



Hürthle-cell neoplasms of the thyroid: An algorithmic approach to pathologic diagnosis in light of molecular advances

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ABSTRACT

Our understanding of neoplasia is evolving at a rapid pace in these exciting times, where recent molecular pathology advances are reinforcing and fine tuning morphological divisions and classification. Thyroid gland neoplasia in general, and Hürthle-cell neoplasms in particular, are no exception in the current era of histopathology-molecular biology paradigm. In this review paper, we discuss the rationale that led pathologists in the past to separate Hürthle-cell neoplasms into its own dedicated diagnostic category, and provide an algorithmic approach to the differential diagnosis of oncocytic lesions of the thyroid. This review will also shed light on the current WHO classification of Hürthle-cell neoplasms in light of molecular advances that justify histopathologic distinctions.

The Oncocyte

An “oncocyte” is an enlarged polygonal cell with an abundant granular eosinophilic cytoplasm, round nuclei with even chromatin pattern and prominent nucleoli. The term “oncocyte” was first defined by Hamprel in 1931, and a tumor arising from these cells in a parotid gland was termed “oncocytoma” in 1932 by Jaffe.^{1, 2} Electron microscopy studies have shown that this peculiar cytoplasmic eosinophilia is due to the accumulation of numerous enlarged and globular mitochondria.^{3–5} It has been shown that oncocytic change represents a cellular response due to mitochondrial adaptive homeostasis. Cells from different organs undergo oncocytic change including breast, salivary gland, thyroid, parathyroid, pituitary and kidney. The oncocytes are named differently in each organ. For example, they are referred to as apocrine cells in the breast and as Hürthle cells in the thyroid gland. The latter being a misnomer, since Karl Hürthle described parafollicular cells (now known as C-cells) in the thyroid gland⁶; however, it was actually Askanazy who first described oncocytic change in follicular cells of the human thyroid gland.⁷ Interestingly, varying degrees of cytoplasmic eosinophilia can occur in thyroid follicular cells. It will not be farfetched to assume that this most likely represents an early adaptive cellular change eventually resulting in classic Hürthle cell morphology.

Hürthle-cell neoplasms of the thyroid gland: general pathologic concepts

The thyroid gland can show oncocytes either as few scattered cells lining follicles, cellular aggregates, or well-defined unencapsulated, partially or completely encapsulated nodules. The cellular aggregates are commonly encountered in chronic lymphocytic thyroiditis as focal proliferations confined to a lobe or scattered throughout the entire gland. Similarly, oncocytic change or cellular proliferations can also be seen in euthyroid-multinodular and diffuse or nodular toxic goiter.

Hürthle-cell neoplasms are rare compared to other follicular cell derived thyroid tumors. In the 2004 edition of WHO Classification of the Endocrine Organs,⁸ the oncocytic tumors of the thyroid gland were classified under the umbrella term of “Oncocytic Follicular Neoplasms”; however, this was changed to “Hürthle (oncocytic) cell tumors” in the current version published in 2017.⁹ As defined by WHO, Hürthle-cell neoplasms are composed of 75% or greater Hürthle cells.⁹

In the early literature, some authors considered all Hürthle-cell neoplasms of the thyroid gland as malignant, because some of the lesions classified as benign Hürthle-cell neoplasms behaved in a malignant fashion.^{10–12} However, these findings were later disputed by reports based on strict application of diagnostic criteria to differentiate between benign and malignant Hürthle-cell neoplasms and meticulous clinical follow-up.^{13–17}

In this review, we describe an algorithmic approach (Figs. 1 and 2)

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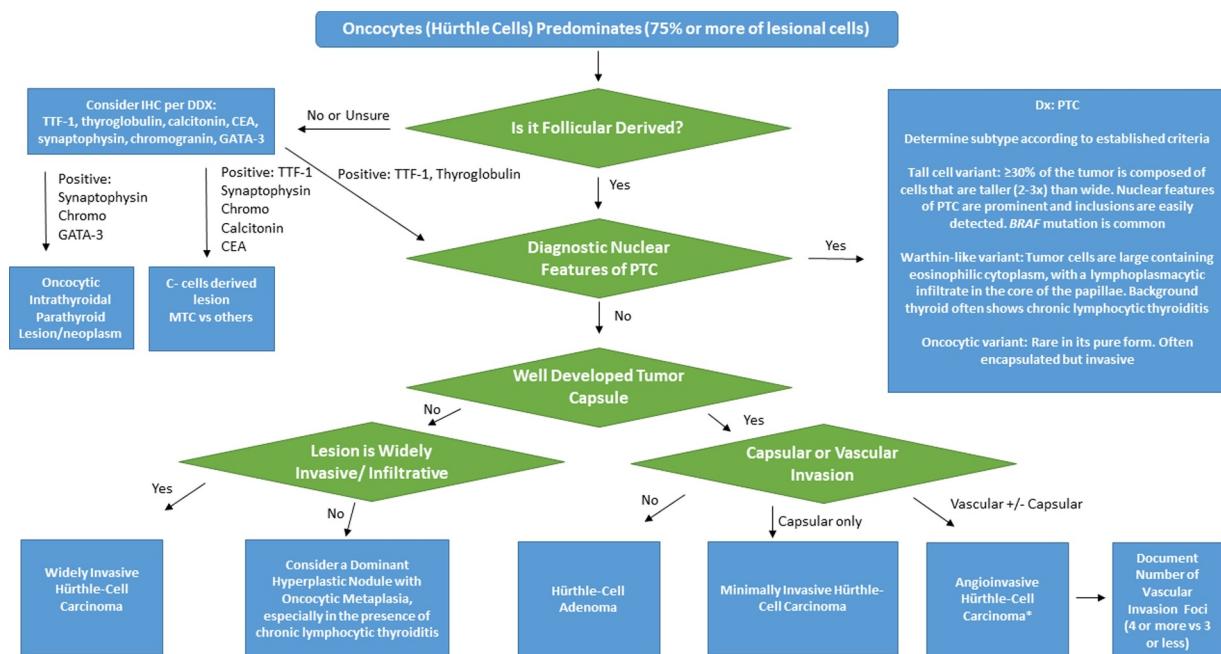


Fig. 1. An algorithmic approach to oncocytic lesions in surgical pathology.

* Some authors combine focal angioinvasion (less than 4 foci) with minimally invasive follicular carcinoma, while they consider extensive angioinvasion with widely invasive follicular carcinoma. Our personal approach is similar to the 2017 WHO with separating angioinvasive follicular or Hürthle-cell carcinoma into its own category.

Abbreviations: Chromo: chromogranin. Dx: Diagnosis. DDX: Differential diagnosis. IHC: Immunohistochemistry. MTC: Medullary thyroid carcinoma. PTC: Papillary thyroid carcinoma.

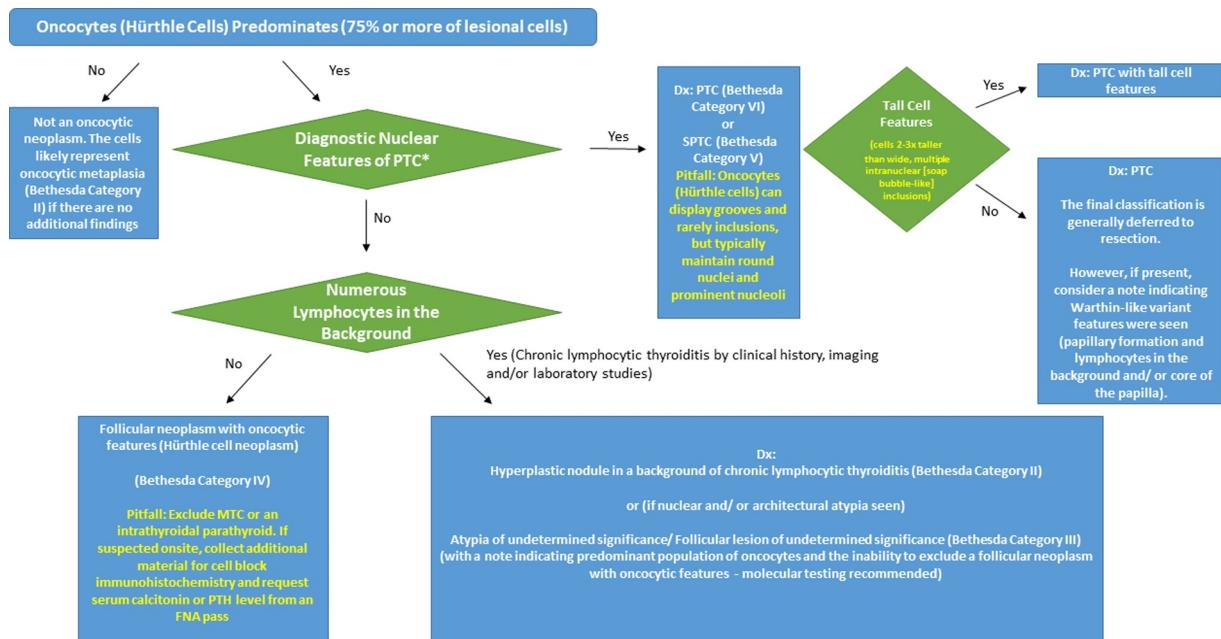


Fig. 2. An algorithmic approach to oncocytic lesions in cytopathology.

* Nuclear crowding, elongation, chromatin clearing, nuclear grooves, and inclusions.

Abbreviations: Dx: Diagnosis. MTC: Medullary thyroid carcinoma. PTC: Papillary thyroid carcinoma. PTH: Parathyroid hormone. SPTC: Suspicious for papillary thyroid carcinoma.

to the pathologic diagnosis of Hürthle-cell neoplasms of the thyroid gland, in light of the recent molecular advances.

The following should be considered in diagnosing a Hürthle-cell neoplasm of the thyroid gland.

Establishing the origin of a Hürthle/oncocytic thyroid lesion

Answering this basic question is the first step to avoid pitfalls while evaluating Hürthle /oncocytic lesions of the thyroid, especially in limited tissue samples procured by fine-needle aspiration (FNA) and core biopsies. Oncocytic change can be encountered in medullary thyroid carcinoma and intrathyroidal parathyroid lesions.

Table 1

The differential diagnosis of oncocytic lesion in the thyroid.

Diagnosis	Morphological Clues and Immunohistochemistry
Medullary thyroid carcinoma	Eccentric nucleus with salt and pepper chromatin, tumor cells spindling, presence of amyloid, positive for calcitonin, CEA, TTF-1, synaptophysin, and chromogranin
Intrathyroidal parathyroid with oncocytic features	Trabecular or cord-like arrangement of cells, centrally placed small nucleus without a prominent nucleoli, presence of other cell types (chief cells or clear cells), positive for GATA-3 and PTH
Hürthle cell adenoma	Follicular growth, colloid, centrally placed large hyperchromatic nuclei with prominent nucleoli, encapsulated with no capsular or vascular invasion, positive for TTF-1, thyroglobulin, Pax-8
Hürthle cell carcinoma	Similar to Hürthle cell adenoma but with capsular or vascular invasion, widely infiltrative variants may not be encapsulated
Oncocytic variant of PTC (including tall cell and Warthin-like variants)	Nuclear features of PTC regardless of growth pattern, positive for TTF-1, Pax-8, thyroglobulin

Morphological clues that help establish follicular cell origin mainly in FNA samples is the presence of background colloid, oncocytic follicular cells partially rimmed by blebs of colloid, AKA “flame cells”, and absence of clinical, and radiological features that suggest non-follicular lesions. Clinical history of hypercalcemia (although some oncocytic parathyroid adenomas have minimal increase in serum parathyroid hormone levels) or ultrasound findings may suggest an intrathyroidal parathyroid.^{18–20} In this scenario, performing PTH levels on FNA needle rinse or immunohistochemistry can establish the correct diagnosis. The presence of amyloid or spindled tumor cells may serve as clues to the possibility of an oncocytic variant of medullary thyroid carcinoma.^{21–26} Fortunately, this question is easily resolved by a limited immunohistochemistry panel in ambiguous cases (Table 1).

Pathologic features

Histopathologic features

Hürthle-cell neoplasms of the thyroid are rare. The differential diagnosis includes papillary thyroid carcinoma variants demonstrating oncocytic cytoplasm, oncocytic variant of medullary thyroid carcinoma and oncocytic parathyroid lesions (see Table 1 and Figs. 3 and 4).

Hürthle-cell carcinomas (HCCA) are more common in men and older patients. Most HCCAs are large tumors and demonstrate aggressive clinical behavior compared to non-Hürthle follicular carcinomas.^{27–29}

On gross pathologic examination, a majority of Hürthle-cell neoplasms are partially or completely encapsulated solitary nodules, and can show a distinct mahogany brown cut surface and a central scar similar to oncocytic lesions of other organs. Most HCCAs measure greater than 3.0 cm; however, smaller tumors have been reported.³⁰ Intratumoral hemorrhage and partial or complete tumor infarction due to a preoperative FNA are commonly encountered in these tumors.³¹

On light microscopy, Hürthle-cell neoplasms can display varying growth patterns, the most common being solid and trabecular. However, Hürthle-cell neoplasms can also demonstrate micro and macrofollicular, pseudopapillary and rarely true papillary growth patterns. Concentric calcifications of the luminal colloid resembling psammoma bodies can be seen in Hürthle-cell neoplasms.^{16,17,28,32,33} Furthermore, nuclear grooves and rarely poorly formed intranuclear pseudo-inclusions can also be seen in Hürthle cells, especially in cytology preparations.³⁴ These features may pose diagnostic challenges in differentiating a Hürthle-cell neoplasm from papillary thyroid carcinoma (see also the differential diagnosis section). Clear-cell change is

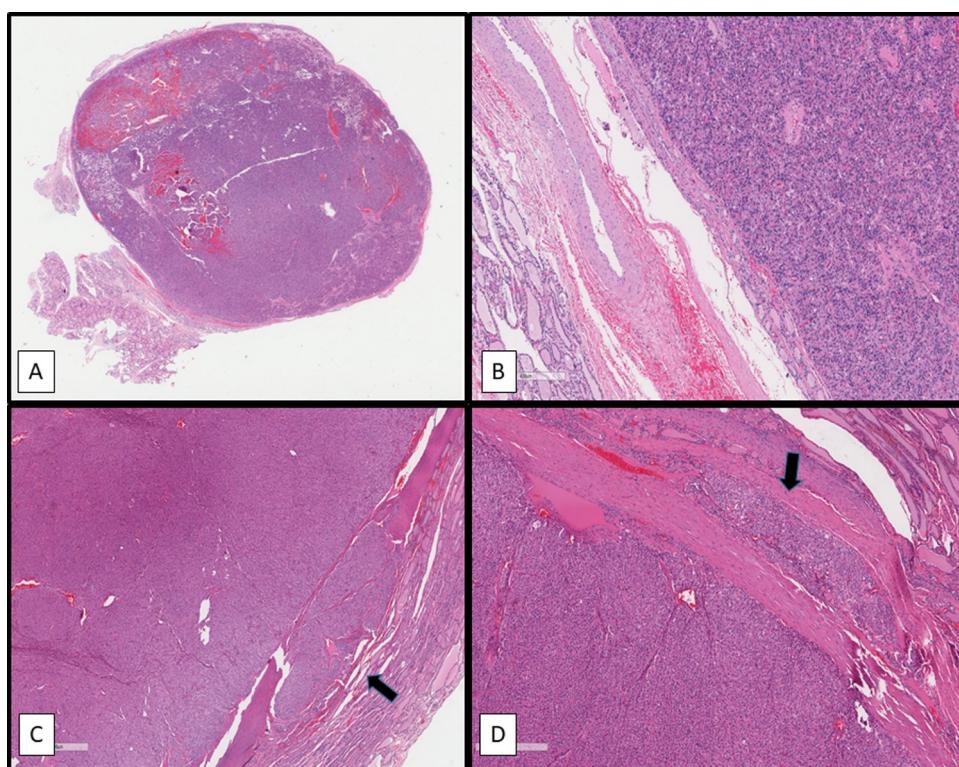


Fig. 3. Hürthle-cell adenoma (A and B) and carcinoma (C and D). Black arrows indicate foci of capsular (C) and vascular (D) invasion.

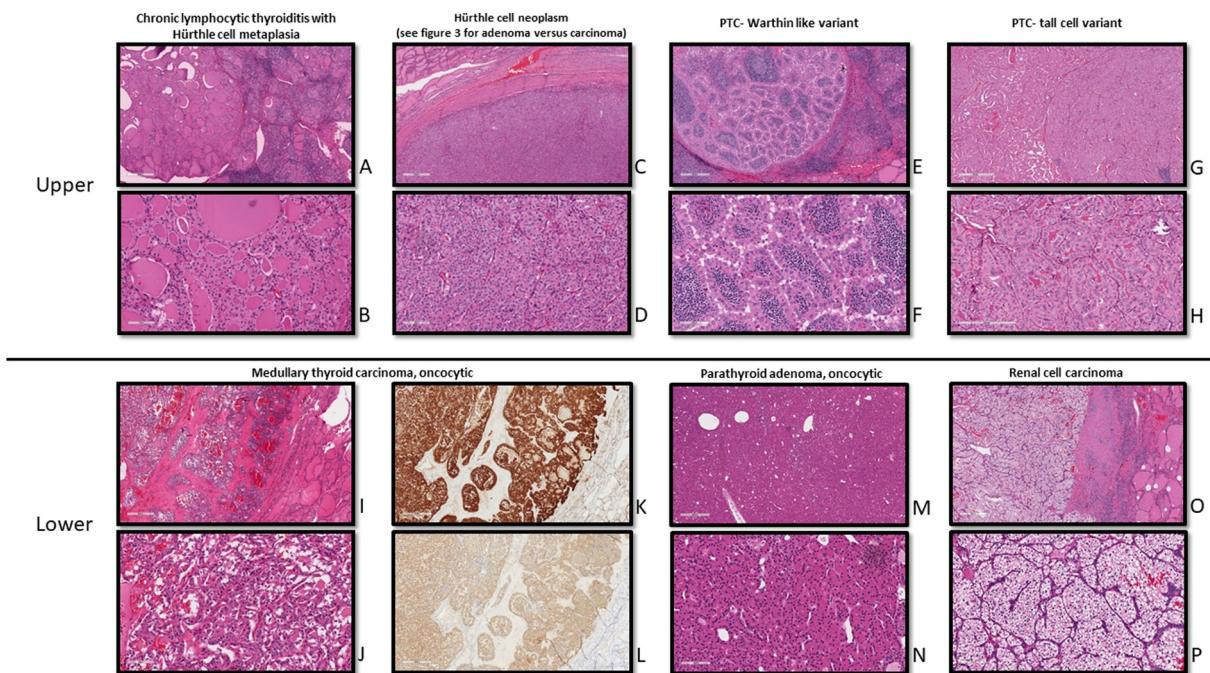


Fig. 4. Oncocytic lesions in the thyroid gland. Upper panel: follicular derived lesions. Lower panel: Non-follicular derived lesions. Chronic lymphocytic thyroiditis with Hürthle-cell metaplasia, 400x (A) and 2000x (B). Hürthle-cell neoplasm, 400x (C) and 2000x (D). Papillary thyroid carcinoma, Warthin-Like Variant, 400x (E) and 2000x (F). Papillary thyroid carcinoma, tall cell variant, 400x (G) and 2000x (H). Medullary thyroid carcinoma, 400x (I), 2000x (J), CEA (K) and calcitonin (L). Oncocytic parathyroid, 400x (M) and 2000x (N). Metastatic clear renal cell carcinoma, 400x (O) and 2000x (P).

Abbreviations: PTC: Papillary thyroid carcinoma.

not uncommon in Hürthle-cell neoplasms; however, metastasis from a clear cell carcinoma should be entertained if the tumor mostly consists of clear cells (Fig. 4O and 4P).³³

The presence or absence of invasive characteristics in an encapsulated Hürthle-cell neoplasm (i.e. invasion into the tumor capsule and/or capsular vessels [angioinvasion]) distinguishes Hürthle-cell adenoma from carcinoma (Fig. 3). Tumor size, nuclear atypia, cellular pleomorphism and mitoses in a Hürthle-cell neoplasm are not independent predictors of malignancy. Similar to follicular thyroid carcinoma, HCCA is classified to minimally invasive, encapsulated angioinvasive and widely invasive.⁹ The minimally invasive terminology is restricted for tumors showing only tumor capsule invasion, whereas the encapsulated angioinvasive tumors are further subdivided into tumors showing limited (less than four vessels) or extensive (equal to or greater than four vessels) vascular invasion.^{9, 35}

The diagnosis of HCCA as well as non-Hürthle cell follicular carcinoma based solely on tumor capsule invasion is controversial, as none of these cases (which have been adequately sampled) have shown malignant behavior on clinical follow-up. This controversy is further inflamed by what constitutes capsular invasion; whether it is invasion into the tumor capsule only or one which invades into and through the tumor capsule.³⁶ In our practice, we do classify adequately sampled (sampling of the entire capsule and tumor interface) Hürthle-cell neoplasms as minimally invasive showing either invasion into or through the entire width of the tumor capsule.^{37–41}

All pathologists agree that a Hürthle-cell neoplasm associated with invasion either into the vessels within the tumor capsule or beyond is a HCCA. It has been shown that vascular invasion is an important risk factor for tumor recurrence and distant metastasis. Specifically, higher risk of tumor recurrence is noted in cases involving four or more vessels.³⁵ The criteria for vascular invasion include an endothelial lined tumor thrombus attached to a vessel wall. However, the pathologic criteria for vascular invasion should be applied strictly to avoid over-diagnosis. Because Hürthle-cell neoplasms are richly vascularized and stroma-poor, manipulation during surgery and gross pathologic

examination can lead to displacement of the tumor cells within capsular vessels mimicking angioinvasion.³⁶

It is well-known that post FNA changes (AKA Worrisome Histologic Alteration Following Fine-Needle Aspiration – WHAFFT) are more common in Hürthle-cell neoplasms compared to other thyroid lesions^{31,42}. These can range from reactive nuclear atypia, inflammation and vascular proliferation around the needle track to necrosis and infarction. In rare instances, it may be difficult to differentiate between Hürthle-cell adenoma and carcinoma in cases with near total or complete infarction. It has been recommended to diagnose these as “Hürthle-cell neoplasm with extensive infarction without definite evidence of invasive features” with a comment explaining the difficulties in definite pathologic characterization.²⁸

The widely invasive HCCA does not pose any diagnostic challenges as it is associated with extensive invasion into the tumor capsule, involvement of surrounding thyroid and angioinvasion.⁹

The dedifferentiation of HCCA to poorly differentiated and anaplastic carcinoma has been reported.^{43–48} The poorly differentiated oncocytic carcinomas usually present as large tumors with insular/trabecular/solid growth patterns, marked nuclear pleomorphism, mitoses and foci of confluent necrosis. Some cases may also show a small cell component. These tumors show readily identifiable foci of capsular and vascular invasion.^{28,33,49}

Cytopathologic features

In FNA specimens one can readily recognize the spectrum of oncocytic change. At one end of the spectrum are the follicular cells with small to medium round nuclei without prominent nucleoli and without sharp cytoplasmic outlines, and on the other are the so called Hürthle cells as classic polygonal cells with abundant and granular cytoplasm, sharp cytoplasmic borders and housing an enlarged round nucleus with prominent nucleolus.^{34,50} On Diff-Quik (Romanowsky type) stain, the cytoplasm appears purple and the cytoplasmic granules are easier to identify (Fig. 5A). On Papanicolaou-stained smears, the nuclear chromatin is coarse and the prominent nucleoli is striking (Fig. 5B).

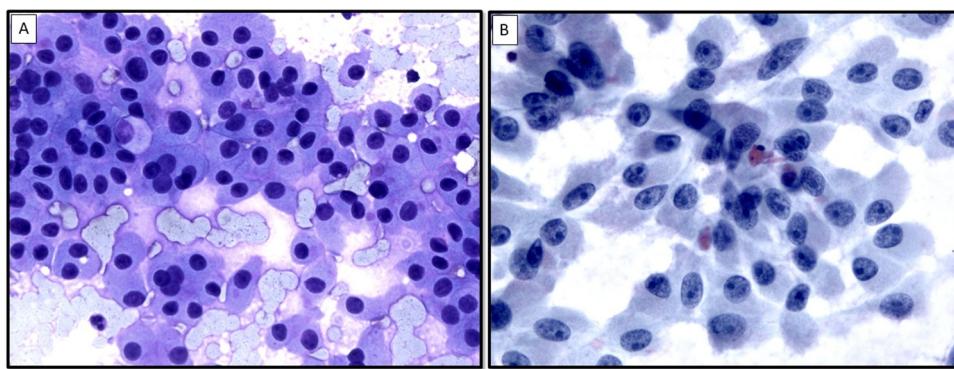


Fig. 5. Hürthle cells in cytology Diff-Quik (A) and Pap stains (B).

As aforementioned, oncocytic follicular cells can be seen in aspirates of adenomatoid nodules in the setting of euthyroid goiter, chronic lymphocytic thyroiditis and toxic nodular goiter. An aspirate containing >75% of oncocytic follicular cells or Hürthle cells should be classified as “suspicious for or consistent with follicular neoplasm with oncocytic features (Bethesda Category IV)”. The majority of FNA specimens from Hürthle cell neoplasms (adenoma or carcinoma) are hypercellular and contain Hürthle cells in cohesive groups, monolayer sheets and three dimensional structures. Follicle formation can also be seen in aspirates of Hürthle-cell neoplasms.³⁴ In most cases, background colloid is either scant or absent; however, watery colloid has been reported in aspirates of HCCA, especially those containing macrofollicles.⁵¹

Random nuclear atypia in the form of nuclear enlargement, multi-nucleation, cellular pleomorphism and prominent nucleoli can be encountered in aspirates of Hürthle-cell neoplasms.⁵² Presence of intracytoplasmic holes/lumina and transgressing vessels among cell groups have also been reported in FNA specimens of Hürthle-cell neoplasms compared to non-neoplastic proliferations.^{53,54}

Differential diagnosis

As noted above, the differential diagnosis of an oncocytic thyroid neoplasm includes papillary thyroid carcinoma variants with oncocytic features, oncocytic variant of medullary thyroid carcinoma, oncocytic lesions of the parathyroid gland and metastatic tumors.

Papillary thyroid carcinoma (PTC) is unique among other malignancies because its diagnosis is based on well characterized nuclear features. The importance of these nuclear features was identified early in the surgical pathology literature.^{55,56} They include nuclear enlargement and crowding, chromatin clearing, nuclear membrane irregularities, nuclear grooves and pseudoinclusions. The three main papillary thyroid carcinoma variants with abundant eosinophilic cytoplasm are: Tall cell variant (TCV), Warthin-like variant, and oncocytic variant. PTC-TCV tumor cells are at least 2-3 times longer than wide, and therefore appear columnar. The nuclear features in this variant are well-developed and pose no diagnostic problems. The Warthin-like variant shows a prominent lymphocytic infiltrate involving the tumor papillary fronds, resembling Warthin tumor in salivary glands. The oncocytic variant is the least common, especially in its pure form, and most controversial. It shows complex branching papillae and contains greater than 75% oncocytic tumor cells with nuclear features of PTC. A recent study compared the oncocytic variant of PTC with classic PTC.⁵⁷ The oncocytic variant of PTC occurs in older patients and the tumor size is larger. However, recurrence and disease specific survival did not differ between the two, which is supported by others as well.^{58,59}

Oncocytic Variant of Medullary Thyroid Carcinoma is rare; and shows nested or trabecular growth pattern with interspersed intratumoral hyaline stroma. The tumor cells demonstrate characteristic neuroendocrine nuclear chromatin. The diagnosis of medullary carcinoma can be confirmed by immunoreactivity for TTF-1, calcitonin,

synaptophysin, chromogranin, and CEA, but not for thyroglobulin or pax-8 (Fig. 1).^{21–26}

Parathyroid glands can show few poorly circumscribed collections of oncocytic cells. Oncocytic parathyroid adenomas are rare (Fig. 4M and 4N). It may be difficult to differentiate between oncocytic parathyroid neoplasms and Hürthle-cell neoplasms, especially in FNA samples. Ancillary studies such as PTH analysis of the FNA aspirate and/or immunostains are helpful in arriving at a correct diagnosis. A limited number of oncocytic parathyroid carcinomas have been reported in the literature.^{18,19,60,61}

Clinical behavior

There is much debate in the literature regarding the clinical behavior, prognosis and optimal treatment of HCCA.^{27, 62, 63} Some studies have reported that the biologic behavior of HCCA is considerably different from non-Hürthle follicular carcinoma; the former with more frequent lymph nodes and distant metastasis. Others conclude that the prognosis of HCCA is similar to non-Hürthle counterparts when disease stage is taken into account.^{14–16,27,45,63,64} While HCCA may lose their ability to uptake radioactive iodine, total thyroidectomy followed by radioactive iodine still remain the standard of care for most patients.^{29,63,65}

Molecular characterization of Hürthle-cell carcinoma

The morphological distinction of Hürthle-cell carcinoma from the non-Hürthle follicular carcinoma is supported by the unique molecular profiles of both malignant neoplasms.^{66–69} Follicular carcinoma frequently shows RAS (K-RAS, H-RAS, N-RAS) mutations. However, these mutations are rather infrequent (10–15%) in HCCA. BRAF V600E mutation is almost non-existent in HCCA, in contrast to PTC. The molecular profile of HCCA has been studied extensively.^{60,68–83} The molecular changes of HCCA can be divided into the following (Table 2):

Abnormal number of chromosomes

Corver et al were among the first to describe the near haploid genome unique to HCCA.⁷¹ None of the papillary and non-Hürthle follicular thyroid carcinomas studied displayed this unique karyotype. Interestingly, chromosome 7 maintained a heterozygous status in the analyzed cohort of neoplasms. These unique findings were also supported by subsequent studies.^{60,75, 78} More recently, two large studies by Ganly et al and Gopal et al evaluated 56 and 70 cases of HCCA, respectively. Both studies identified the unique near-haploid genome of HCCA. Chromosome 7 also gained additional copies in many HCCAs. These additional copies were due to genome wide duplication that lead to loss of heterozygosity in the haploid chromosomes. The BRAF gene is housed on chromosome 7, and the additional copies provide alternative methods of gene amplification/ overexpression. Other chromosomes that frequently maintained heterozygous status with or without

Table 2

The molecular profile of Hürthle cell carcinoma (Follicular Thyroid Carcinoma, Oncocytic Variant).

Study (No. of cases)	Method	Findings
Tuin K et 2019 ³⁵	DNA variant analysis, Sanger sequencing	Others <i>TERT</i> (24%), <i>RAS</i> (3%), <i>ETV6-NTRK3</i> (3%)
Gopal RK 2018 ⁷⁰	Whole exome sequencing	Chromosomal aberrations Near haploid genome (54%) Genome wide duplication (20%) Loss of heterozygosity (LOH) with heterozygous chromosomes 5, 7, 12, and 20
	FISH analysis	Mitochondrial DNA Mutations in genes coding complex I (60%) <i>ND5</i> (20%), <i>ND2</i> (15%), <i>ND4</i> (15%), <i>ND1</i> (13%), <i>ND3</i> (3%), <i>ND4L</i> (3%), <i>ND6</i> (3%) Others: <i>CO1</i> (5%), <i>CO2</i> (5%), <i>CYB</i> (3%) MtDNA common deletion not detected
	Imaging flow cytometry	Others <i>TERT</i> (32%), <i>DAXX</i> (17%), <i>NFI</i> (17%), <i>TP53</i> (12%), <i>ATM</i> (10%), <i>NRAS</i> (7%), <i>CDKN1A</i> (7%), <i>KMT2D</i> (7%), <i>ARHGAP35</i> (5%), <i>BRAF</i> (5%), <i>TSHR</i> (2%), <i>EIF1AX</i> (2%), <i>MEN1</i> (2%), <i>PTEN</i> (5%), <i>MTOR</i> (2%), <i>MSH6</i> (2%), <i>NFE2L2</i> (2%), <i>HERC2</i> (2%), <i>ARHGEF3</i> (2%), <i>FBXO31</i> (2%), <i>CSF2RA</i> (2%), <i>DMD</i> (2%), <i>KDM6A</i> (2%), <i>NFE2L2</i> (2%), <i>CACNA1C</i> (2%), <i>KRAS</i> (2%) <i>PAX8-PPARγ</i> fusion (2%)
Ganly I 2018 ⁵⁶	Whole exome sequencing	Chromosomal aberrations Near haploid genome with or without genome wide duplication and loss of heterozygosity Chromosome 5, 7, 12 duplication
	RNA sequencing	Mitochondrial DNA (71%) <i>ND1</i> , <i>ND2</i> , <i>ND4</i> , <i>ND4L</i> , <i>ND5</i> , <i>CYB</i> , <i>TL1</i> , <i>CO1</i> , <i>CO2</i> , <i>CO3</i> , and <i>ATP6</i> Almost all mtDNA encoding genes were involved except <i>ND6</i> and <i>ATP8</i>
	FISH analysis	Others Overall, 4,293 somatic mutations. Commonly mutated genes include: <i>TERT</i> (22%), <i>MADCAM1</i> (20%), <i>EIF1AX</i> (11%), <i>ERBB2</i> (11%), <i>UBXN11</i> (9%), <i>NRAS</i> (9%), <i>HLA-A</i> (9%), <i>NFI</i> (9%), <i>ALKBHZ</i> (7%), <i>FAT1</i> (7%), <i>SIRPA</i> (7%), <i>NBPFI</i> (7%), <i>TP53</i> (7%), <i>ATM</i> (5%), <i>PTEN</i> (4%), <i>RET</i> (4%), <i>MET</i> (4%), <i>KRAS</i> (4%), <i>HRAS</i> (2%) <i>PIK3CA</i> (2%) Other commonly mutated genes include: <i>OR4L1</i> , <i>ATXN1</i> , <i>FAM171B</i> , <i>POMZP3</i> , <i>HRCT1</i> , <i>DLX6</i> , and <i>FRG2B</i> No <i>BRAF</i> mutations Recurrent in-frame coding rearrangements: <i>CHCHD10_VPREB3</i> , <i>HEPHL1_PANX1</i> , <i>TMEM233_PRKAB1</i> , <i>ACSS1_APMAP</i> , <i>RSPH6A_DMWD</i> , <i>DUOXA1_DUOX2</i> , <i>OSGIN1_NECAZ2</i> , <i>BCAP29_SLC26A4</i> , <i>TFG_GPR128</i> Gene expression profiling shows significant difference between widely invasive Hürthle cell carcinoma and minimally invasive Hürthle cell carcinoma
Chindris et al 2015 ⁶¹	Targeted <i>TERT</i> sequencing	Others <i>TERT</i> (13%)
Evangelisti C et al 2015 ¹⁴	Targeted <i>TP53</i> , <i>BRAF</i> , <i>RAS</i> , and nuclear genes coding mitochondrial complex I subunits sequencing RT-PCR for <i>RET-PTC3</i> <i>FISH</i> for <i>PAX8-PPARγ</i>	Mitochondrial DNA <i>ND1</i> (21%), <i>ND2</i> (7%), <i>ND4</i> (14%), <i>ND5</i> (7%), <i>CYB</i> (7%) Nuclear genes coding mitochondrial complex I subunits <i>NDUFA12</i> , <i>NDUFB6</i> Others <i>TP53</i> (14%), <i>NRAS</i> (7%), <i>PAX8-PPARγ</i> (7%), No <i>RET-PTC</i>
Kasaian K 2015 ⁷⁴	Whole genome sequencing and sanger sequencing of <i>MEN1</i> in two cases	Chromosomal aberrations Loss of chromosomes 1, 2, 3, 4, 6, 8, 9, 11, 14, 15, 16, 21, and X (ie: near haploid genome) Extra copies in remaining chromosomes
	Targeted <i>MEN1</i> sequencing	Others <i>MEN1</i> (4.2%), <i>EWSR1</i> , <i>MSH2</i> , <i>BRCA1</i>
Qasem E et al 2015 ³	Targeted <i>TERT</i> and <i>BRAF</i> sequencing	Others <i>TERT</i> (30%) No <i>BRAF V600E</i> mutation
Wei S 2015 ⁹	Massive parallel sequencing of 47 genes	Others <i>TP53</i> (44%), <i>PTEN</i> (22%)
Corver WE 2014 ⁷	Genome wide SNP array	Chromosomal aberrations Near haploid genome (43%) with or without genome wide duplication Heterozygous chromosomes 5, 7, and 12
	Flow cytometry	Mitochondrial DNA <i>ND1</i> (14%), <i>ND3</i> (14%), <i>CO1</i> (14%)
	MtDNA sequencing	Others No <i>BRAF</i> , <i>HRAS</i> , <i>NRAS</i> , or <i>IDH1</i> or <i>IDH2</i> mutations
Ganly I 2013 ¹⁹	Mass spectrometry-based genotyping assays of common genes RT-PCR of common fusions comparative genomic hybridization Gene expression profiling	Chromosomal aberrations Chromosomes 5,7,10,12,17 and 20 gains Chromosomal losses: 4p16, 5p15-5q35, 6p25, 7p15-7q36, 8p21-23, 10p13, 12p13-q24, 16q23, 17p13-q25 Chromosomal gains: 4q24, 6p23, 6q26, 7p15, 9q33, 13q14, 22 Others <i>RAS</i> (16%) No <i>RET</i> , <i>BRAF</i> , <i>PIK3CA</i> mutations No <i>RET</i> , <i>PPARG</i> rearrangement No <i>PIK3CA</i> amplifications Gene expression profiling shows significant difference between widely invasive Hürthle cell carcinoma and minimally invasive Hürthle cell carcinoma or Hürthle cell adenoma

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Table 2 (continued)

Study (No. of cases)	Method	Findings
Landa I 2013 ²⁵	Targeted <i>TERT</i> sequencing	Others <i>TERT</i> (16%)
Corver WE 2012 ¹⁰	Flow cytometry FISH SNP array and LAIR analysis	Chromosomal aberrations Near-haploid with or without genome wide duplication and subsequent loss of heterozygosity Heterozygous chromosome 7 Others <i>PIK3CA</i> (c.86276A.G, p.H1047R) (10%)
De Vries MM 2012 ¹⁷	FISH analysis for <i>RET-PTC</i> and <i>PAX8-PPARγ</i> Targeted <i>BRAF</i> , <i>HRAS</i> , and <i>NRAS</i> sequencing	Others <i>RET-PTC</i> (38%) <i>PAX8-PPARγ</i> (27%) <i>NRAS</i> (6%) No <i>BRAF</i> or <i>HRAS</i> mutations Mitochondrial DNA <i>ND1</i> , <i>ND2</i> , <i>ND4</i> , <i>ND5</i> , <i>ATP6</i>
Gasparre G 2007 ²²	mtDNA sequencing	Mitochondrial DNA <i>GRIM-19</i> (9%) Others No <i>BRAF</i> mutation or <i>RET/PTC</i> rearrangement
Máximo V 2005 ¹¹	Targeted GRIM-19 Sequencing LOH analysis in 19p13.2 region	Mitochondrial DNA <i>GRIM-19</i> (9%) Others No <i>BRAF</i> mutation or <i>RET/PTC</i> rearrangement
Máximo V 2002 ¹³	mtDNA sequencing	Mitochondrial DNA <i>ND1</i> , <i>ND3</i> , <i>ND5</i> , <i>ND6</i> , <i>ATP6</i> , <i>CYB</i> , <i>CO1</i> , <i>CO2</i> , <i>CO3</i> MtDNA common deletion (100% cases) MtDNA D-loop somatic point mutations (53.9%)

additional copies were chromosomes 5 and 12.

Corver et al. have recently showed that this near haploid genome may be due to reactive oxygen species secondary to mitochondrial DNA mutations involving complexes of the respiratory chain.⁸⁴ The mitochondrial DNA mutations lead to glycolysis dependent glucose metabolism with accumulation of reactive oxygen species. Interestingly, the near haploid genome and mitochondrial DNA mutations were shown to be early events (truncal events) in the evolution of HCCA, shared between the primary tumor, recurrence, and metastasis.⁷⁷

Mutations targeting the mitochondrial DNA

Most mitochondrial DNA mutations involve genes encoding Complex I of the mitochondrial respiratory chain, which is composed of seven hydrophobic complex I subunits encoded by mitochondrial DNA genes (*ND1-ND6* and *ND4L*). All were shown to be mutated in HCCA, albeit with varying frequencies.^{60,68,69,74,76,77} Gopal et al have shown that *ND5* is the most frequently involved, followed by *ND2*, *ND4*, and *ND1*. Other mitochondrial genes were mutated as well, including *CO1*, *CO2*, *CO3*, *CYB*, and *ATP6* being consistently mutated across different studies.^{60,68,74,76,77} Mitochondrial DNA mutations are not unique to Hürthle-cell neoplasms, as studies have shown that papillary thyroid carcinoma, non-Hürthle follicular carcinoma, and other malignancies can show these mutations.^{60,68,76} The common deletion in mitochondrial DNA, which deletes between nucleotides 8,470 and 13,447 of the human mtDNA, was initially thought to be more prevalent in Hürthle-cell neoplasms⁶⁸; however, recent studies have not corroborated these findings.^{60,77}

Mutations in specific nuclear DNA genes

Among the different genes mutated in HCCA, the *TERT* promoter is frequently mutated (range 13–32%).^{70,74,77,79–81} It should be noted that *TERT* promoter mutation is also described in other aggressive thyroid malignancies and is not unique to HCCA. Interestingly, minimally invasive HCCAs are less likely to show *TERT* promoter mutation compared to widely invasive/ angioinvasive HCCAs.^{70,74} Some authors classify Hürthle-cell carcinomas into minimally invasive (including those with focal (< 4 foci) of angioinvasion) and widely invasive (including those with ≥ 4 foci of angioinvasion). In that classification, minimally invasive Hürthle-cell carcinomas with *TERT* mutations also show focal angioinvasion.^{70,74} It is our practice to separate angioinvasive Hürthle-cell carcinomas into its own category in line with the WHO 2017 recommendation (Fig. 1). HCCA can also display mutations in

other genes involved in telomere length (*DAXX* and *ATRX*).⁷⁷

Ganly et al showed that genes mutated in HCCA were clustered in signal transduction and tumor suppressor pathways.⁷⁴ The MAP kinase and mTOR pathways were involved in 55% of HCCA. These genes included *EIF1AX*, *NFI*, *PTEN*, *TSC1/2*, *EIF*, *RAS*, *EGFR*, *ERBB2*, *PDGFR*, *TSHR*, *MET*, and *RET*. DNA damage and repair pathway genes were mutated in 38% of tumors, namely, *ATM*, *TP53*, *CHEK2*, *CDKN1A*, *TLK1*, *E2F1*, *PML*, *RB1*, *XPC*, *ERCC%*, *BRCA1*, *XRCC3*, *MSH2*, *MSH3*, *PMS*, *POLE*, *FANCB*, and *FANCD2*. Epigenetic modification mutations were also seen in 59% of tumors, such as *SWI/SNF*, *ISWI/CHD*, *ARID1A*, *CREBBP*, *BRD7*, *KMT2C*, *NSD1*, *EZH1*, *HCAT7*, *SIRT6*, *PHF2*, *DNMT1*, among others. Mutations also occurred in ephrin genes regulating angiogenesis (16%), including *EPHA1*, *EPHA2*, *EPHA6*, *EPHA7*, *EPHA10*, *EPHA8*, *EPHB2*, and *EPHB4*.

Summary

- Hürthle-cell neoplasms are divided into Hürthle-cell adenoma and Hürthle-cell carcinoma. The latter is further divided into encapsulated minimally invasive, angioinvasive, and widely invasive Hürthle-cell carcinoma.
- Establishing follicular origin of oncocytic lesions in the thyroid is important to avoid diagnostic pitfalls in diagnosing medullary thyroid carcinoma, intrathyroidal parathyroid lesion(s), and secondary tumors of the thyroid gland.
- Papillary thyroid carcinoma variants can exhibit oncocytic cytoplasm such as tall cell, Warthin-like, and oncocytic variants.
- Near genome haplodization is unique to Hürthle-cell neoplasms and maybe secondary to mitochondrial mutations.
- *TERT* promotor mutations are more frequent in aggressive (angioinvasive and widely invasive) Hürthle-cell carcinoma.

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