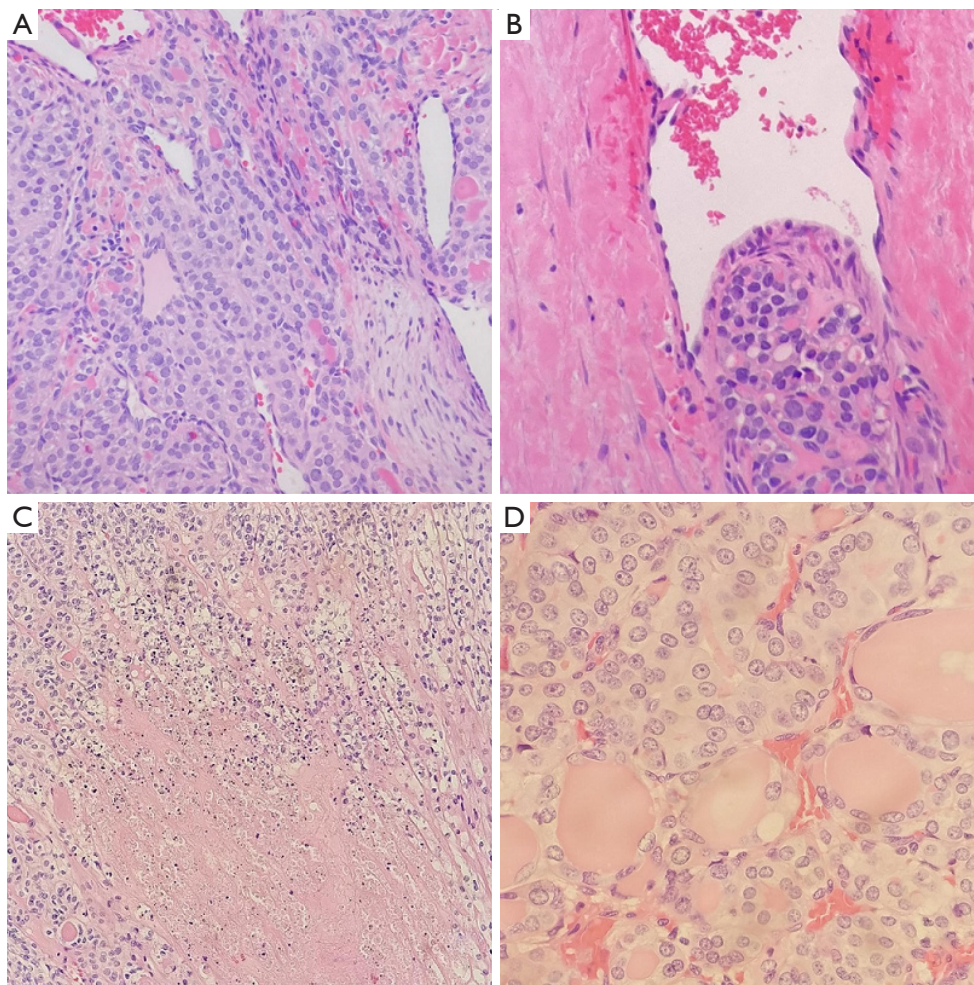
Figure 3 Nerve invasion (A, 200×), lymphovascular invasion (B, 400×), necrosis (C, 200×), and background papillary thyroid carcinoma (D, 400×). All images are stained with H&E.

WDTC and ATCs. Extrathyroidal, regional, and distant spread are common, with studies showing 70%, >50%, and 43% respectively (7). The etiology of this tumor is poorly understood.

The Turin criteria (3 features) are used for definitive diagnosis of PDTC. The growth pattern is shown to be trabecular, solid, or insular (Turin criteria 1). There must be no nuclear features of PTC (Turin criteria 2) and the tumor must have at least one of the three following features: convoluted nuclei, ≥3 mitotic figures per 10 high power fields (HPF), or necrosis (Turin criteria 3) (7). These criteria are necessary due to the ambiguous histopathologic features of the tumor as it can have focal areas of WDTC amongst prominent areas of atypia that do not reach the levels seen in ATC.

Given the rarity of this tumor type in younger age groups, it was suspected to be associated with DICER syndrome caused by a mutation in the *DICER1* gene. The *DICER1* gene is responsible for encoding the DICER endoribonuclease protein that is important in the control of protein translation, specifically microRNAs (miRNAs). As a component of the RNA induced silencing complex (RISC), DICER's role is to transform the single stranded pre-miRNAs into double stranded RNAs that can then be transported and bind to specific mRNAs. The binding of the miRNA to the mRNA prevents the attachment of ribosomes and subsequently, any protein translation that would have followed (2,8). It is thought that the absence of a functional DICER protein may contribute to either a loss of function in tumor suppressor genes, or a gain-of-function
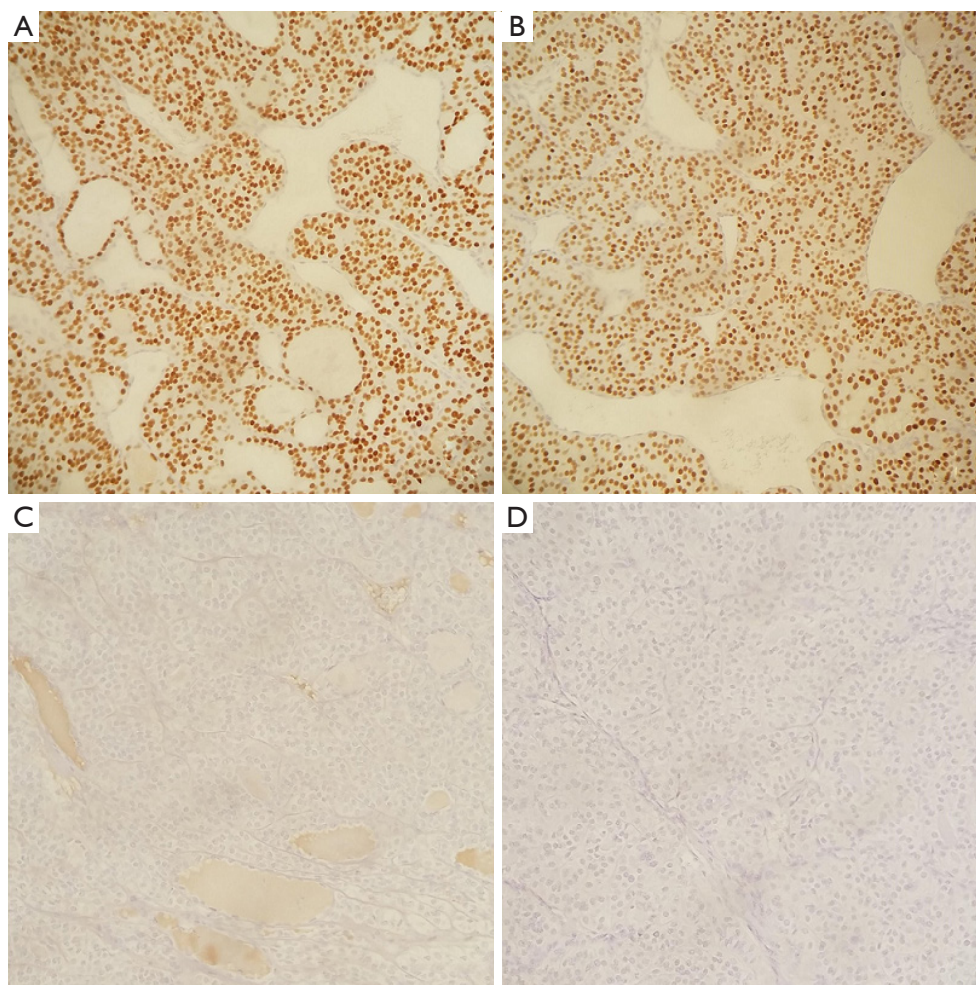
Figure 4 Immunohistochemistry: PAX8 stain (A, 200×), TTF1 stain (B, 200×), synaptophysin stain (C, 200×), and chromogranin stain (D, 200×).

in oncogenic genes that are regulated by the presence of inhibitory miRNA. Defective, downregulated, or even over expression of miRNAs resulting from a *DICER1* mutation may lead to the development of cancer, or a poor prognosis in cancer (9-11).

DICER syndrome is a rare disorder characterized by a predisposition to multiple cancers including but not limited to Sertoli-Leydig tumors, pleuropulmonary blastoma, multinodular goiter, cystic nephromas, and PDTC (2). One study showed that PDTC in young patients is likely due to a distinct etiology associated with *DICER1*, although its sample size was extremely limited. 5 out of 6 patients <21 years with PDTC were found to have *DICER1* mutations and no additional known mutations associated with other thyroid neoplasms. Three out of the five patients died

within 8–24 months due to their disease (3). In this case, it is recommended to do a genetic evaluation in order to better assess the patient's potential for developing additional malignancies and overall prognosis.

A notable finding in this case was the presence of a *MYC* amplification using FISH. *MYC* is a well-known proto-oncogene that shows overexpression in approximately 20% of all human cancers and is a poor prognostic indicator (12,13). Chernock *et al.* (3) found that 1 of 6 patients <21 years diagnosed with PDTC did not have a *DICER1* mutation, but it was not mentioned if there were any additional mutations or other genetic abnormalities that may be implicated in oncogenesis. Current studies have not shown a direct connection between MYC and PDTCs, but it can be assumed as *MYC* amplifications have been found in
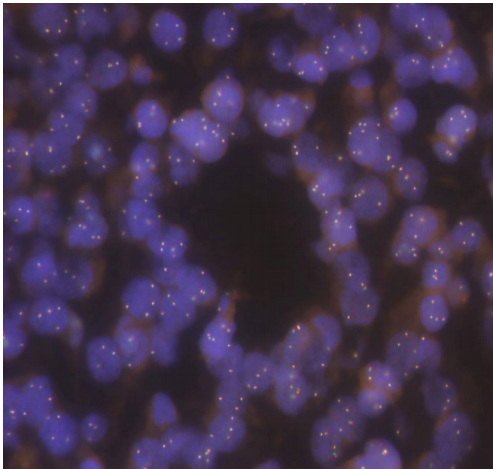
Figure 5 *MYC* amplification as shown by fluorescent *in-situ* hybridization showing greater than 4 MYC signals per cancer cell nucleus.

PTCs and may encourage dedifferentiation. These findings support prior evidence that PDTC can be a progression from WDTC (14). The *MYC* amplification's significance in this case can be implied by its frequent association with other cancers, but definitive significance is uncertain. Unfortunately, since follow-up with the patient was lost, it cannot be determined what the clinical outcome of the treated disease was, as well as the implications of the genetic findings. As research on PDTC continues, the etiology of the disease may become clearer, especially in its poorly understood occurrence in younger populations. Currently our understanding of PDTC etiology in children is limited, and additional studies focused on genetic markers may prove useful.

Notably, a recently published case of a metastatic PDTC arising from a PTC in a 28-year-old patient had shown a germline *MET* mutation (15). The *MET* gene encodes a receptor tyrosine kinase, which in tandem with the MET ligand, hepatocyte growth factor (HGF), stimulates the MET-HGF pathway resulting in cellular proliferation, mitosis, apoptosis and motility (15,16). Mutations in the well-known *MET* oncogene can also be found in a variety of cancers not unlike *MYC* gene amplifications and may play a role in their dedifferentiation. Our case was negative for any *MET* abnormalities.

Additional findings of undetermined clinical significance in the genetic analysis of this patient include mutations in the *KMT2A*, *NF2*, *BLM*, *MPL*, and *PRKDC* genes. *KMT2A* is an epigenetic regulator and is one of the most frequently mutated histone methyltransferases. Mutations of this gene are considered clinically insignificant, but its mutations have been implicated in the pathogenesis of thyroid cancer (17). According to AACR project Genie, *NF2* mutations are common in ATC (8.06%) and not in PDTC (0.64%). This is for patients with ATC that have *NF2* mutations (in the tumor) without NF2 disease (18). *BLM* and *MPL* mutations have also been identified in thyroid cancer, although their correlation is unknown. *BLM* encodes for helicase enzymes and *MPL* encodes for thrombopoietin receptor protein (19). PRKDC is a DNA double stranded break repair enzyme, and its amplification has been implicated in a variety of cancers. There is no documented correlation with the development of thyroid cancer. These variant mutations in addition to *MYC* amplification may support the overall picture of PDTC arising from a previous WDTC through acquired mutations. Research by Landa *et al.* shows evidence of cumulative mutations as a major factor in the development of PDTC or ATC that may be of clinical value (20).

This case provides a histopathological overview of PDTC as well as relevant insight into the possible etiology of this poorly understood disease. Although one small study has shown a connection between PDTC and DICER1 syndrome, this case did not, and instead describes another means for young patients to develop this tumor type. We hope this case report is useful in further defining genetic risk factors and assists in identification of PDTC, especially in young patients.

Practice points

PDTC is a rare thyroid cancer with its own distinct criteria for diagnosis. It mainly presents in older adults but can occur in children and adolescents.

The etiology of PTDC can be associated with a genetic component such as *DICER1* mutations, or in this case, *MYC* amplification. It can arise from a preexisting well-differentiated thyroid carcinoma or *de novo*.

DICER1 mutation should be strongly suspected when a PTDC is diagnosed in children or adolescents, although more research is needed as this case shows that it is not always present.

The Turin criteria help a great deal in diagnosing this type of cancer due to its ambiguity.

Self-assessment questions

(I) Which IHC stain's positivity can help differentiate

between PDTC and ATC while maintaining its specificity for a tumor of thyroid origin?
    A. PAX8
    B. TTF1
    C. Chromogranin
    D. Synaptophysin
    E. Calcitonin
(II) Per the TURIN guidelines, which of the following is one of the diagnostic criteria for PDTC?
    A. Lymphovascular invasion
    B. Distant metastasis
    C. Presence of nuclear features such as nuclear clearing, nuclear grooves, and pseudonuclear inclusion bodies.
    D. >3 mitotic figures/10 HPF.
    E. Arising from a background of a WDTC.

Answers

(I) B—TTF1 (ATC almost always loses its TTF1 staining but retains PAX8).
(II) D—>3 mitotic figures/10 HPF (one of the 3 sub-criteria with the other two being necrosis and convoluted nuclei. These can be present either individually or together).

appropriately investigated and resolved. All procedures performed in this study were in accordance with the ethical standards of the institutional and/or national research committee(s) and with the Helsinki Declaration (as revised in 2013). Informed Consent was unable to be obtained from the patient as follow-up was lost. The patient's contact information was unreliable at the time this case report was being written.

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