Differentiated Thyroid Carcinoma in Pediatric Age: Genetic and Clinical Scenario

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Introduction: Follicular-derived differentiated thyroid carcinoma (DTC) is the most common endocrine and epithelial malignancy in children. The differences in the clinical and pathological features of pediatric vs. adult DTC could relate to a different genetic profile. Few studies are currently available in this issue, however, and most of them involved a limited number of patients and focused mainly on radiation-exposed populations.

Materials and Methods: We considered 59 pediatric patients who underwent surgery for DTC between 2000 and 2017. RET/PTC rearrangement was investigated with fluorescent in situ hybridization and real-time polymerase chain reaction. Sequencing was used to analyze mutations in the BRAF, NRAS, PTEN, PIK3CA genes, and the TERT promoter. The pediatric patients’ clinical and molecular features were compared with those of 178 adult patients.

Results: In our pediatric sample, male gender and age <15 years coincided with more extensive disease and more frequent lymph node and distant metastases. Compared with adults, the pediatric patients were more likely to have lymph node and distant metastasis, and to need second treatments (p < 0.01). In all, 44% of the pediatric patients were found to carry molecular alterations. RET/PTC rearrangement was confirmed as the most frequent genetic alteration in childhood DTC (24.6%) and correlated with aggressive features. BRAFV600E was only identified in 16% of the pediatric DTCs, while NRASQ61R, NRASQ61K, and TERTC250T mutations were very rare.

Conclusions: Pediatric DTC is more aggressive at diagnosis and more likely to recur than its adult counterpart. Unlike the adult disease, point mutations have no key genetic role.

Keywords: childhood, thyroid cancer, BRAF, TERT, RAS
INTRODUCTION

Thyroid carcinoma is the most common malignant neoplasm of the endocrine system. Papillary (PTC) and follicular (FTC) carcinomas are the most frequently seen histological types (accounting for more than 90% of cases) and belong to the family of differentiated carcinomas (DTC) originating from follicular thyroid cells (1).

Though less frequent than the adult type, DTC can also develop in pediatric age. There are significant clinical, pathological, and molecular differences between pediatric and adult patients, however. DTC in the pediatric population must consequently be characterized as a distinct entity with a separate diagnostic pathway and different treatment (2).

At the time of diagnosis, children are more likely than adults to present with advanced disease (3, 4). The incidence of distant metastases is estimated to be 25% in children, with lymph node involvement ranging from 40 to 80% in various studies (3, 5, 6). Positive lymph nodes and distant metastases from thyroid cancers are amenable to surgical resection and radioactive iodine (RAI) therapy, and these interventions have produced favorable outcomes in the pediatric patient population (5).

Our study aimed to characterize pediatric DTC in a large monocentric series of patients, most of them not exposed to radiation, focusing on: (i) clinical features and outcomes; (ii) molecular profile, with particular reference to the study of point mutations of the \( \text{BRAF} \), \( \text{RAS} \), \( \text{PTEN} \) genes, and \( \text{RET/PTC} \) translocations; (iii) correlations between clinical and molecular findings; and (iv) comparisons between the clinical and molecular profile of pediatric DTC and that of a large series of adult thyroid carcinoma cases coming from our center.

MATERIALS AND METHODS

Patients

Patients under 18 years of age with a histological diagnosis of DTC were retrospectively reviewed for this study, enrolling a series of 59 pediatric patients who underwent thyroid surgery between 2000 and 2017. Thirty of the Fifty-Nine patients (51%) also underwent neck dissection to remove the central/lateral lymph nodes.

All study samples were retrospectively selected from the electronic archives of the Surgical Pathology and Cytopathology Unit at Padua University. Clinical and histopathological data were obtained from electronic databases.

All patients underwent total thyroidectomy. Only information concerning their clinical presentation, pathological staging and histology was available for 8 patients; for the other 51, we had information concerning their clinical presentation, pathological staging and histology, local and systemic therapy, recurrences, subsequent treatments, and outcome. The follow-up was a mean 7 years (minimum 1, maximum 16), median 5.9 years.

After surgery, RAI ablation therapy was administered to 48/51 patients (94%), with a median dose of 100 mCi (range 30–200 mCi). Judging from their medical history, only 2 patients been exposed to radiation during childhood for the treatment of a previous tumor: one patient underwent fractioned total body irradiation (TBI) followed by allogenic bone marrow transplantation for acute lymphoblastic leukemia 11 years before the DTC development; the other patient underwent fractioned TBI plus chemotherapy followed by allogenic bone marrow transplantation for severe aplastic anemia 4 years before the DTC development.

The pediatric patients’ clinical and molecular characteristics were compared with those of 178 consecutive adult patients who underwent surgery for PTC between 2007 and 2010, and were followed up at the Radiotherapy Unit and Endocrinology Unit in Padua. The TNM 7th edition (7) was applied to all patients to ensure a correct comparison of their pathological and clinical characteristics.

All studies were performed in accordance with the guidelines proposed in the Declaration of Helsinki: the local ethical committee (Ethical Committee for the Clinical Experimentation of the Hospital of Padua) approved our study protocol (Ref. 3388) and all patients (including the parent/guardian on behalf of the minor) gave their written informed consent.

Outcomes

Patients were classified as being in remission if, at the time of their latest follow-up, their suppressed thyroglobulin (Tg) was <1 ug/L, Tg antibodies were negative, neck US was free of suspicious signs, and there were no pathological findings on any other imaging studies performed for clinically indicated reasons or any other modality or in any biopsy specimen. Patients with persistent disease at the time of their latest follow-up were classified as having either biochemically or structurally persistent disease, i.e., those with suppressed and/or stimulated Tg levels >1 ug/L but no structural evidence of disease were classified as having biochemically persistent disease, while those with structural evidence of disease (with or without abnormal Tg levels) were classified as having structurally persistent disease. Clinical status at final follow-up reflected not only the initial response to total thyroidectomy and RAI ablation therapy, but also the potential effects of continued levothyroxine suppressant therapy, any additional surgery or radiation therapy, and the passage of time.

Molecular Analysis

**BRAF, NRAS, PTEN, PIK3CA, and TERT Analysis**

For all patients, DNA was extracted from formalin-fixed, paraffin-embedded (FFPE) tissues. The samples were deparaffinized as follows: 30 min in xylene; 15 min in 99% alcohol, 15 min in 95% alcohol; 15 min in 70% alcohol; 15 min in 50% alcohol; several washes in distilled water. DNA was extracted from the sample using the “DNeasy Blood and Tissue Handbook” extraction kit (Qiagen, Milano, Italy), according to the manufacturer’s protocol. The \( \text{BRAF} \) gene (exon 5), \( \text{NRAS} \) (exon 3), \( \text{PTEN} \) (exons 5 and 8), \( \text{PIK3CA} \) (exons 9 and 20), and the \( \text{TERT} \) promoter were amplified using the polymerase chain reaction (PCR) technique. Mutation analyses were performed using direct sequencing ABI PRISM (Applied Biosystems, Foster City, California), as described elsewhere (8).
**RET/PTC Rearrangement Analysis**

RET/PTC translocations were investigated with both fluorescence in situ hybridization (FISH) and real-time polymerase chain reaction (RT-PCR).

To identify RET/PTC rearrangements (either 10q11.2 inversions or translocations), FISH was performed using the REPEAT-FREE POSEIDON RET (10q11) break-apart probe (Kreatech Diagnostics, Amsterdam, Netherlands) on FFPE samples. The FISH procedure was completed according to the Kreatech protocol with some modifications, as previously described (9). Slides were examined using an Olympus BX-61 microscope (Olympus, Hamburg, Germany) with appropriate fluorescence excitation/emission filters. Signals were recorded by a CCD camera (Olympus Digital Camera). For microscopic examination, at least 100 intact and non-overlapping cell nuclei were scored for the presence of a split signal. Only cells with two overlapping signals, or one split and one overlapping signal, were counted to ensure that only complete cell nuclei had been scored. The signal pattern was interpreted as follows: interphase nuclei with two co-localized green/red fusion signals identified normal chromosomes 10, while separate red and green signals and green/red fusion signals indicated rearranged RET. To establish the cut-off for RET/PTC rearrangements, we performed a FISH analysis on ten samples of normal thyroid parenchyma and 100 nuclei were scored for the presence of a split signal. As previously reported, the cut-off was calculated as the mean + 3 SD of RET rearranged cells (10).

For the RT-PCR analysis, tissue cores were deparaffinized with xylene at 50°C for 3 min. Total RNA extraction was done using the RecoverAll kit (Ambion, Austin, Texas, USA) according to the manufacturer’s instructions. The nucleic acid extracted was analyzed with RT-PCR in an ABI-PRISM 7900HT Sequence Detector (Applied Biosystems, Milan, Italy) with the EntroGen Thyroid Cancer Mutation Analysis Panel Kit (EntroGen, Inc., Woodland Hills, California), which detects RET/PTC1 (fusion between RET and CCDC6 genes), and RET/PTC3 (fusion between RET and NCOA4 genes). Fusion detection reactions were performed with a 1-step procedure that combines complementary DNA synthesis and RT-PCR. The resulting RT-PCR amplification curves were visualized using Sequence Detection Software rel. 2.4 (Applied Biosystems).

**Statistical Analysis**

The personal, clinical, and histopathological data were summarized using rates and percentages. All statistical analyses were performed using the MedCalc software (rel. 11.6.0). The distributions of the continuous variables were examined and the data were consequently summarized. The Mann-Whitney test and the t-test for independent samples were used to assess the differences in continuous variables (age at diagnosis, tumor size, and follow-up) between the two groups (pediatric vs. adult patients). The χ²-test was used to compare the variables within subgroups of the same series. Multivariate analysis was performed, using logistic regression, to confirm the independent role of different histopathological variables associated with final outcome. Differences were considered statistically significant when p was < 0.05.

**RESULTS**

**Clinical Features of Pediatric DTC Cases**

The main clinical features of each pediatric patient are shown in Table 1.

Average age at diagnosis was 14.4 ± 2.9 years (range 6.4–17.8 years): 29/59 patients (49%) were <15 years old (defined as “children”), and 30/59 (51%) were ≥15 (defined as “adolescents”). There were 16/59 males (27%), and 43/59 females (73%), with a male/female ratio of 1:2.7.

At histological analysis, 42/59 (71.2%) cases were the classic variant of PTC (cv-PTC), 7/59 (11.8%) were the follicular variant of PTC (fv-PTC), 4/59 (6.8%) were the sclerosing variant of PTC (sv-PTC), 1/59 (1.7%) was the tall cell variant of PTC (tcv-PTC), 3/59 (5.1%) were cases of follicular thyroid carcinoma (FTC), and 2/59 (3.4%) were poorly-differentiated thyroid carcinomas (PDTC).

Data on tumor size were available for 56/59 (95%) patients (no clear definition of tumor size was provided for 3 patients). The median size of the primary tumor was 20.0 mm (range 3–58 mm): 9/56 tumors (16.0%) were ≤10 mm (microcarcinomas), and 47/56 (84.0%) were >10 mm in size. As suggested by Nikita et al., the lesions’ size was further classified as follows: 26/56 (46.4%) were <20 mm; 27/56 (48.2%) were 20–40 mm; and 3/56 (5.4%) were >40 mm (11).

Multifocality was detected in 25/59 patients (42.4%), extra-thyroid extension in 25/59 (42%), and vascular invasion in 48/59 (81.4%). There were lymph node metastases in the lateral compartment in 20/59 patients (33.9%). Distant metastases were identified on WBS in 10/48 patients who underwent RAI: the site involved was the lung in 9/10 and the mediastinum in 1/10.

No significant differences in tumor size emerged between the male and female patients (with a median tumor size of 24.6 mm and 20.1 mm, respectively; p = 0.23). The extent of the disease (T) was gender-related, however: 10/16 males (63%) had T3 or T4 disease, while the proportion among females was 22/43 (51%; p < 0.01). Gender was also significantly associated with the presence of distant metastases, and therefore with the most advanced stage: 6/16 (38%) males had metastatic disease, as opposed to only 4/43 (9%) females (p = 0.03). No significant associations came to light between histotype, multifocality, lymph node involvement, need for second treatment, or final disease status.

Table 2 shows the main pathological and clinical characteristics correlated with the presence/absence of metastases for all 59 patients. The 8 patients for which we had no information about their treatment and follow-up were included because they started with a low-risk disease. They were considered as M0-x.

The presence of distant metastases was found significantly associated with younger age (p = 0.01), larger and more extensive tumors (p < 0.01), and cervical lymph node metastases (p = 0.02).

On multivariate analysis, only T4 status was identified as an independent predictor of distant metastases (OR 43.75, CI: 4.04 to 474.27).
### TABLE 1 | Clinic pathological features.

| Patient | Age at the diagnosis (years) | Gender | Tumor size (Mm) | Histological variant | T | N | M | Second treatment | Disease status |
|---------|-----------------------------|--------|----------------|---------------------|---|---|---|-----------------|----------------|
| 1       | 17                          | F      | 8              | cv-PTC              | 1a| X | X | 0               | BD             |
| 2       | 17                          | M      | 30             | fv-PTC              | 2 | 0 | 0 | 0               | Excellent      |
| 3       | 17                          | F      | 25             | cv-PTC              | 2 | 1b| 0 | Surgery/RAI     | Excellent      |
| 4       | 17                          | F      | 25             | cv-PTC              | 3 | 0 | 0 | n.a.            | n.a.           |
| 5       | 17                          | M      | 10             | cv-PTC              | 1a| 1a| 0 | n.a.            | n.a.           |
| 6       | 9                           | F      | 6              | sv-PTC              | 1a| 1a| 0 | Excellent       | Excellent      |
| 7       | 11                          | F      | 21             | cv-PTC              | 3 | 1b| 0 | Excellent       | Excellent      |
| 8       | 17                          | F      | 22             | cv-PTC              | 2 | 1a| 0 | Surgery         | Excellent      |
| 9       | 18                          | F      | 13             | cv-PTC              | 1b| X | 0 | n.a.            | n.a.           |
| 10      | 18                          | F      | 12             | cv-PTC              | 1b| 1a| 0 | Excellent       | Excellent      |
| 11      | 16                          | F      | 24             | cv-PTC              | 3 | 1b| 0 | RAI             | BD             |
| 12      | 15                          | F      | 15             | cv-PTC              | 3 | 1b| 1 | RAI             | Excellent      |
| 13      | 17                          | F      | 25             | cv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 14      | 13                          | M      | n.a.           | PDTC                | 4a| 1b| 1 | RAI/Surgery     | BD             |
| 15      | 13                          | F      | 50             | PDTC                | 3 | 1b| 1 | RAI             | SD             |
| 16      | 8                           | M      | 20             | cv-PTC              | 3 | 1b| 1 | RAI             | SD             |
| 17      | 8                           | F      | 20             | cv-PTC              | 3 | 0 | 0 | n.a.            | n.a.           |
| 18      | 16                          | F      | 12             | cv-PTC              | 1b| 1b| 0 | 0               | Excellent      |
| 19      | 17                          | F      | 3              | cv-PTC              | 1a| X | 0 | n.a.            | n.a.           |
| 20      | 14                          | M      | 12             | cv-PTC              | 1b| 1b| 0 | 0               | Excellent      |
| 21      | 15                          | M      | 35             | FTC                 | 2 | X | 1 | 0               | Excellent      |
| 22      | 12                          | F      | 36             | cv-PTC              | 3 | 1b| 1 | Surgery/RAI     | SD             |
| 23      | 17                          | F      | 18             | cv-PTC              | 3 | 1a| 0 | 0               | Excellent      |
| 24      | 17                          | F      | 13             | cv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 25      | 9                           | M      | 42             | cv-PTC              | 4a| 1 | 1 | RAI             | SD             |
| 26      | 14                          | F      | 40             | cv-PTC              | 3 | 1a| 0 | Surgery         | BD             |
| 27      | 13                          | F      | 11             | cv-PTC              | 3 | 1a| 0 | Surgery         | Excellent      |
| 28      | 14                          | F      | 20             | cv-PTC              | 3 | 1b| 0 | RAI             | SD (Neck)      |
| 29      | 9                           | F      | 35             | cv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 30      | 12                          | M      | 22             | cv-PTC              | 4a| 1a| 1 | Surgery/RAI     | SD             |
| 31      | 14                          | F      | 3              | cv-PTC              | 1a| 1 | 0 | 0               | Excellent      |
| 32      | 14                          | M      | 10             | cv-PTC              | 3 | 0 | 0 | Surgery/RAI     | BD             |
| 33      | 17                          | F      | 11             | cv-PTC              | 3 | 0 | 0 | 0               | Excellent      |
| 34      | 17                          | F      | 35             | fv-PTC              | 2 | 0 | 0 | 0               | Excellent      |
| 35      | 11                          | F      | 20             | cv-PTC              | 3 | 1a| 0 | 0               | Excellent      |
| 36      | 14                          | F      | 30             | cv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 37      | 15                          | F      | 12             | cv-PTC              | 3 | 1a| 0 | Surgery         | Indeterminate  |
| 38      | 17                          | F      | 25             | fv-PTC              | 2 | 0 | 0 | 0               | Excellent      |
| 39      | 15                          | F      | 15             | cv-PTC              | 1b| 0 | 0 | 0               | Excellent      |
| 40      | 15                          | F      | 12             | fv-PTC              | 1b| 0 | 0 | 0               | n.a.           |
| 41      | 17                          | M      | 20             | FTC                 | 4 | 1 | 1 | RAI             | Excellent      |
| 42      | 6                           | F      | 15             | fv-PTC              | 1b| 0 | 0 | n.a.            | n.a.           |
| 43      | 13                          | M      | n.a.           | sv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 44      | 18                          | F      | 22             | cv-PTC              | 3 | 1b| 0 | 0               | Excellent      |
| 45      | 16                          | M      | 38             | cv-PTC              | 2 | 0 | 0 | 0               | Excellent      |
| 46      | 17                          | F      | 7              | tv-PTC              | 1a| 0 | 0 | n.a.            | n.a.           |
| 47      | 13                          | F      | 25             | fv-PTC              | 3 | 1a| 0 | 0               | Excellent      |
| 48      | 16                          | F      | 7              | cv-PTC              | 1a| 0 | 0 | 0               | BD             |
| 49      | 17                          | F      | 15             | cv-PTC              | 1b| 0 | 0 | 0               | Excellent      |
| 50      | 17                          | F      | 8              | cv-PTC              | 1a| 1a| 0 | Surgery         | BD             |
During the follow-up, 18/51 (35.3%) patients received further treatments, including: second surgery in 5/18 (27.8%) patients; RAI in 8/18 (44.4%), and both in 5/18 (27.8%). Positive lymph nodes and distant metastases were associated with a higher likelihood of undergoing a second treatment ($p = 0.04$ and $p = 0.002$, respectively).

Among the 51 patients with follow-up data, the final state of the disease was remission in 34/51 (66.7%) cases, persistent disease in 16/51 (31.3%); 8 with biochemically evident disease, and 8 with structurally evident disease); and for one patient who recently had surgery for a recurrence in the lateral neck the final status was classified as undetermined (restaging is ongoing).

Of the 8 patients with a final status of structurally evident disease, 6 had distant metastases at: 5/6 had stable lung disease, and 1 had progression in the lung with lesions failing to uptake RAI or 18-FDG. The other 2 patients with structurally evident disease had a local progression and are currently being reassessed for further surgery.

Disease status at the end of the follow-up correlated with disease stage: 9/41 patients in stage I (22%) had...
Molecular Characterization

In all, 26/59 patients (44%) of our series were carriers of molecular alterations.

Point Mutations

Mutually-exclusive point mutations were found in 13/50 (26%) tissue samples from which the genetic material could be amplified. No point mutations were documented in patients under 10 years old, while there was 1 among 13 (8%) 10- to 14-year-olds, and 12 among 13 (92%) 15- to 18-year-olds.

Among the samples that could be tested for the presence of BRAF mutations, the V600E mutation was found in 8/50 (16%) cases. All mutations belonged to samples of cv-PTC (7/8) and tcv-PTC (1/8). No other BRAF mutations were identified.

Among the samples that were examined for the presence of NRAS mutations, a total of 4/52 (8%) molecular events came to light. The Q61K mutation was found in 2 samples and the Q61R mutation in other 2. All mutations belonged to samples of fv-PTC. Among the samples tested for the presence of TERT promoter mutations, the C250T mutation was only found in one PTC sample (1/44).

Among the samples successfully amplified for PCR, no mutations were identified in exons 5 or 8 of PTEN, or in exons 9 or 20 of PIK3CA.

RET/PTC Rearrangements

Among the samples that could be tested for the presence of RET/PTC rearrangements, 14/57 (24.6%) cases were found to carry the translocation (Figure 2). RT-PCR data analysis confirmed the results obtained with the FISH method. Interestingly, 1 of these 14 patients had both the RET/PTC translocation and a TERT promoter mutation and was classified as a case of PDTC.

Correlation Between Clinical and Molecular Features

BRAFV600E was related to the need for a second treatment: 4/6 BRAFV600E-mutated patients (66.7%) underwent a second treatment, as opposed to only 8/35 (22.9%) in the group of patients without this mutation (p = 0.04). No information on any second treatment or follow-up were available for 2/8 BRAFV600E-mutated patients. Two of the 8 BRAFV600E-mutated patients (25%) were <15 years old, while 6/8 (75%) were ≥15–18 years old (p = 0.11) (Table 4). No significant association was found between BRAFV600E mutation status and sex, tumor size, histology, multifocality, lymph node and distant metastasis, stage, or disease outcome.

NRAS mutations were associated with histology: 4/4 samples were fv-PTC. Significant associations were found between NRAS

| Table 3 | Correlation with clinic-pathological features and outcome. |
|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|-----------------|
| Tumor size (mm) | Total (51)      | Structural disease (8/51) | Biochemical disease (8/51) | Indeterminate (1/51) | Remission (34/51) | P-value         |
| 1                | 12 (23.5%)      | 0/12 (0%)        | 3/12 (25%)       | 0/12 (0%)         | 9/12 (75%)       | 0.05           |
| 2                | 9 (17.6%)       | 0/9 (11.1%)      | 0/9 (0%)         | 0/9 (0%)          | 9/9 (100%)       |
| 3                | 24 (47.1%)      | 5/24 (20.8%)     | 3/24 (12.5%)     | 1/24 (4.2%)       | 15/24 (62.5%)    | 0.01           |
| 4                | 6 (11.8%)       | 3/6 (50%)        | 2/6 (33.3%)      | 0/6 (0%)          | 1/6 (16.7%)      |
| Lymph nodal metastasis | Structural disease (8/51) | Biochemical disease (8/51) | Indeterminate (1/51) | Remission (34/51) |
| N0–N1a          | 30 (58.8%)      | 1/30 (3.3%)      | 4/30 (13.4%)     | 1/30 (3.3%)       | 24/30 (80%)      |
| N1b             | 19 (37.3%)      | 7/19 (36.8%)     | 3/18 (15.8%)     | 0/19 (0%)         | 9/19 (47.4%)     | 0.0008          |
| Distant metastasis | Structural disease (8/51) | Biochemical disease (8/51) | Indeterminate (1/51) | Remission (34/51) |
| M1              | 10 (19.6%)      | 6/10 (60%)       | 1/10 (10%)       | 0/10 (0%)         | 3/10 (30%)       |
| TNM staging     | Structural disease (8/51) | Biochemical disease (8/51) | Indeterminate (1/51) | Remission (34/51) |
| I               | 41 (80.4%)      | 2/41 (4.9%)      | 7/41 (17.1%)     | 1/41 (2.4%)       | 31/41 (75.6%)    | 0.0003          |
| II              | 10 (19.6%)      | 6/10 (60%)       | 1/10 (10%)       | 0/10 (0%)         | 3/10 (30%)       |
| Second treatment | Structural disease (8/51) | Biochemical disease (8/51) | Indeterminate (1/51) | Remission (34/51) |
|                 | 18 (35.3%)      | 7/18 (38.9%)     | 5/18 (27.8%)     | 1/18 (5.6%)       | 5/18 (27.8%)     | 0.0001          |

Bold values are the statistically significant associations.
mutations and the absence of vascular invasion or lymph node metastases.

The only patient carrying the TERT promoter mutation (with a concomitant RET/PTC rearrangement, as mentioned above) had a locally advanced PDTC with cervical lymph node metastases (N1b) and distant metastases. At the end of the follow-up (46 months) the patient had persistent disease.

As mentioned earlier, RET/PTC rearrangements were found in 14 patients: 5/14 were RET/PTC 3 (35.7%), and 9/14 (64.8%) were RET/PTC 1. The two patients with a history of exposure to radiation carried this translocation (1 with RET/PTC 1, the other with RET/PTC 3). As for histotype, 2 patients were cases of PDTC, 10 were cv-PTC, and 2 were sv-PTC. RET/PTC rearrangement was found significantly correlated with aggressive features such as lymph node metastases (p = 0.01), and RET/PTC 3, in particular, correlated with both cervical lymph node involvement (N1b) and distant metastases (p = 0.02 and p = 0.03, respectively). Moreover, 4/5 patients with RET/PTC 3 rearrangements needed a second treatment during their follow-up (p = 0.03). As for the 8 patients with a final status of structurally evident disease, 1 was a young female with lung progression whose lesions showed no uptake of RAI or 18-FDG, and she had the RET/PTC 3 translocation.

The Clinical and Molecular Profile of Pediatric vs. Adult Thyroid Cancer

The main clinical, pathological and histopathological characteristics and outcomes of the samples of pediatric (children and adolescents) and adult patients are summarized in Table 4.

A more advanced disease at diagnosis was apparent in the younger patients: 22/29 patients <15 years old (75.8%) had T3 or T4 tumors, as opposed to 10/30 (33.3%) older pediatric patients (p < 0.01). Lymph node metastases were also more common in the former (24/28; 85.7%) than in the latter (14/27; 51.9%, p = 0.007). Of the 10 patients with distant metastases, 8 were <15 years old (p = 0.01). These two subgroups showed no significant differences in terms of primary tumor size, multifocality, histotype, need for second treatment, or disease status at the end of the follow-up, though only 56% of the younger group vs. 81% of the older one reached disease remission at the end of the follow-up.

In the adult population, 138/178 (77.5%) patients were female and 40/178 (22.5%) were male, with a male: female ratio of 1:3.5. The sex distribution was homogeneous between the pediatric and adult groups (p = 0.46). The average age of the adults was 48.6 ± 13.1 years (range 22–81 years).

The pediatric patients had larger lesions than the adults (median 23.6 vs. 19.3 mm); only 26/56 (46.4%) of the former, as opposed to 134/177 (75.7%) of the latter had a tumor <20 mm in diameter (p = 0.0004).

Lymph node metastases were found in 40/59 pediatric patients (67.8%) and 75/178 adults (42.1%) (p < 0.001). Distant metastases were more common in the pediatric patients too, involving 10/51 (19.6%) as opposed to 7/178 adults (3.9%) (p < 0.001). All distant metastases in both groups involved the lung, with the sole exception of a 17-year-old male with mediastinal disease.

The children and adolescents were significantly more likely to have a second treatment: 18/51 pediatric patients (35.3%) and 18/178 adults (10.1%) had further surgery or radiometabolic therapy (p < 0.0001). Age was found associated with final disease status: 34/51 pediatric patients (66.7%) and 157/176 adults (89.2%) were in remission (p < 0.0001). Among the adult patients, 4/176 (2.3%) died of their disease, while no pediatric patients died of cancer-related causes.

As regards the molecular profile, the rate of BRAFV600E mutations was significantly higher in adult PTC than in pediatric PTC, with 107/178 (60%) and 8/50 (16%), respectively (p < 0.0001). Although NRAS mutations emerged in 8% of the pediatric patients (4/52) as opposed to 2.2% (4/178) of the adults, this difference was not statistically significant.

FIGURE 1 | Correlation between lateral cervical (N1b) lymph node metastases and disease status in pediatric population. The Kaplan-Meier curve shows N1b significantly associated with persistent/recurrent disease at the end of the follow-up (p = 0.01).

DISCUSSION

DTC is rare in the pediatric population and its clinical, pathological and molecular characteristics differ from those of adult DTC (12). In children and adolescents, it has a more aggressive presentation at diagnosis, and a higher frequency of lymph node and distant metastases (13–18). There is also a greater risk of disease recurrence in the pediatric population. The prognosis is nonetheless excellent and the mortality rate is very low (2, 19).

Some Authors argue that the clinical and pathological differences between pediatric and adult DTC are due to a different genetic profile (20). The most common molecular event in children and adolescents (particularly after exposure to radiation) is represented by RET/PTC rearrangements, while point mutations in BRAF, RAS or TERT’ promoter genes are the main genetic drivers of tumorigenesis in adults (21). Somatic mutations (which are infrequent in pediatric age) are usually
associated with cellular dedifferentiation, genetic instability, and a reduced NIS iodine transporter expression. The paradox of pediatric DTC—with its worse presentation at diagnosis, but good outcome—could therefore be explained by a greater differentiation of the tumor cells and a consequently better response to radiometabolic treatments and TSH suppression therapy (22, 23) and also by a lower frequency of BRAF mutations (24). Regarding the potential effects of continued levothyroxine suppression, TSH contributes to the regulation of thyrocyte differentiation by modulating thyroid gene levels (25).

The clinical and histological features seen in our pediatric population are similar to those already described in other studies. The majority of our patients were female, with males and younger patients presenting with more advanced disease. Male gender coincides with more aggressive tumor features in terms of the extent of the primary tumor at diagnosis, dissemination outside the thyroid and to distant sites. On the other hand, males did not have significantly larger tumors, nor did they differ from females in terms of final outcome (26–28). Younger patients had a more aggressive tumor at diagnosis in terms of extent, lymph node involvement and distant metastases than older pediatric patients although, here again, age did not affect final outcome in our sample. In addition, the prevalence of DTC was higher among females in both subgroups, but the female/male ratio was higher for the older than for the younger pediatric patients, although the difference was not statistically significant (4:1 vs. 1.9:1). This trend is in line with findings in adult DTC, and could be influenced by hormonal aspects, and the impact of estrogens in particular. These data seem to suggest that DTC in childhood is locally more aggressive and should be considered as a distinct clinical entity from DTC in adolescence.

In the literature, the frequency of lymph node metastases in the pediatric population varies between 50 and 75%, while for distant metastases it ranges between 6 and 33% (5, 17, 19, 29, 30).
Our findings confirm this aggressiveness at diagnosis, with a prevalence of 67.7% for lymph node positivity and 19.6% for distant metastases. In our series, DTC dissemination outside the thyroid gland emerged as the main factor associated with the risk of distant metastases.

As expected, lymph node and distant metastases were significantly associated with a higher chance of undergoing further treatment during the follow-up.

The optimal management of neck disease depends on several factors, including the size and site of disease, previous treatments, disease progression rate, and the patient's age. Some patients in our sample known to have persistent disease were treated months or even years afterwards; 9 underwent second surgery, with or without further RAI, and all except one of them achieved at least a structurally evident remission.

Distant metastases in pediatric cases usually involve micronodular lung lesions with an excellent RAI uptake, characteristics that can explain why distant metastases in children are more amenable and responsive to RAI therapy than those in adults. There were 10 young patients with lung metastases in our sample and, after one or more RAI treatments, 3 obtained a complete remission, 1 had only biochemically evident disease, and 6 had persistent lung metastases, with only one patient in progression. Our multivariate analysis indicated that only the presence of distant metastases independently correlated with persistent disease (OR 13.95 % CI 2.19 to 77.03).

As emphasized by recent ATA guidelines (2), our data highlight the importance of a tailored assessment of children and adolescents with DTC. The aim should be to identify already at the initial diagnosis which pediatric patients would benefit from more aggressive surgical and radiometabolic approaches, and a closer follow-up. Clinical status at final follow-up reflects not only initial response to total thyroidectomy and RAI ablation, but also the potential effects of continued levothyroxine suppression and/or further surgery or RAI therapy over time.

Comparing our cohort of pediatric patients with an adult PTC population treated and followed up at the same center confirmed the greater aggressiveness at diagnosis of pediatric DTC. Tumor size was significantly larger in patients <18 years old, and there was a greater frequency of lymph node metastases (67.7 vs. 47.5%) and distant metastases (19.6 vs. 3.9%) than in adults. In terms of survival, however, the prognosis for pediatric DTC is excellent, and better than for the adult counterpart.

As for the molecular aspects, pediatric patients and adults clearly showed a different genetic profile. Even though almost all of the patients in our series had no history of exposure to radiation, RET/PTC rearrangements were confirmed as the most common genetic alteration in pediatric DTC (24.6%). RET/PTC rearrangements, and RET/PTC 3 in particular, were found to correlate with aggressive clinical and pathological features such as lymph node and distant metastases. Patients with the RET/PTC 3 translocation needed more second treatments (p = 0.03), suggesting [as reported in the literature (31)] that this molecular event results in a lower response to RAI and a consequent need for further treatments to achieve disease remission. In fact, the only patient with progressive lung disease in our series who became radio-refractory carries this translocation.

The BRAFV600E mutation is the most common in adult PTC (32–35), while in pediatric patients its prevalence varies between 0 and 37% (20). Our study is in line with previous reports: this mutation was found in 16% of our pediatric patients and 60.1% of our adult sample. Although the association was not statistically significant, we found that the frequency of BRAF mutations increased with age at diagnosis, even in the pediatric population, as seen in previous studies: it was 0% in patients under 10 years old, 14% among those between 11 and 14 years old, and 86% among the 15- to 18-yea-olds (21, 36). Several studies identified an association in adult DTC between BRAF mutations and more aggressive clinical, pathological and histopathological features, more recurrences and a higher mortality. This relationship has not been seen in pediatric populations (35, 36).

Our study also found no significant associations between BRAF mutations and gender, tumor size, histotype, multifocality, lymph node metastases, stage of disease or final outcome. The presence of a BRAFV600E mutation nonetheless correlated significantly with the need for a second treatment during the follow-up: among the BRAFV600E-positive patients, 4/6 (67%) received further treatment, as opposed to 11/42 (26%) in the group of patients without this mutation (p = 0.04). This finding prompts us to speculate that, as seen in adults, BRAF mutations may be associated with more aggressive clinical features and a higher risk of recurrence or persistence of disease in the pediatric population too.

Mutations in the RAS gene are rare in pediatric patients with DTC, the prevalence found in various studies to range between 0 and 7% (20). The rarity of this molecular event was confirmed in our study too: only 8% of pediatric patients had a NRAS mutation. In adults, RAS is more frequently mutated in cases of FTC (~40%), and fv-PTC (~15–20%) (37). The association between RAS gene alterations and the fv-PTC histology was confirmed in our pediatric series as well. In the adult control group, the frequency of Nras gene mutations was 3%, i.e., lower than reported in the literature—a discrepancy probably due to the fact that only 6% of PTCs in our adult population were the follicular variant. In our pediatric population, some significant associations emerged between Nras gene mutations and the clinical and pathological features of the DTC. NRAS-mutated DTC correlated with older age, no vascular invasion, and no lymph node or distant metastases. These data suggest that NRAS mutations do not have a key role in the pathogenesis of pediatric DTC, and might be a hallmark of a subset of less aggressive tumors, as seen in the albeit limited number of cases presenting this mutation in other studies (11, 18, 38).

TERT promoter, PTEN, and PIK3CA gene mutations are rare in adult DTC, while they have been found more frequently in poorly-differentiated carcinomas and anaplastic carcinoma (39–42). In pediatric age, though few studies have been conducted, the prevalence is even lower (1%) (38). This is consistent with the observation of a greater differentiation of tumor cells in pediatric DTC.

Interestingly, our one pediatric patient carrying a TERT mutation had advanced-stage disease at diagnosis, multifocality,
bilateral extra-thyroid extension, areas of poorly-differentiated carcinoma, and lymph node and distant metastases. He also had a concomitant RET/PTC translocation. The boy underwent second treatments involving both surgery and RAI, achieving a final disease status of biochemically persistent disease. This picture is consistent with previous reports in adult series: in thyroid carcinoma, TERT promoter mutations are associated with more aggressive histopathological features and a worse prognosis (40).

In conclusion, our study showed that DTC in pediatric age has different clinical, pathological and prognostic features from its counterpart in adults: it is more aggressive at diagnosis and carries a greater risk of persistence/recurrence. Within the pediatric group, special attention should be paid to male patients under 15 years old, as they are associated with a more advanced disease at diagnosis, although their final disease status does not seem to be affected by gender or age. As for the molecular profile, there are substantial differences between pediatric and adult DTC. RET/PTC translocations are the main molecular event in the pediatric population, while BRAFV600E mutations are significantly less common in pediatric DTC than in adults, and they are unassociated with the clinical and histopathological features of the disease. Finally, mutations in the TERT promoter and NRAS, PTF1, and PIK3CA genes are occasional molecular drivers of cancer in the pediatric population. Unlike the case in adults, point mutations do not have a key genetic role in children, even those not exposed to ionizing radiation.

DATA AVAILABILITY

The raw data supporting the conclusions of this manuscript will be made available by the authors, without undue reservation, to any qualified researcher.

ETHICS STATEMENT

This study was carried out in accordance with the recommendations of Padova Hospital Ethical Committee protocol No. 58403 with written informed consent from all subjects. All subjects gave written informed consent in accordance with the Declaration of Helsinki.

AUTHOR CONTRIBUTIONS

FG and FV: study concept and design, analysis and interpretation, drafting of the manuscript, and final approval of the version to be published. CM and GP: study concept and design, supervision, and final approval of the version to be published and agreement with all aspects of the work. SCe, SB, LB, SW, MR, MI, and SCA: substantial contributions to data acquisition and interpretation, critical revision of the manuscript, and final approval of the version to be published. JM: substantial contributions to data acquisition. CC: analysis and interpretation, critical revision of the manuscript, final approval of the version to be published, and agreement with all aspects of the work.

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REFERENCES

1. Merino MJ. Thyroid cancer. In: Strayer DS, Rubin E, editors. Rubin’s Pathology. 7th ed. Baltimore, MD; Philadelphia, PA: Wolters Kluwer (2015). p. 1192–6.
2. Francis GL, Waguespack SG, Bauer AL, Angelos P, Benvenga S, Cerutti JM, et al. Management guidelines for children with thyroid nodules and differentiated thyroid cancer. Thyroid. (2015) 25:716–59. doi: 10.1089/thy.2014.0460
3. Dinauer CA, Breuer C, Rivkees SA. Differentiated thyroid cancer in children: diagnosis and management. Curr Opin Oncol. (2008) 20:59–65. doi: 10.1097/CCO.0b013e3282f0220
4. Halac I, Zimmerman D. Thyroid nodules and cancers in children. Endocrinol Metab Clin North Am. (2005) 34:725–44. doi: 10.1016/j.ecl.2005.04.087
5. Grigsby PW, Gal-or A, Michalski JM, Doherty GM. Childhood and adolescent thyroid carcinoma. Cancer. (2002) 92:724–9. doi: 10.1002/cncr.10725
6. Demirdichk YE, Demirdichk EP, Reiners C, Biko J, Mine M, Saenko VA, et al. Comprehensive clinical assessment of 740 cases of surgically treated thyroid cancer in children of Belarus. Ann Surg. (2006) 243:525–32. doi: 10.1097/01.sla.0000205977.74806.0b
7. UICC. Thyroid gland. In: Sobin LH, Gospodarowicz MK, Wittekind C, editors. TNM Classification of Malignant Tumours. 7th ed. Oxford: Wiley-Blackwell Publishers (2009). p. 58–62. doi: 10.1002/9780471420194.tnmcc08.pub2
8. Pennelli G, Vianello F, Barollo S, Pezzani R, Merante Boschin I, Pelizzo MR, et al. BRAF(K601E) mutation in a patient with a follicular thyroid carcinoma. Thyroid. (2011) 21:1593. doi: 10.1089/thy.2011.0120
9. Colato C, Vicentini C, Cantara S, Pedron S, Brazzarella P, Marchetti I, et al. Break-apart interphase fluorescence in situ hybridization assay in papillary thyroid carcinoma: on the road to optimizing the cut-off level for RET/PTC rearrangements. Eur J Endocrinol. (2015) 172:571–82. doi: 10.1530/EJE-14-0930
10. Unger K, Zurnadzhy L, Walch A, Mall M, Bogdanova T, Braselmann H, et al. RET rearrangements in post-Chernobyl papillary thyroid carcinomas with a short latency analysed by interphase FISH. Br J Cancer. (2006) 94:1472–7. doi: 10.1038/sj.bjc.6603109
11. Nikita ME, Jiang W, Cheng SM, Hantash FM, McPhaul MJ, Newbury RO, et al. Mutational analysis in pediatric thyroid cancer and correlations with age, ethnicity, and clinical presentation. Thyroid. (2016) 26:227–34. doi: 10.1089/thy.2015.0401
12. Dermody S, Walls A, Harley EH Jr. Pediatric thyroid cancer: an update from the SEER database 2007-2012. Int J Pediatr Otorhinolaryngol. (2016) 89:121–6. doi: 10.1016/j.ijporl.2016.08.005
13. Chow SM, Law SC, Mendenhall WM, Au SK, Yau S, Mang O, et al. Differentiated thyroid carcinoma in childhood and adolescence - clinical course and role of radioiodine. Pediatr Blood Cancer. (2004) 42:176–83. doi: 10.1002/pbc.10410
14. Newman KD, Black T, Heller G, Azizkhan RG, Holcomb GW 3rd, Sklar C, et al. Differentiated thyroid cancer: determinants of disease progression in...
patients <21 years of age at diagnosis: a report from the Surgical Discipline Committee of the Children’s Cancer Group. Ann Surg. (1998) 227:533–41. doi: 10.1097/00000658-199804000-00014

15. Popovtzer A, Shpitzer T, Bahar G, Feinmesser R, Segal K. Thyroid cancer in children: management and outcome experience of a referral center. Otolaryngol Head Neck Surg. (2006) 135:88–4. doi: 10.1016/j.otohns.2006.04.004

16. Welch Dinauer CA, Tuttle RM, Robie DK, McClellan DR, Svec RL, Adair C, et al. Clinical features associated with metastasis and recurrence of differentiated thyroid cancer in children, adolescents and young adults. Clin Endocrinol. (1998) 45:619–28. doi: 10.1046/j.1365-2265.1998.00584.x

17. Agac Ay A, Kutun S, Cetin A. Are the characteristics of thyroid cancer different in young patients? J Pediatr Endocrinol Metab. (2014) 27:497–502. doi: 10.1515/jpem-2013-0192

18. Gertz RJ, Nikiforov Y, Rehrauer W, McDaniel L, Lloyd RV. Mutation in BRAF and other members of the MAPK pathway in papillary thyroid carcinoma in the pediatric population. Arch Pathol Lab Med. (2016) 140:134–9. doi: 10.5858/arpa.2014.0612-OA

19. Kiratli PO, Volkan-Salanci B, Gunay EC, Varan A, Akyuz C, Buyukpamukcu M. Thyroid cancer in pediatric age group: an institutional experience and review of the literature. J Pediatr Hematol Oncol. (2013) 35:93–7. doi: 10.1097/MPH.0b013e3182755d9e

20. Cordioli MI, Moraes L, Cury AN, Cerutti JM. Are we really at the dawn of cancer in children: management and outcome experience of a referral center. Ann Surg. (1998) 45:619–28. doi: 10.1046/j.1365-2265.1998.00584.x

21. Cordioli MI, Moraes L, Bastos AU, Besson P, Alves MT, Delcelo R, et al. Fusion oncogenes are the main genetic events found in sporadic papillary thyroid carcinomas from children. Thyroid. (2017) 27:182–8. doi: 10.1089/thy.2016.0387

22. Faggiano A, Coulot J, Bellon N, Talbot M, Caillou B, Ricard M, et al. Age-dependent variation of follicular size and expression of iodine transporters in human thyroid tissue. J Nucl Med. (2004) 45:232–7.

23. Patel A, Jiang S, Dogra S, Terrell R, Powers PA, Fenton C, et al. Differentiated thyroid carcinoma that express sodium-iodide symporter have a lower risk of recurrence for children and adolescents. Pediatr Res. (2002) 52:737–44. doi: 10.1203/00006450-200211000-00021

24. Durante C, Puixeddu E, Ferretti E, Morisi R, Moretti S, Bruno R, et al. BRAF mutations in papillary thyroid carcinomas inhibit genes involved in iodine metabolism. J Clin Endocrinol Metab. (2007) 92:2840–3. doi: 10.1210/jc.2006-2707

25. Bruno R, Ferretti E, Tosi E, Arturi F, Giannasio P, Mattei T, et al. Modulation of thyroid-specific gene expression in normal and nodular human thyroid tissues from adults: an in vivo effect of thyrotropin. J Clin Endocrinol Metab. (2005) 90:5692–7. doi: 10.1210/jc.2005-0800

26. Russo M, Malandrino P, Moleti M, Vermiglio F, D’Angelo A, La Rosa G, et al. Differentiated thyroid cancer in children: heterogeneity of predictive risk factors. Pediatr Blood Cancer. (2018) 65:e27226. doi: 10.1002/pbc.27226

27. Pires BP, Alves PA Jr, Bordallo MA, Bulzico DA, Lopes FP, Farias T, et al. Prognostic factors for early and long-term remission in pediatric differentiated thyroid carcinoma: the role of sex, age, clinical presentation, and the newly proposed American Thyroid Association Risk Stratification System. Thyroid. (2016) 26:1480–7. doi: 10.1089/thy.2016.0302

28. Balachandar S, La Quaglia M, Tuttle RM, Heller G, Ghosein RA, Sklar CA. Pediatric differentiated thyroid carcinoma of follicular cell origin: prognostic significance of histologic subtypes. Thyroid. (2016) 26:219–26. doi: 10.1089/thy.2015.0287

29. Wang JT, Huang R, Kuang AR. Comparison of presentation and clinical outcome between children and young adults with differentiated thyroid cancer. Asian Pac J Cancer Prev. (2014) 15:7271–5. doi: 10.7314/APJCP.2014.15.17.7721

30. Park S, Jeong JS, Ryu HR, Lee CR, Park JH, Kang SW, et al. Differentiated thyroid carcinoma of children and adolescents: 27-year experience in the Yonsei University Health System. J Korean Med Sci. (2013) 28:693–9. doi: 10.3346/jkms.2013.28.5.693

31. Prescott JD, Zeiger MA. The RET oncogene in papillary thyroid carcinoma. Cancer. (2015) 121:2137–46. doi: 10.1002/cncr.29044

32. Kimura ET, Nikiforova MN, Zhu Z, Knauf JA, Nikiforov YE, Fagnin JA. High prevalence of BRAF mutations in thyroid cancer: genetic evidence for constitutive activation of the RET/PTC-RAS-BRAF signaling pathway in papillary thyroid carcinoma. Cancer Res. (2003) 63:1454–7.

33. Pelizzo MR, Dobrinja C, Casal Ide E, Zane M, Lora O, Tonioati A, et al. The role of BRAF(V600E) mutation as poor prognostic factor for the outcome of patients with intrathyroid papillary thyroid carcinoma. Biomed Pharmacother. (2014) 68:413–7. doi: 10.1016/j.biopha.2014.03.008

34. Xing M. BRAF mutation in thyroid cancer. Endocr Relat Cancer. (2005) 12:245–62. doi: 10.1677/eRC.1.0978

35. Frasca F, Nucera C, Pellegriti G, Gangemi P, Attard M, Stella M, et al. BRAF(V600E) mutation and the biology of papillary thyroid cancer. Endocr Relat Cancer. (2008) 15:191–205. doi: 10.1677/ERC-07-0212

36. Oishi N, Kondo T, Nakazawa T, Mochizuki K, Inoue T, Kasaï K, et al. Frequent BRAF V600E and absence of TERT promoter mutations characterize sporadic pediatric papillary thyroid carcinomas in Japan. Endocr Pathol. (2017) 28:103–11. doi: 10.1007/s12022-017-9470-y

37. An JH, Song KH, Kim SK, Park KS, Yoo YR, Yang JH, et al. RAS mutations in indeterminate thyroid nodules are predictive of the follicular variant of papillary thyroid carcinoma. Clin Endocrinol. (2015) 82:760–6. doi: 10.1111/cen.12579

38. Alzahrani AS, Murugan AK, Qasem E, Alswailem M, Al-Hindi H, Shi Y. Single point mutations in pediatric differentiated thyroid cancer. Thyroid. (2017) 27:189–96. doi: 10.1089/thy.2016.0339

39. Xing M. Genetic alterations in the phosphatidylinositol-3 kinase/Akt pathway in thyroid cancer. Thyroid. (2010) 20:697–706. doi: 10.1080/10506590903502570

40. Penna GC, Vaisman F, Vaisman M, Sobrinho-Simoes M, Soares P. Molecular markers involved in tumorigenesis of thyroid carcinoma: focus on aggressive histotypes. Cytogenet Genome Res. (2016) 150:194–207. doi: 10.1159/000456576

41. Bamford S, Dawson E, Forbes S, Clements J, Pettett R, Dogan A, et al. THE COSMIC (catalogue of somatic mutations in cancer) database and website. Br J Cancer. (2004) 91:353–8. doi: 10.1038/sj/bjc.6601894

42. Wu G, Mambo E, Guo Z, Hu S, Huang X, Gollin SM, et al. Uncommon mutation, but common amplifications, of the PIK3CA gene in thyroid tumors. J Clin Endocrinol Metab. (2005) 90:4688–93. doi: 10.1210/jc.2004-2281

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