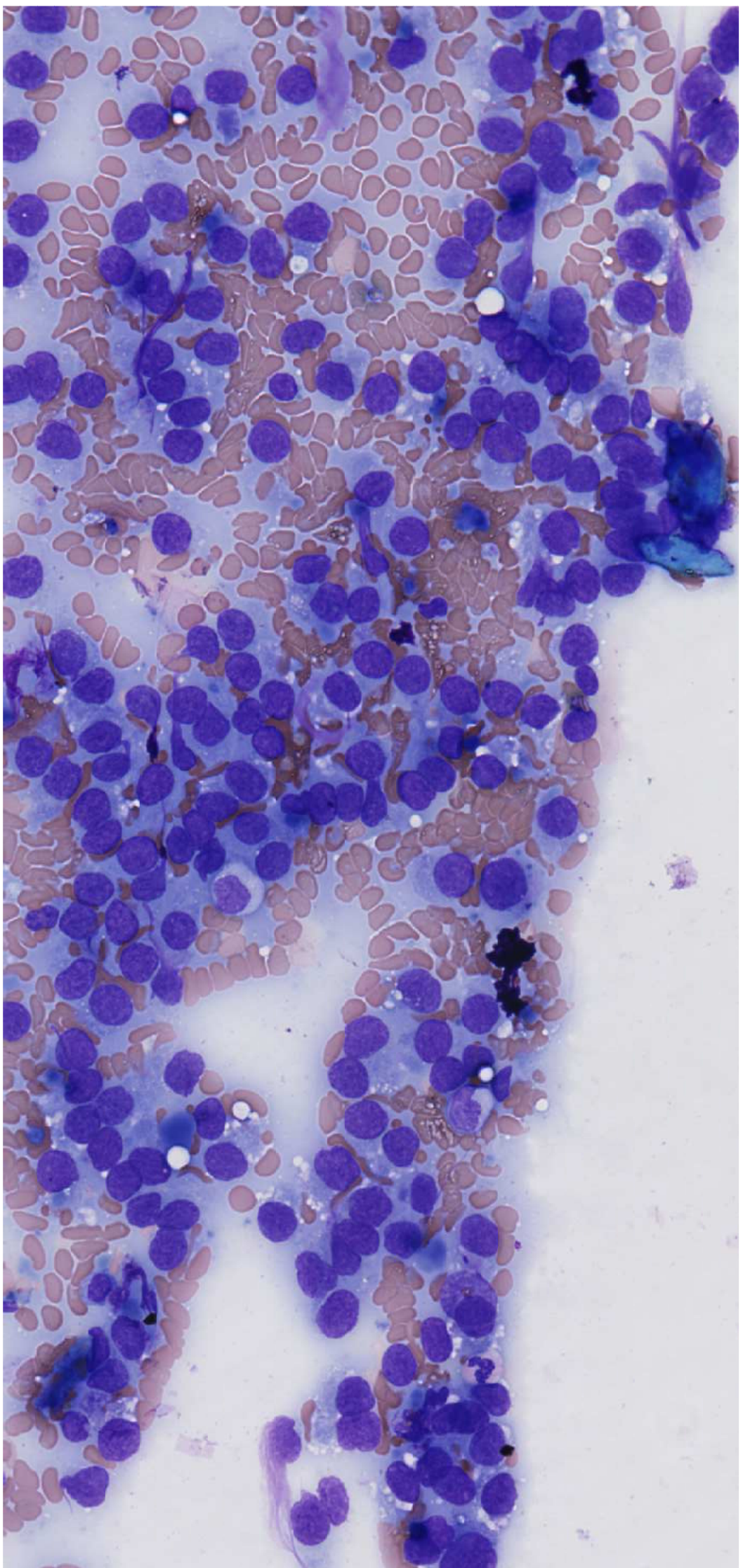


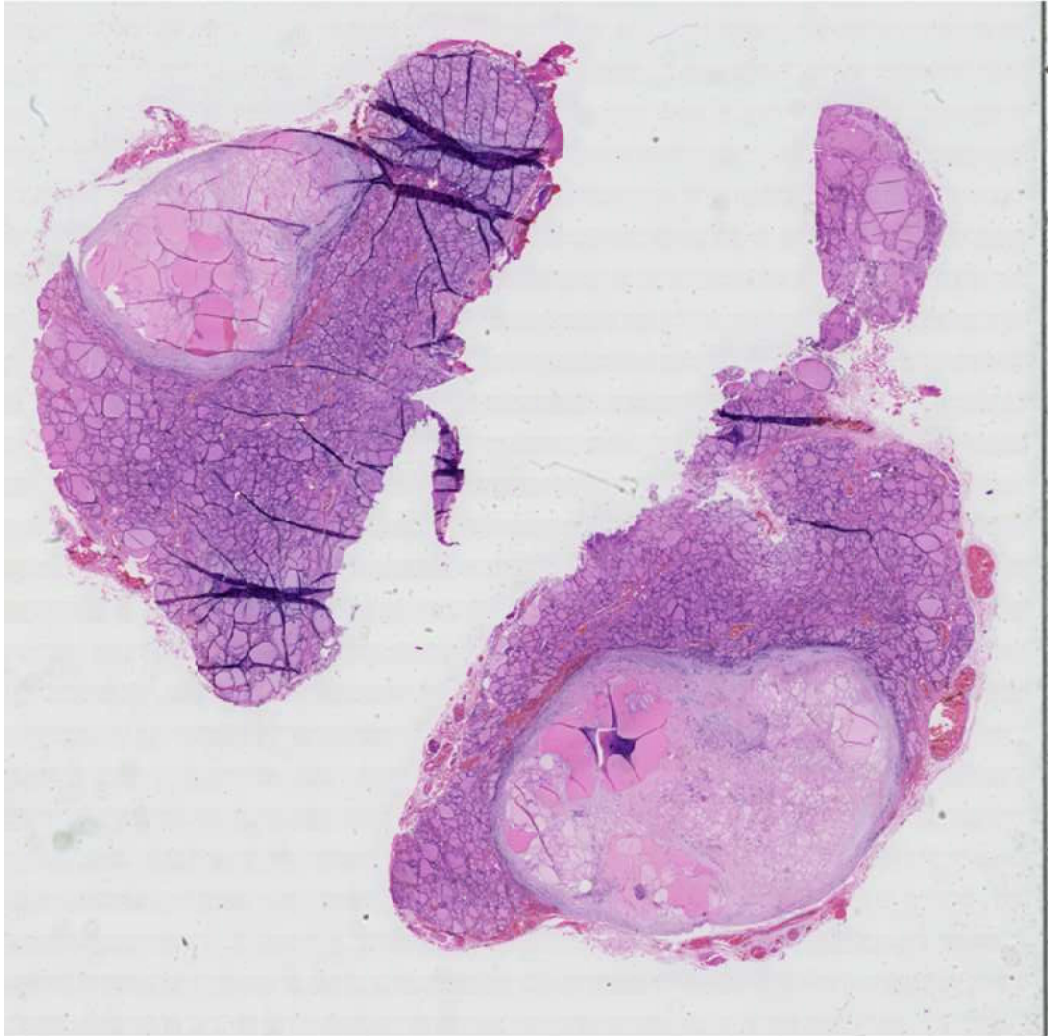
CASE 3

Single 1.8 cm thyroid nodule of the left lobe in a 16 year-old young woman; previous FNA Thy3a (Bethesda III). Left lobectomy

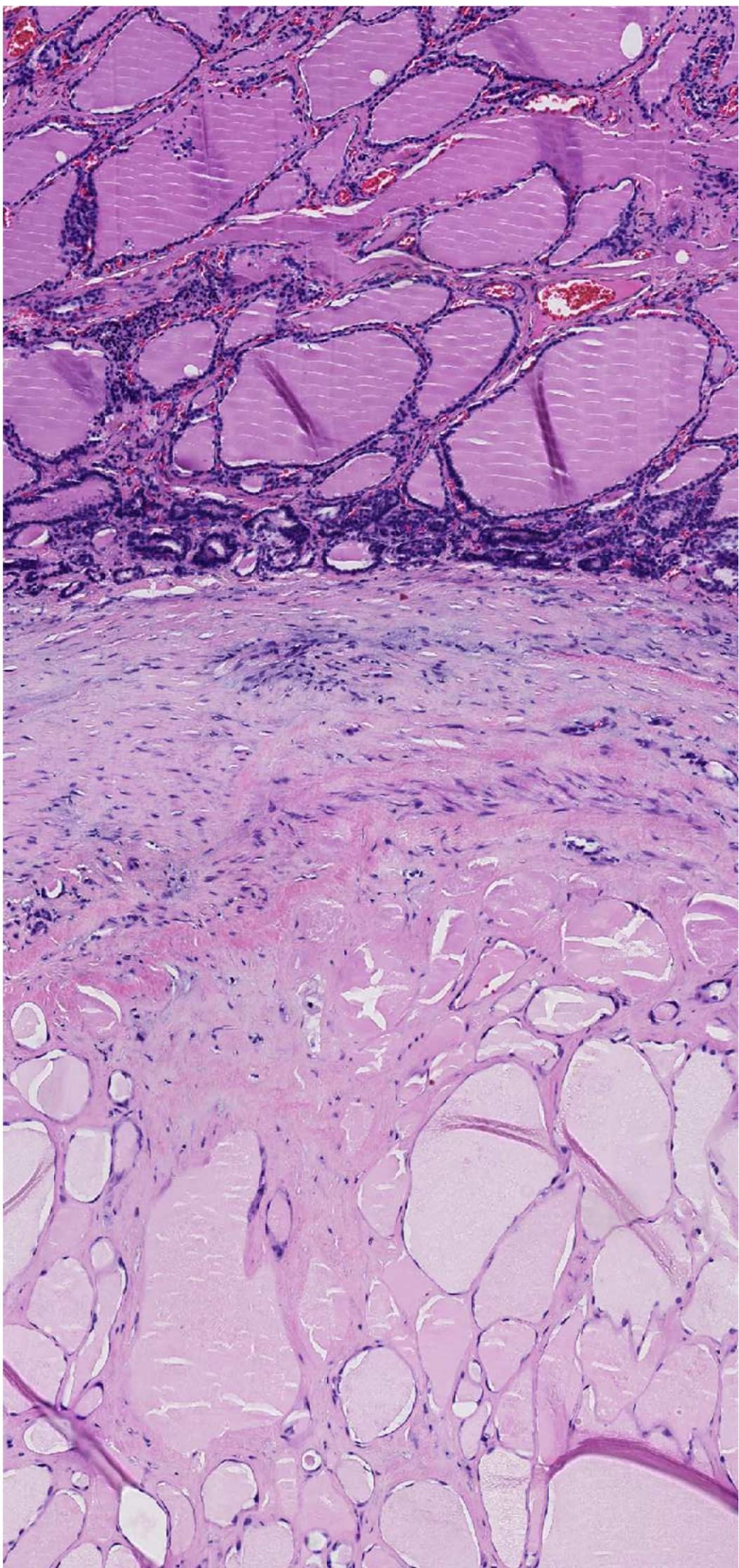
Giovanni Tallini, MD

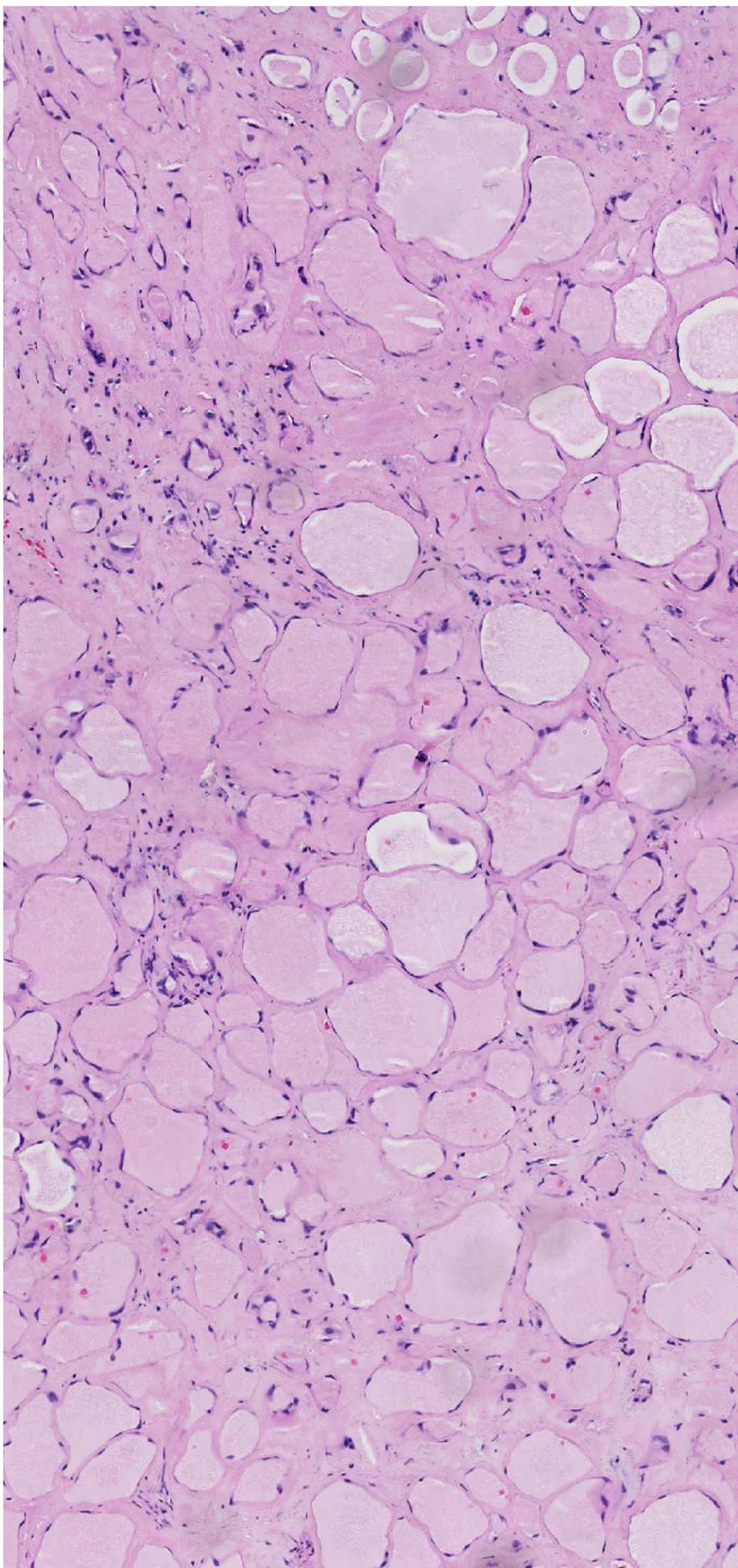
Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

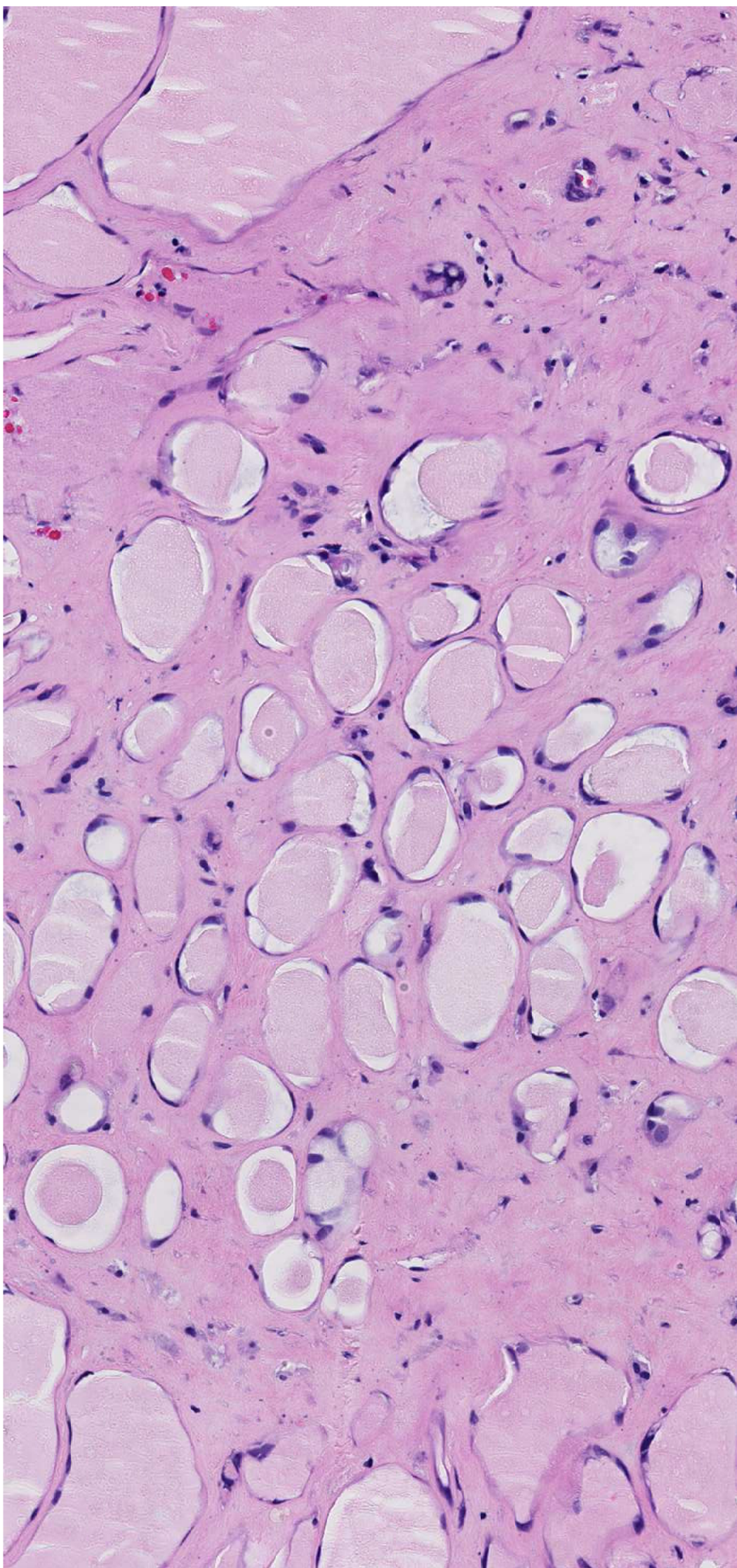


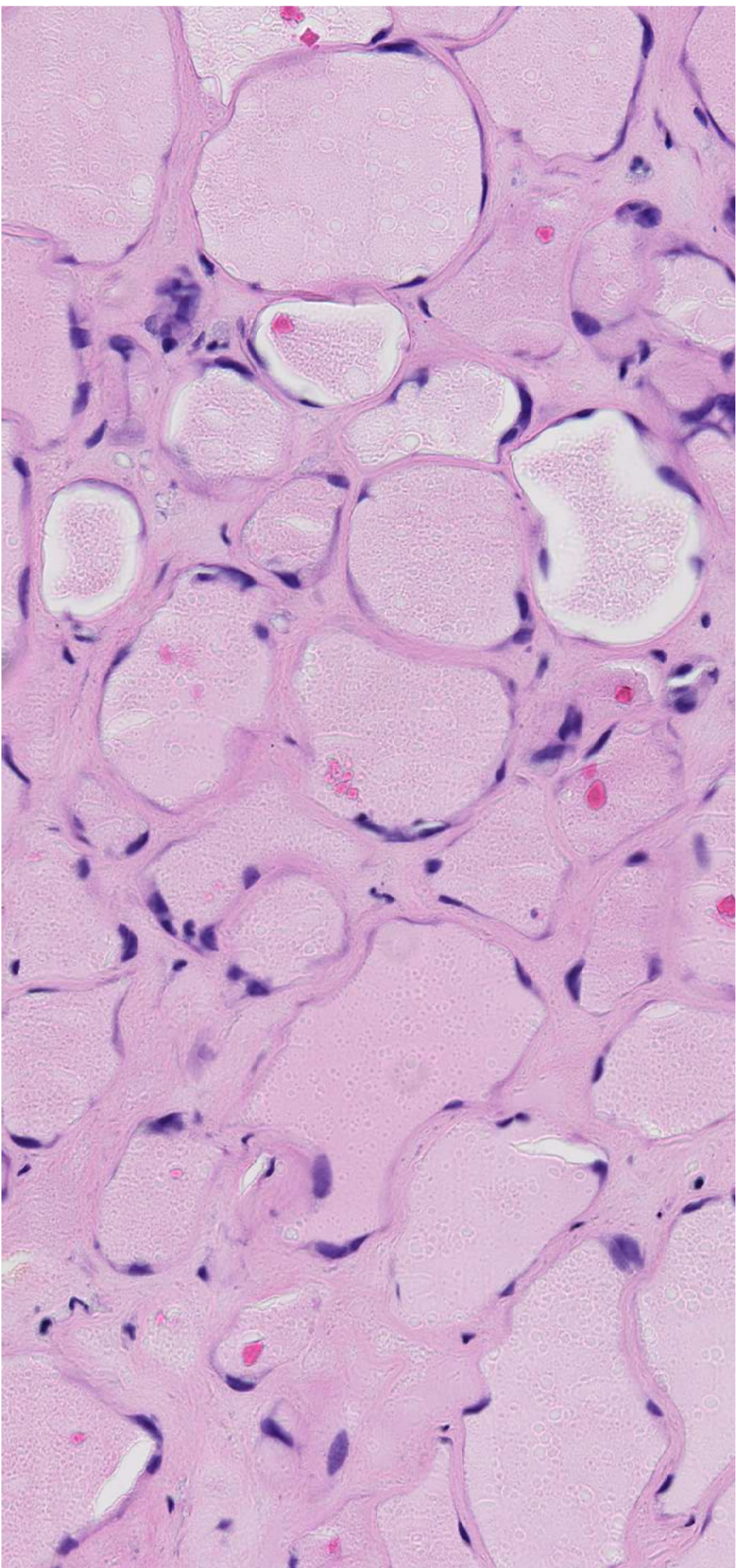












CASE 3

Follicular adenoma with atrophic follicles ("vanishing follicles")

Giovanni Tallini, MD

Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

CASE 1

Tumor tissue was analyzed by NGS: DICER1 p.Glu1705Lys (c.5113G>A, Esone 24, VAF 39%), Pathogenic ACMG variant (<https://varsome.com/>), not present in perilesional tissue (i.e. not a germline mutation)

Giovanni Tallini, MD

Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

CASE 1

Points for discussion

- Which type of thyroid nodules are DICER1 mutated?
- Should patients with somatic DICER1 mutation in thyroid nodules be tested for DICER1 syndrome?

Giovanni Tallini, MD

Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

CASE 1

Points for discussion

- Which type of thyroid nodules are DICER1 mutated?
- Should patients with somatic DICER1 mutation in thyroid nodules be tested for DICER1 syndrome?

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Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

DICER1-mutated tumors/nodules

Macrofollicular Variant of Follicular Thyroid Carcinoma: A Rare Underappreciated Pitfall in the Diagnosis of Thyroid Carcinoma
 Bongiovanni M, Sykiotis GP, La Rosa S, Bisig B, Trimech M, Missiaglia E, Gremaud M, Salvatori Chappuis V, De Vito C, Sciarra A, Foulkes WD, Pusztaszneri M
 Thyroid. 2020 Jan;30(1):72-80

Poorly differentiated thyroid carcinoma of childhood and adolescence: a distinct entity characterized by DICER1 mutations
 Chernock RD, Rivera B, Borrelli N, Hill DA, Fahiminiya S, Shah T, Chong AS, Aqil B, Mehrad M, Giordano TJ, Sheridan R, Rutter MM, Dehner LP, Foulkes WD, Nikiforov YE
 Mod Pathol. 2020 Jul;33(7):1264-1274

Prevalence and Spectrum of DICER1 Mutations in Adult-onset Thyroid Nodules with Indeterminate Cytology
 Chong AS, Nikiforov YE, Condello V, Wald AI, Nikiforova MN, Foulkes WD, Rivera B
 J Clin Endocrinol Metab. 2021 Mar 25;106(4):968-977.

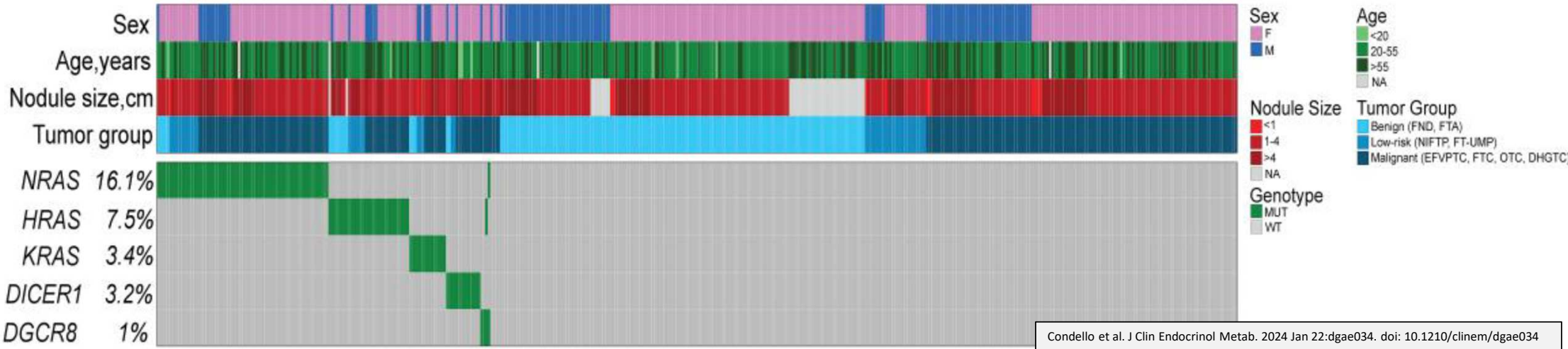
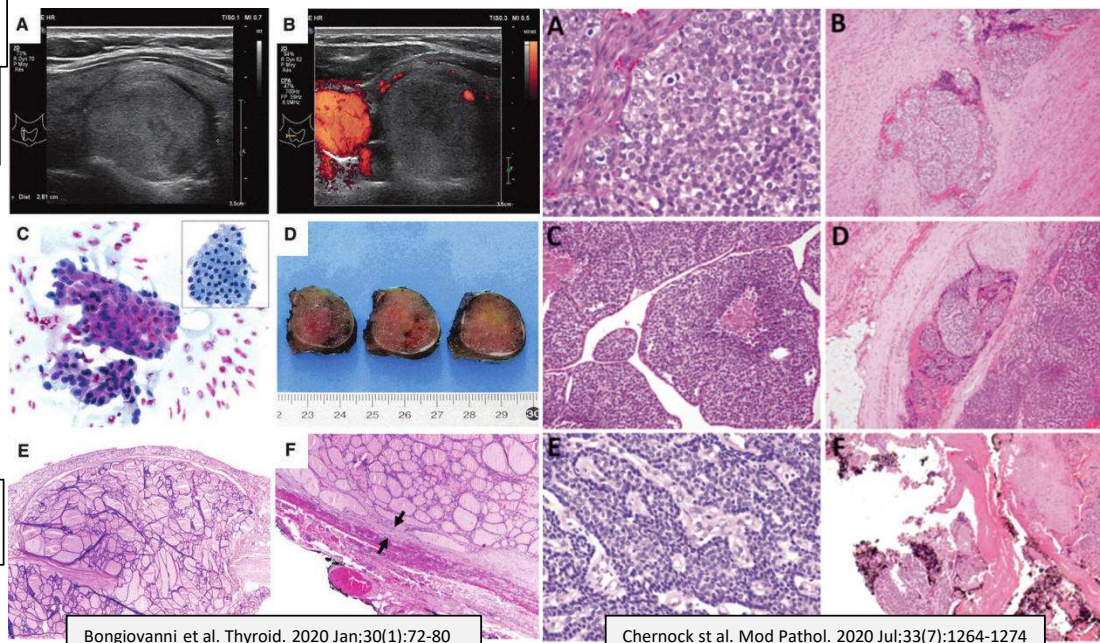
Macrofollicular variant follicular thyroid tumors are DICER1 mutated and exhibit distinct histological features
 Juhlin CC, Stenman A, Zedenius J
 Histopathology. 2021 Oct;79(4):661-666

Expanding the spectrum of thyroid carcinoma with somatic DICER1 mutation: a survey of 829 thyroid carcinomas using MSK-IMPACT next-generation sequencing platform
 Ghossein CA, Dogan S, Farhat N, Landa I, Xu B
 Virchows Arch. 2022 Feb;480(2):293-302

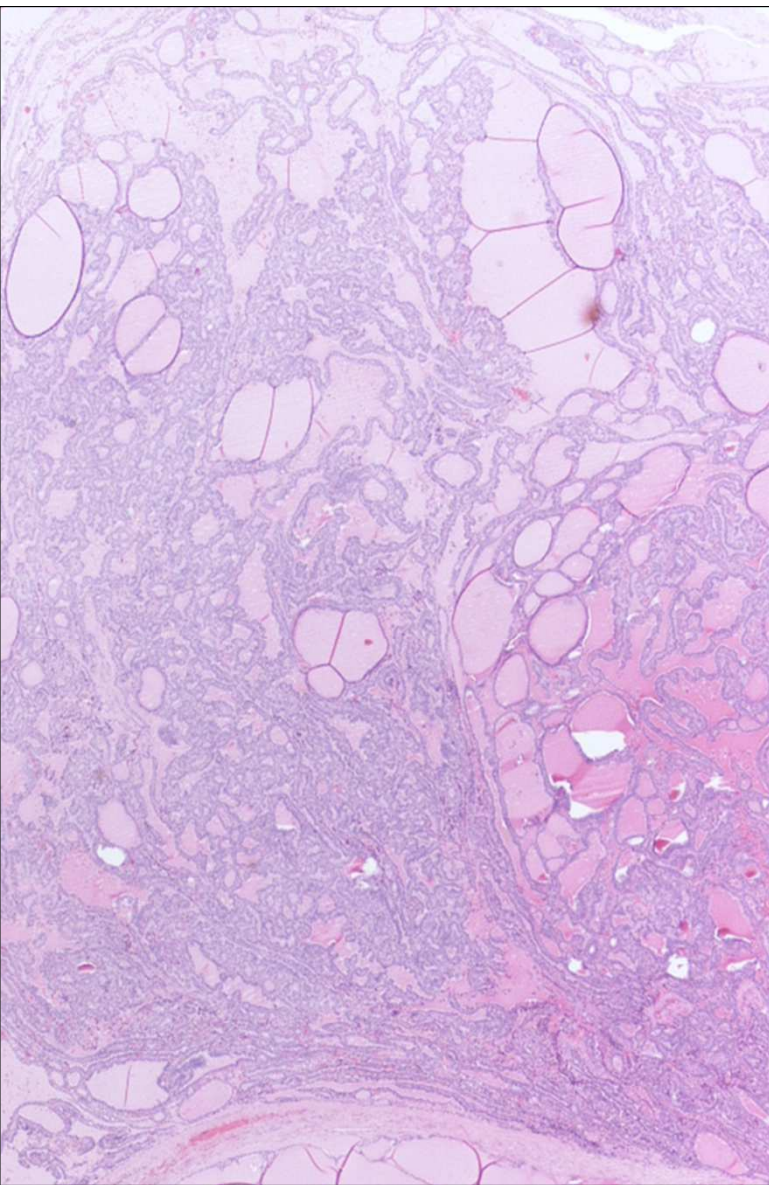
Evaluation of the Molecular Landscape of Pediatric Thyroid Nodules and Use of a Multigene Genomic Classifier in Children
 Gallant JN, Chen SC, Ortega CA, Rohde SL, Belcher RH, Nettekville JL, Baregamian N, Wang H, Liang J, Ye F, Nikiforov YE, Nikiforova MN, Weiss VL
 JAMA Oncol. 2022 Sep 1;8(9):1323-1327

Prevalence, Molecular Landscape and Clinical Impact of DICER1 and DGCR8 Mutated Follicular-Patterned Thyroid Nodules
 Condello V, Poma AM, Macerola E, Vignali P, Paulsson JO, Zedenius J, Basolo F, Juhlin CC
 J Clin Endocrinol Metab. 2024 Jan 22:dgae034. doi: 10.1210/clinem/dgae034

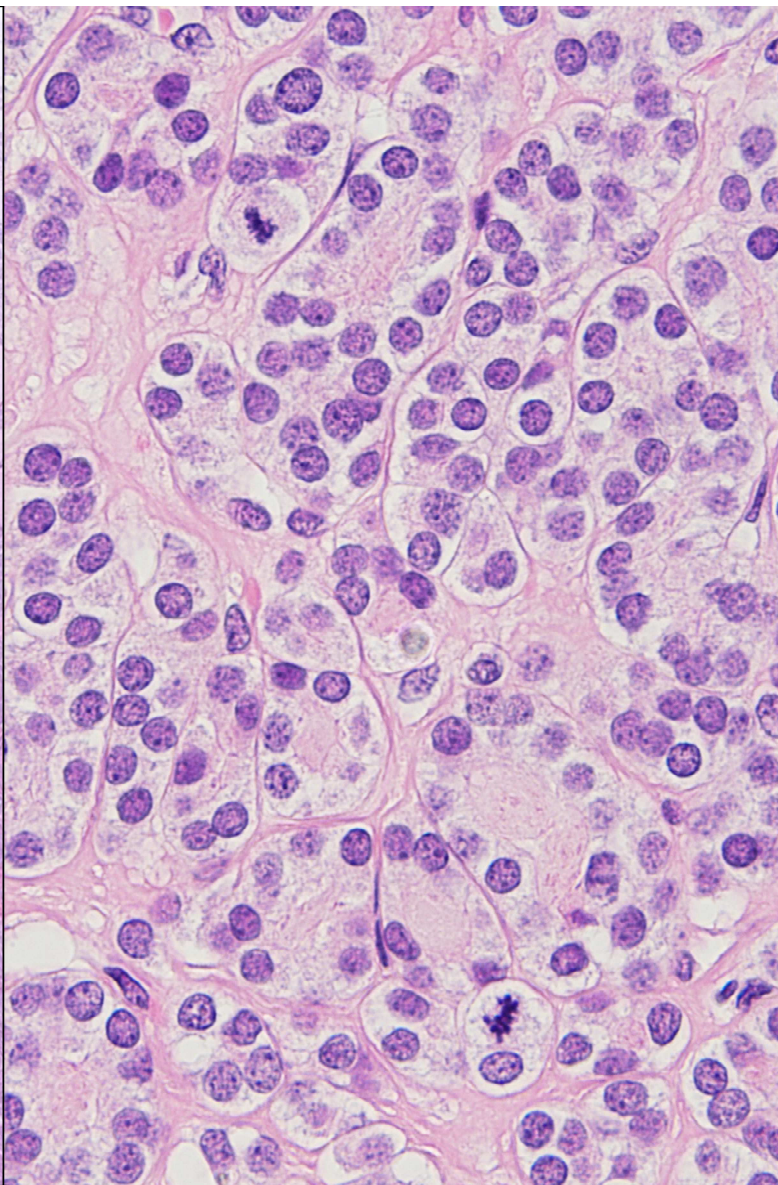
Histological clues of DICER1 mutations in thyroid nodules
 Jung CK, Liu Z, Hirokawa M, Bychkov A
 Virchows Arch. 2024 Oct;485(4):755-757



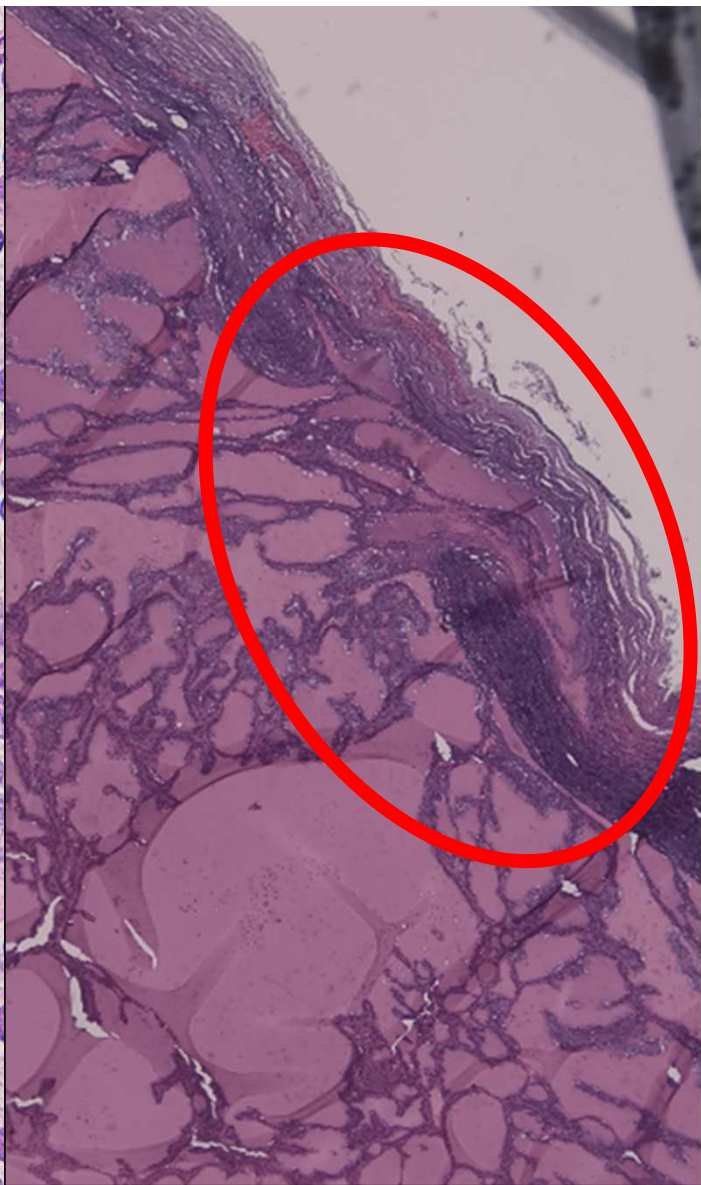
Condello et al. J Clin Endocrinol Metab. 2024 Jan 22:dgae034. doi: 10.1210/clinem/dgae034



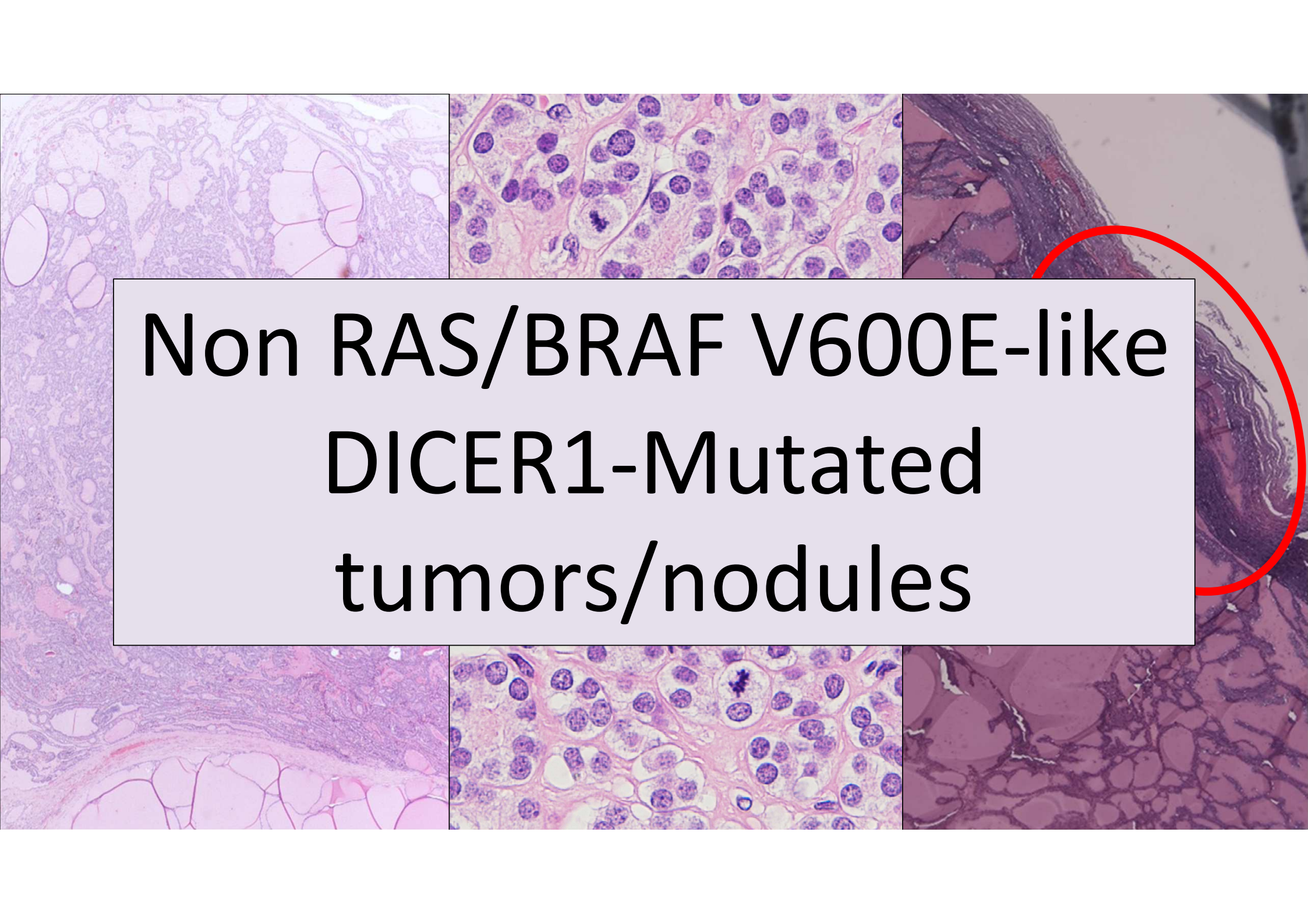
Follicular Nod. Disease



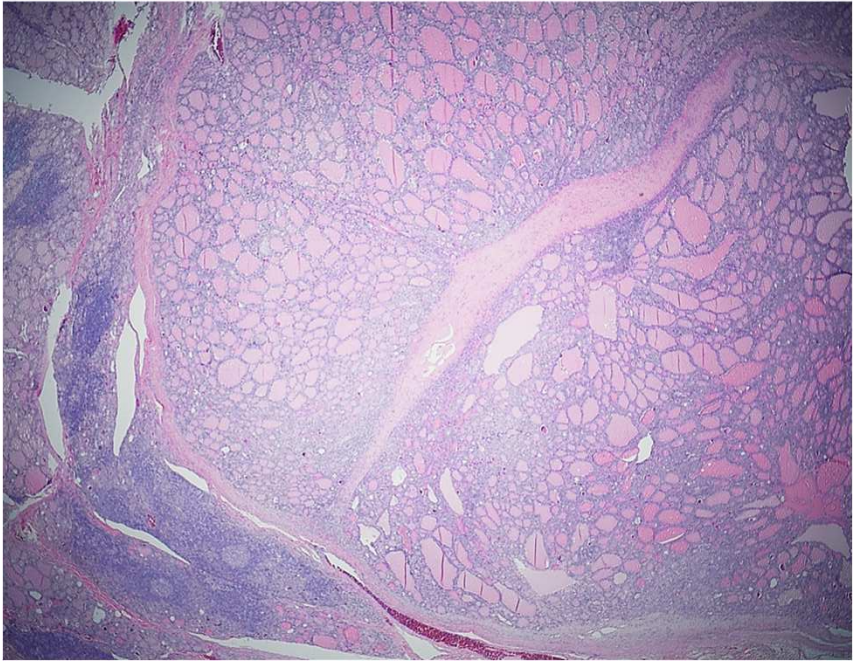
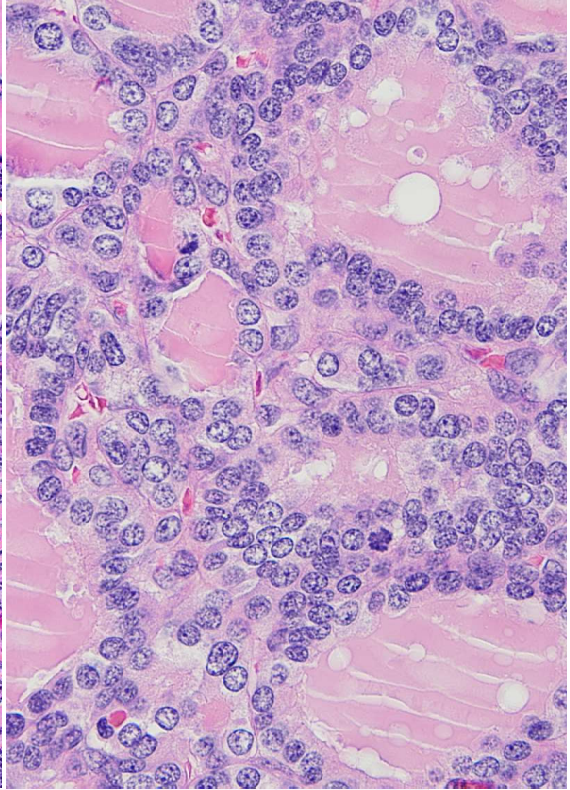
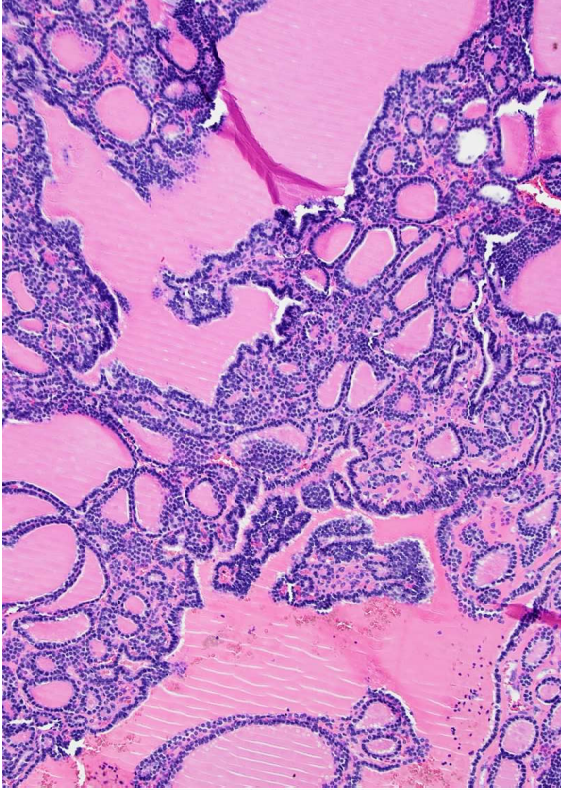
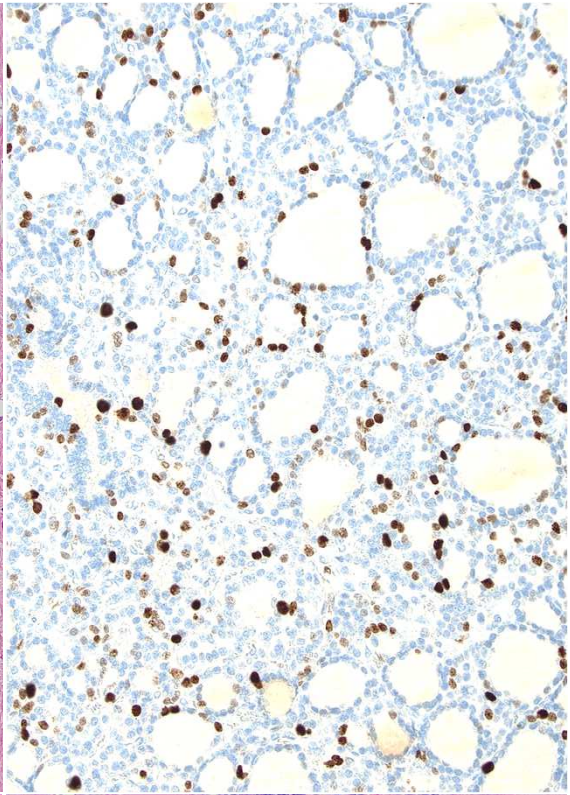
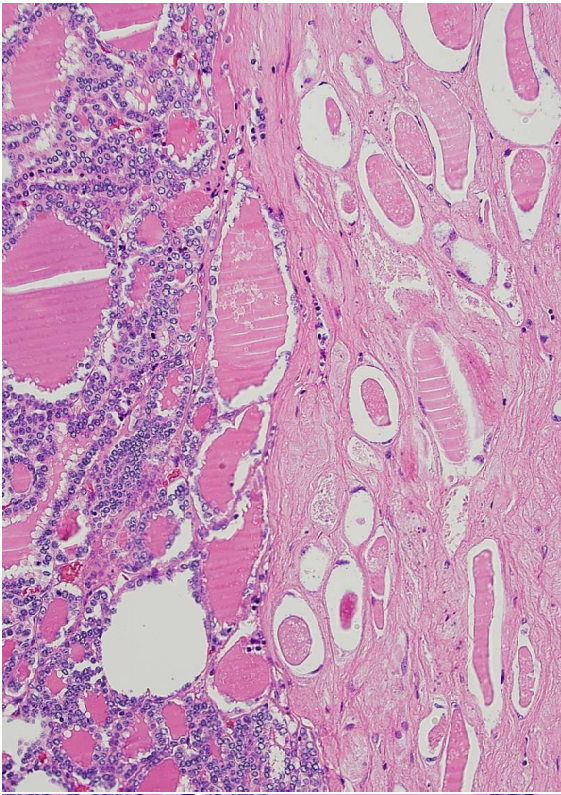
Follicular Adenoma

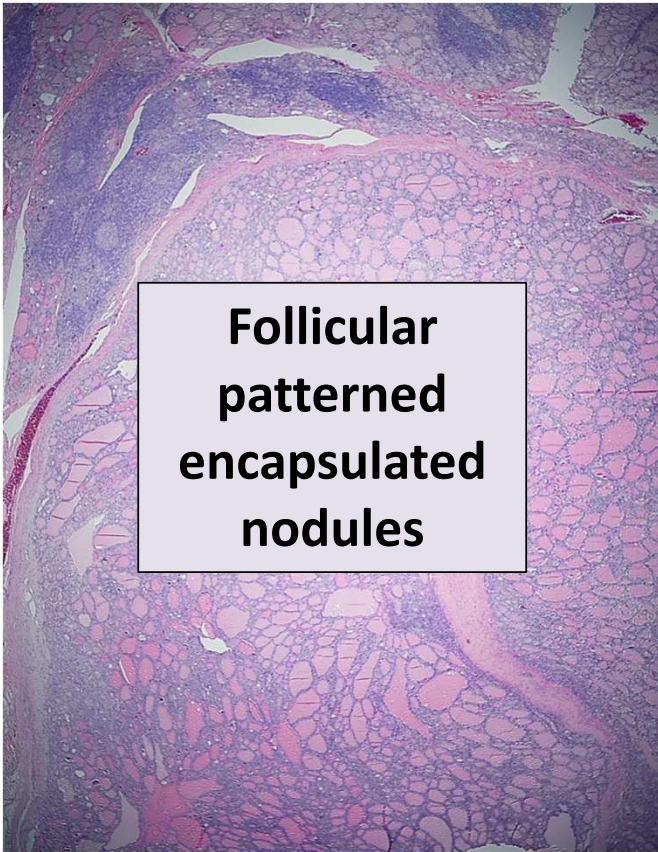


Foll. Ca. (Pap. architecture)

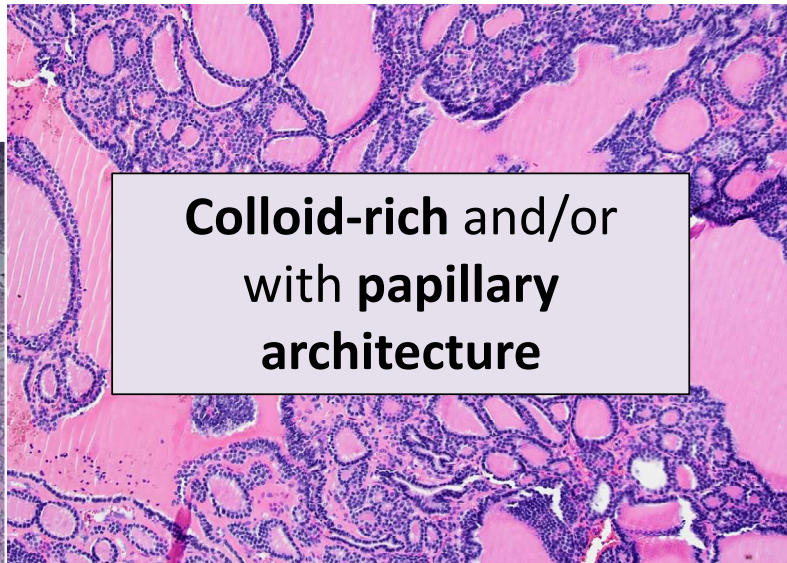
The image is a composite of six histological panels. The top row shows three panels: a low-magnification view of a tumor with glandular structures, a high-magnification view of glandular epithelial cells with hyperchromatic nuclei, and a high-magnification view of a tumor nodule with a red circle highlighting a specific area. The bottom row shows three panels: a low-magnification view of a tumor with glandular structures, a high-magnification view of glandular epithelial cells with hyperchromatic nuclei, and a high-magnification view of a tumor nodule with a red circle highlighting a specific area.

**Non RAS/BRAF V600E-like
DICER1-Mutated
tumors/nodules**

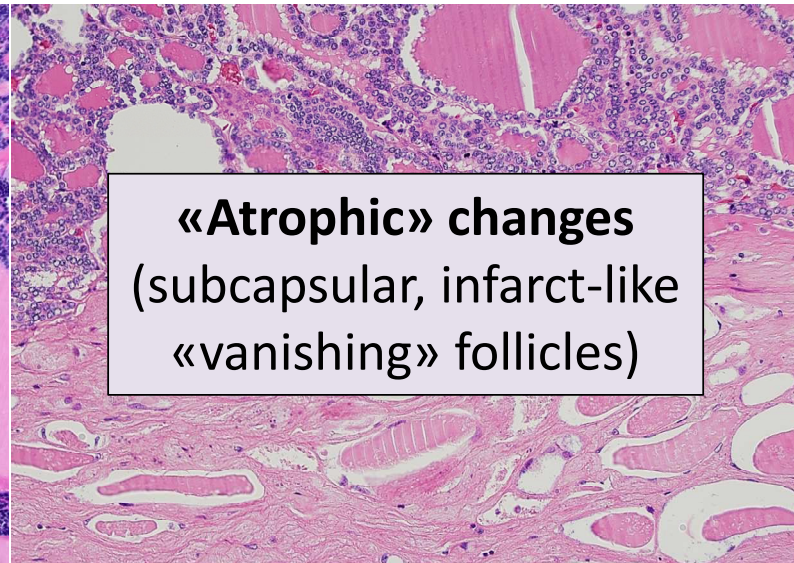




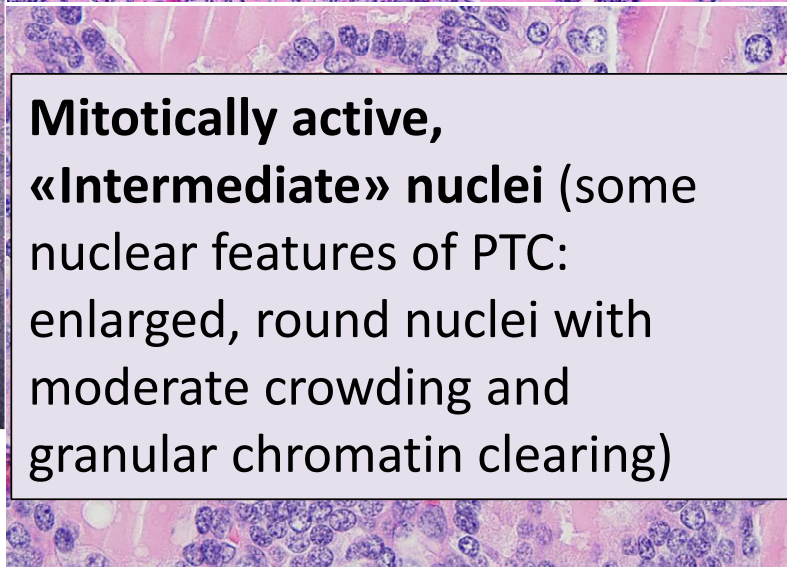
Follicular patterned encapsulated nodules



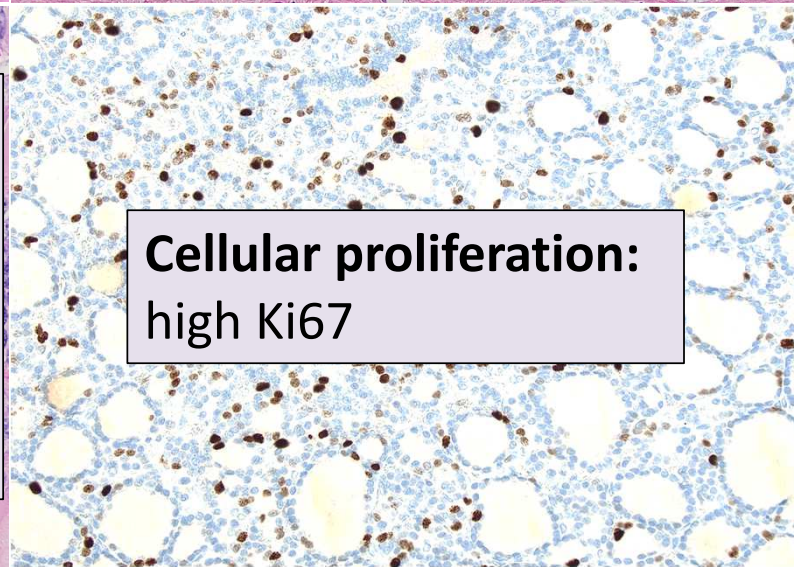
Colloid-rich and/or with papillary architecture



«Atrophic» changes (subcapsular, infarct-like «vanishing» follicles)



Mitotically active, «Intermediate» nuclei (some nuclear features of PTC: enlarged, round nuclei with moderate crowding and granular chromatin clearing)



Cellular proliferation: high Ki67

The background of the slide features several histological images of thyroid tissue. At the top, there are two horizontal strips showing follicular structures with pink-stained colloid and purple-stained nuclei. On the left side, there is a vertical strip showing a cross-section of a thyroid nodule with a capsule. On the right side, there are two vertical strips: the upper one shows follicular architecture with a label 'e' next to it, and the lower one shows a higher magnification of follicular cells with brown-stained nuclei, likely indicating immunohistochemical staining for Ki67.

Somatic DICER1 mutations in thyroid nodules/tumors:

- **Prevalence: 1-3%**, overall (follicular patterned nodules/tumors)
- Common in **pediatric age/young adults**
- Act as RAS-like mutations, and are therefore reproduce **follicular patterned** histology
- **Tumors mostly benign**, but may be malignant (even poorly differentiated)
- Follicular patterned **encapsulated** tumors, often **colloid-rich** and/or with **papillary architecture**; «**Atrophic**» changes (subcapsular, infarct-like vanishing follicles); «**Intermediate**» nuclei (some nuclear features of PTC: enlarged, round nuclei with moderate crowding and granular chromatin clearing)
- **Cellular proliferation** (mitoses, elevated Ki67)

CASE 1

Points for discussion

- Which type of thyroid nodules are DICER1 mutated?
- Should patients with somatic DICER1 mutation in thyroid nodules be tested for DICER1 syndrome?

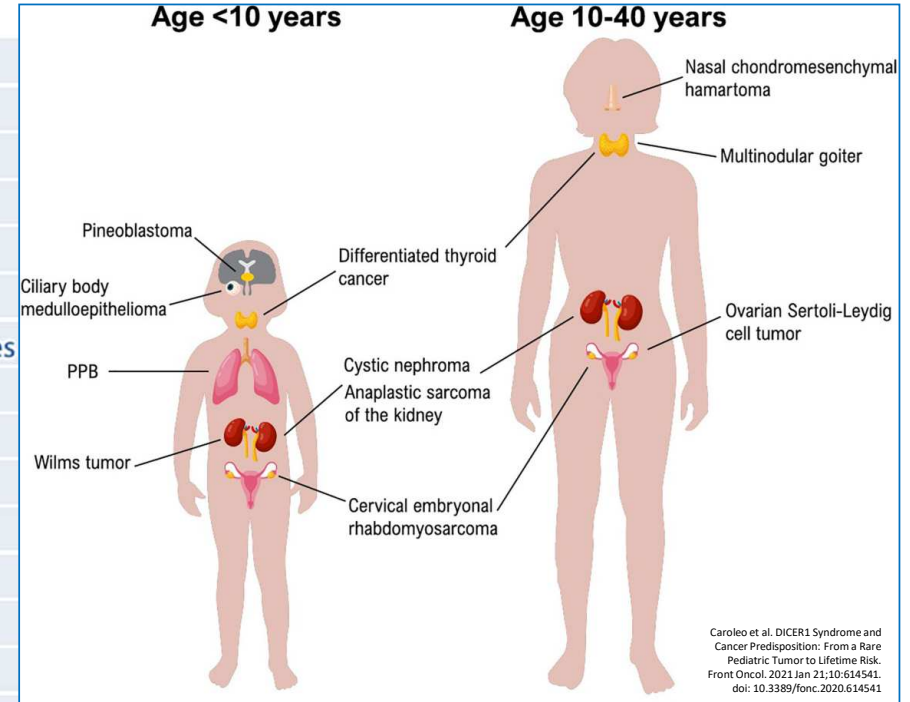
Giovanni Tallini, MD

Anatomic Pathology, University of Bologna Medical Center
giovanni.tallini@unibo.it

DICER1 Syndrome

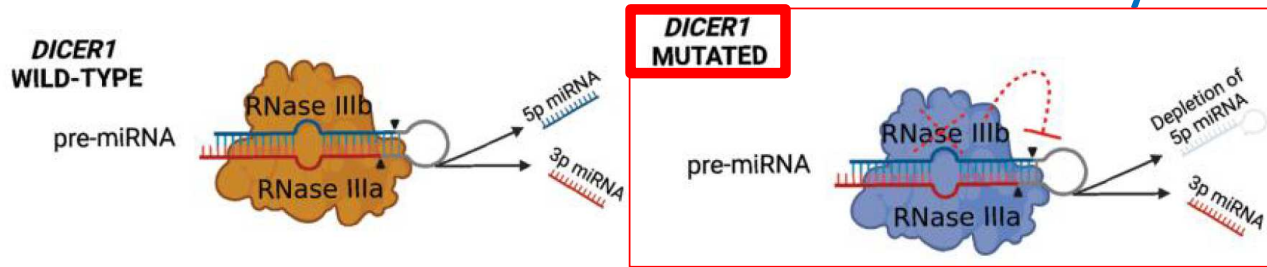
Table 1. *DICER1*-associated neoplasms.

Pleuropulmonary blastoma (PPB) and PPB-like neoplasms
Pleuropulmonary blastoma, type I, IR, II, III
PPB-like Sertoli-Leydig cell tumor of lung
Pediatric cystic neoplasms and <i>DICER1</i> -sarcoma (anaplastic sarcoma of kidney)
Nasal chondromesenchymal hamartoma
Central nervous system sarcoma with rhabdomyosarcoma/PPB III-like features
Sertoli-Leydig cell tumor with and without heterologous features and type I PPB-like features
Peritoneal, ovarian and fallopian tube sarcoma with PPB-like features
<i>DICER1</i> -associated cystic hepatic neoplasm with type I PPB-like features
Cervical embryonal rhabdomyosarcoma
Teratoid and primitive neuroepithelial neoplasms
Cervical-thyroid teratoma
Malignant teratoid neoplasm of sacrococcygeal region
Ciliary body medulloepithelioma
Pituitary blastoma
Pineoblastoma
Embryonal tumor with multilayered rosettes
Thyroid
Multinodular hyperplasia (goiter)
Papillary thyroid carcinoma, invasive follicular variant
Follicular carcinoma, pediatric type
Poorly differentiated thyroid carcinoma, pediatric type
Intestine
Hamartomatous polyp with juvenile polyp-like features

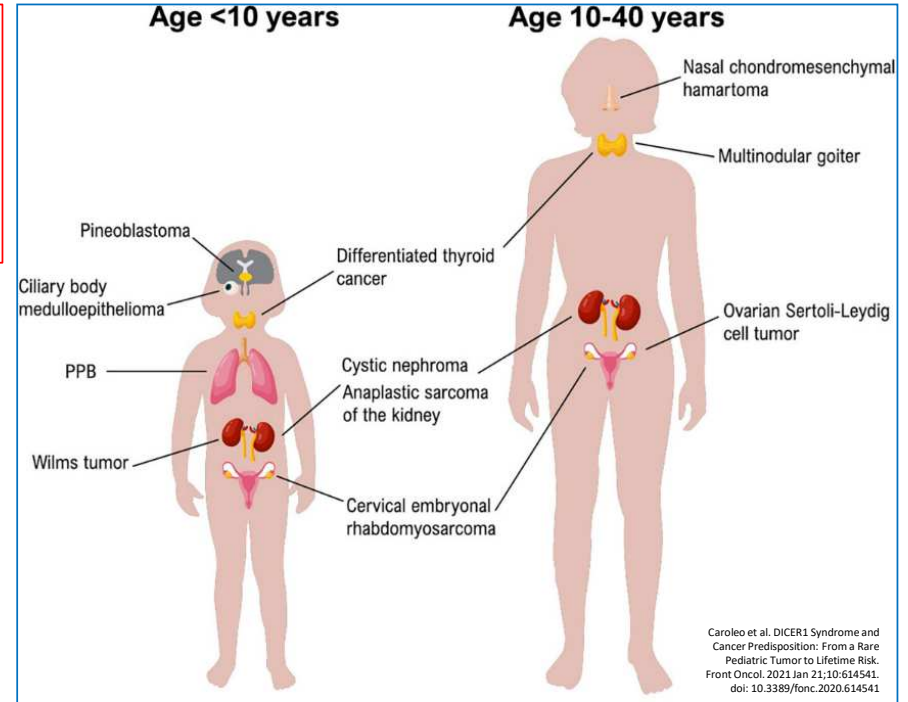


- *DICER1* syndrome (OMIM 606241, 601200) is an autosomal dominant familial tumor predisposition disorder with heterozygous *DICER1* germline mutation
- *DICER1* on chromosome 14q32.13 encodes an RNA endonuclease (Dicer) involved in the post-transcriptional gene expression of over 30% of protein-coding genes by modulating microRNAs
- Reduced penetrance, which likely decreases the rate of familial cases; in cases with pleuropulmonary blastoma, ~80% of *DICER1* germline pathogenic variants are inherited by a parent, ~20% are *de novo*

DICER1 Syndrome



Disabling mutations in RNase III domains of *DICER1* lead to 5p miRNA depletion. Schematic representation of the mechanism by which *DICER1* cleaves pre-miRNA in wild-type and mutated conditions; hotspot mutations in the RNase IIIb domain lead to a significant reduction of 5p strand miRNAs in *DICER1* mutants compared with wild-type cases [Condello et al. J Clin Endocrinol Metab. 2024 Jan 22:dgae034. doi: 10.1210/clinem/dgae034] Disabling mutations in RNase III domains of *DICER1* lead to 5p miRNA depletion. Schematic representation of the mechanism by which *DICER1* cleaves pre-miRNA in wild-type and mutated conditions; hotspot mutations in the RNase IIIb domain lead to a significant reduction of 5p strand miRNAs in *DICER1* mutants compared with wild-type cases [Condello et al. J Clin Endocrinol Metab. 2024 Jan 22:dgae034. doi: 10.1210/clinem/dgae034]



Caroleo et al. *DICER1* Syndrome and Cancer Predisposition: From a Rare Pediatric Tumor to Lifetime Risk. Front Oncol. 2021 Jan 21;10:614541. doi: 10.3389/fonc.2020.614541

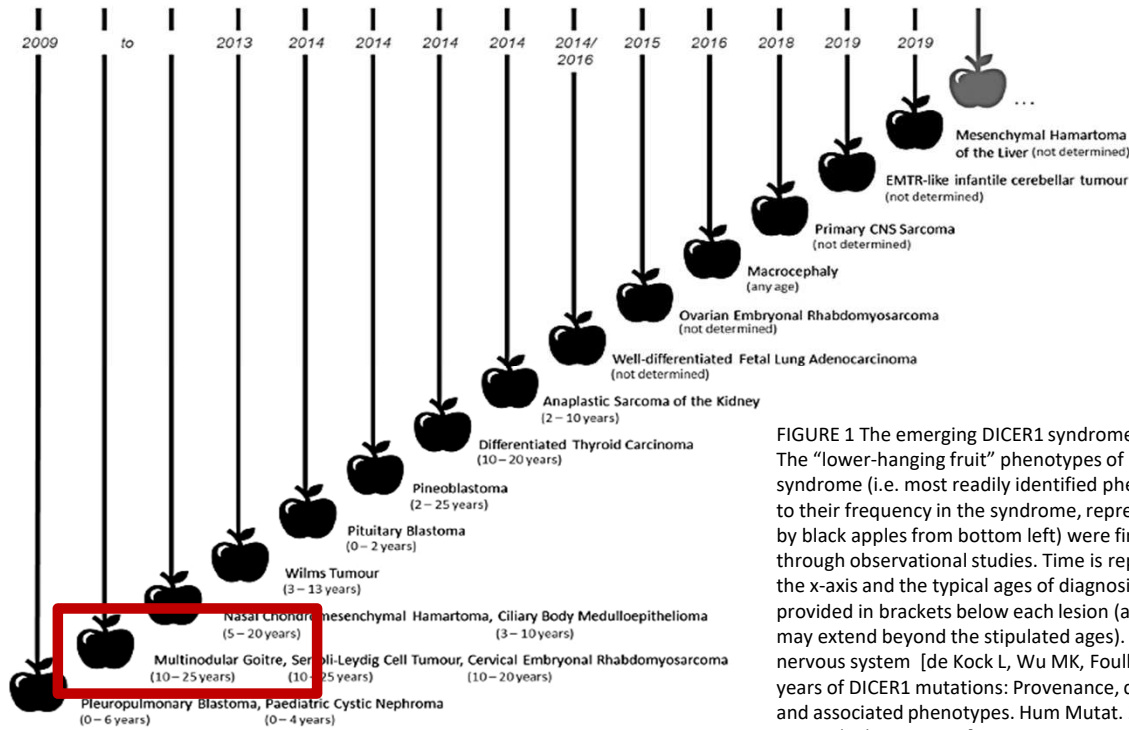


FIGURE 1 The emerging *DICER1* syndrome phenotype. The “lower-hanging fruit” phenotypes of *DICER1* syndrome (i.e. most readily identified phenotypes due to their frequency in the syndrome, represented here by black apples from bottom left) were first noted through observational studies. Time is represented on the x-axis and the typical ages of diagnosis are provided in brackets below each lesion (although risk may extend beyond the stipulated ages). CNS, central nervous system [de Kock L, Wu MK, Foulkes WD. Ten years of *DICER1* mutations: Provenance, distribution, and associated phenotypes. Hum Mutat. 2019 Nov;40(11):1939-1953]

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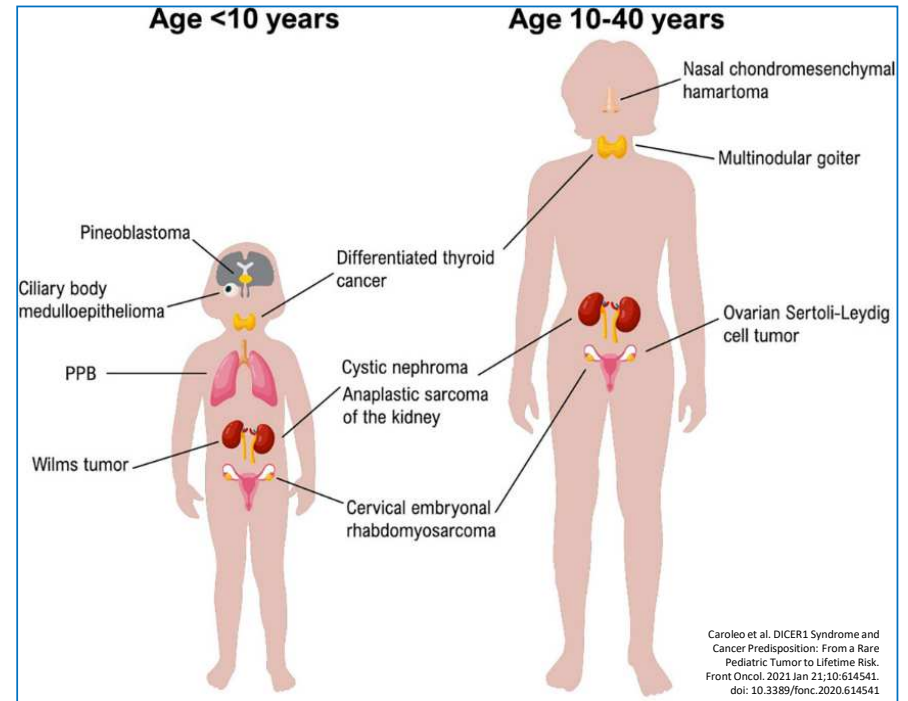
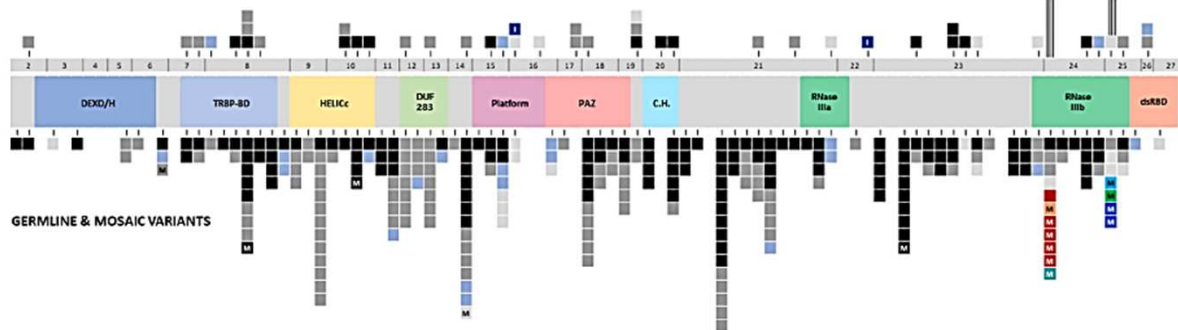
DICER1 Syndrome

PATHOGENIC & LIKELY-PATHOGENIC (Confirmed Somatic)

KEY

Variant Types:	Germline	Somatic	Mosaic
Non-Hotspot Alterations:			
Silent	284	66	5
Missense (non-hotspot)	15	7	1
Nonsense	97	18	1
Frameshift	135	14	3
Splicing	21	4	0
In-frame deletion	6	2	0
Large out-of-frame deletion	2	0	0
Full gene deletion	8	NA	0
Somatic LOH	NA	21	NA
Hotspot Missense Mutations:			
p.E1705	1	356	10
p.D1709	0	44	1
p.D1713	1	86	4
p.G1809	0	3	1
p.D1810	0	51	1
p.E1813	0	43	1
	0	129	2
	285	422	15

Other:
M Mosaic



Caroleo et al. DICER1 Syndrome and Cancer Predisposition: From a Rare Pediatric Tumor to Lifetime Risk. Front Oncol. 2021 Jan 21;10:614541. doi: 10.3389/fonc.2020.614541

FIGURE 4 Pathogenic and likely pathogenic DICER1 alterations published before January 31st, 2019. Only unique-per-family (UPF) germline variants and confirmed-somatic mutations considered pathogenic or likely pathogenic have been plotted along the length of the unfolded DICER1 protein (n = 722). The 422 confirmed somatic events are plotted above the protein, except for the 21 confirmed-somatic LOH events that are shown at the bottom of the figure. LOH, loss of heterozygosity; NA, not applicable [de Kock L, Wu MK, Foulkes WD. Ten years of DICER1 mutations: Provenance, distribution, and associated phenotypes. Hum Mutat. 2019 Nov;40(11):1939-1953]

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DICER1 Syndrome

PATHOGENIC & LIKELY-P

KEY

Variant Types:

Non-Hotspot Alterations:

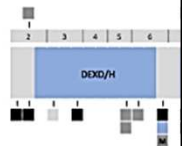
- Silent
- Missense (non-hotspot)
- Nonsense
- Frameshift
- Splicing
- In-frame deletion
- Large out-of-frame deletion
- Full gene deletion
- Somatic LOH

Hotspot Missense Mutations:

- p.E1705
- p.D1709
- p.D1713
- p.G1809
- p.D1810
- p.E1813

Other:

- M Mosaic



GERMLINE & MOSAIC VARIATION

- **DICER1 Putative tumor suppressor** (in most cases biallelic inactivation)
- **Dysregulation of miRNA biogenesis** with reduced expression of 5p miRNAs and increased levels of 3p miRNAs compared with wild-type cases
- **Mutations:**
 - **Hot spot** mutations (in the functionally important RNase III (A and B) region: exons 24-25) are typically somatic, but may represent second hit in patients with germline DICER1 mutation (need to evaluate DICER1 germline: entire gene)
 - **Non-Hot spot** mutations (throughout the gene, usually truncating) can be germline or somatic
 - **Mutually exclusive** with alterations in other thyroid cancer-related genes
- Many patients with DICER1 syndrome have follicular nodular disease and sometimes follicular patterned tumors, **but germline DICER1 mutations are rare in adult patients with incidentally discovered DICER1 hot spot mutated nodules**
- ✓ **Should patients with DICER1 mutated thyroid nodules discovered incidentally, undergo clinical genetics evaluation to rule out DICER1 syndrome?**
 - **Ideally yes** (germline mutations can be transmitted to progeny!), although germline mutations are uncommon in patients with unremarkable medical history

FIGURE 4 Pathol Oncol. 2019;31:31-37. Original mutations color-coded above the protein structure. Foulkes WD. Thyroid cancer phenotypes.

nodromesenchymal
oma

multinodular goiter

ovarian Sertoli-Leydig
cell tumor

et al. DICER1 Syndrome and
Predisposition: From a Rare
Matric Tumor to Lifetime Risk.
J Clin Oncol. 2021 Jan 21;10:614541.
doi: 10.3389/onc.2020.614541

sosomal
heterozygous

endonuclease
expression of over
DNAs
of familial
% of
by a parent, ~

DICER1-Mutated tumors and DICER1 Syndrome

Congenital nodular goitre: TG, TPO, NIS, PDS, IYD, DOUX2, DUOXA2

Nodular goitre debuting at >10 years of age: DICER1, NKX2-1, KEAP1

Altaraihi M et al. Prevalence of Pathogenic Germline DICER1 Variants in Young Individuals Thyroidectomised Due to Goitre - A National Danish Cohort. Front Endocrinol (Lausanne). 2021 Sep 6;12:727970. doi: 10.3389/fendo.2021.727970

Nodular goitre debuting during the first decade: DICER1, RGS12, GRPEL1, CLIC6, WFS1

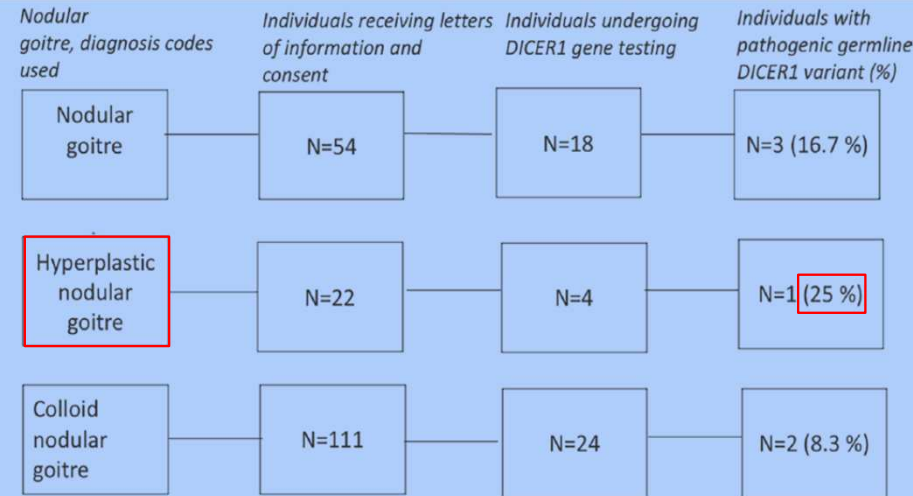
Mutation in 6/46 children/young adults aged ≤ 25 years with thyroidectomy for goiter with germline DICER1 testing (13%)

Table 2 Clinical and genetic characteristics of all DICER1-mutated cases

Case ID	Diagnosis	Sex	Age, years	Nodule size, cm	Architectural pattern	Papillary formations	Atrophic changes	DICER1 hotspot mutation in thyroid lesions (AF%)	Constitutional DICER1 mutation (AF%)	Other mutations apart from DICER1 in thyroid lesions (AF%)
EC 1	FTA	F	19	5.5	Macrofollicular	YES	NO	c.5126 A>G p.D1709G (32%)	NP	None
EC 2	NIFTP	F	22	7.8	Macrofollicular	YES*	NO	c.5126 A>G p.D1709G (30%)	NP	None
EC 3	MFPTC	F	36	2.5	Macrofollicular	YES	YES	c.5439 G>T p.E1813D (63%)	NP	None
EC 4 ¹	PDTC	F	15	2.5	Microfollicular	NO	YES	c.5127 T>G p.D1709E (27%)	c.5464 G>A p.D1822N (VUS)	None
EC 5 ¹	FTA	F	15	1.2	Macrofollicular and solid	NO	YES	c.5437 G>A p.E1813K (15%)	c.5464 G>A p.D1822N (VUS)	None
EC 6	FTA	F	49	1.5	Microfollicular	NO	YES	c.5126 A>G p.D1709G (37%)	NP	None
EC 7	FTA	F	55	2	Macrofollicular	NO	YES	c.5126 A>G p.D1709G (43%)	NP	None
EC 8	DHGTC	M	25	2.8	Macrofollicular	YES	YES	c.5429 A>T p.E1810V (60%)	NP	None
EC 9	MFPTC	F	17	2	Macrofollicular	YES	YES	c.5126 A>G p.D1709G (41%)	WT	None
EC 10	MFPTC	F	22	3.2	Macrofollicular	YES	YES	c.5429 A>T p.E1810V (51%)	NP	None
KI 1	TFND	F	15	12	Macrofollicular	YES	YES	NO	c.4154_4155ins p.Y13* (48%)	None
KI 2	MFPTC	F	13	1.2	Macrofollicular	YES	YES	c.5126 A>G p.D1709G (20%)	c.2830 C>T p.R944* (50%)	FGFR4 p.T179A (51%)
KI 3	MFPTC	F	29	5	Macrofollicular	YES	YES	c.5437 G>C p.E1813Q (39%)	c.2908 G>T p.E790* (40%)	None
KI 4	MFPTC	F	33	1.5	Macrofollicular	YES	YES	c.5439 G>T p.E1813D (39%)	WT	None
KI 5	MFPTC	F	44	2.8	Macrofollicular	YES	YES	c.5428 G>T p.E1810Y (40%)	WT	None
KI 6	DHGTC	F	20	6.5	Microfollicular	NO	NO	c.5437 G>A p.E1813K (27%)	WT	None
KI 7	MFPTC	F	27	2.5	Macrofollicular	YES	YES	c.5126 A>G p.D1709G (11%)	WT	None
KI 8	FTA	F	31	4	Macrofollicular	YES	YES	c.5428 G>T p.E1810Y (42%)	WT	None
KI 9	FTA	F	51	4.2	Macrofollicular	NO	YES	c.5437 G>C p.E1813Q (34%)	WT	None
KI 10	DHGTC	F	17	2.5	Microfollicular	NO	YES	c.5437 G>A p.E1813K (57%)	NP	None
KI 11	FTA	F	30	3.5	Macrofollicular	NO	YES	c.5126 A>G p.D1709G (43%)	WT	None
KI 12	PDTC	M	47	2.5	Trabecular, solid	NO	NO	c.5126 A>T p.D1709V (69%)	WT	None
KI 13	MFPTC	F	27	4.5	Macrofollicular	YES	YES	c.5437 G>C p.E1813Q (25%)	WT	SMA RCB1 p.F25L (46%)
KI 14	TFND	F	41	4.5	Macrofollicular	YES	NO	c.5126 A>G p.D1709G (39%)	NP	None
KI 15	MFPTC	F	38	2.8	Macrofollicular	YES	YES	c.5126 A>G p.D1709G (17%)	NP	PTPN11 p.A72T (11%)
KI 16	TFND	F	23	3.2	Macrofollicular	YES	NO	c.5126 A>G p.D1709G (37%)	NP	KRAS p.Q61L (24%)

Condello V et al. Atrophic changes in thyroid tumors are strong indicators of underlying DICER1 mutations: a bi-institutional genotype-phenotype correlation study. Virchows Arch. 2024 Jul;485(1):105-114

Mutation in 3/15 cases with germline DICER1 testing (20%): 2 minimally invasive follicular carcinoma, 1 follicular nodular disease



Specificity	Phenotypes
High-specificity (much more likely than not to have germline P/LP DICER1)	Pleuropulmonary blastoma (PPB) (including type 1r)
	Pituitary blastoma
	Anaplastic renal sarcoma
	Ciliary body medulloepithelioma
	Cystic nephroma (<18 yrs)
Moderate-specificity (more likely than not to have germline P/LP DICER1)	Embryonal rhabdomyosarcoma (ovarian)
	Embryonal rhabdomyosarcoma (cervix)
	Differentiated thyroid cancer and/or multinodular goiter (<18 years)
	Nasal chondromesenchymal hamartoma
Low-specificity (less likely to have DICER1)	Ovarian Sertoli-Leydig cell tumors
	Ovarian sex-cord stromal tumor of mixed type (specifically, gynandroblastoma)
	Nonparasitic liver cysts (childhood)
	Wilms tumor
	Pineoblastoma
	Cerebral sarcoma
	Lung cysts (<18 yrs)

Hatton JN et al. Specifications of the ACMG/AMP Variant Classification Guidelines for Germline DICER1 Variant Curation. Hum Mutat. 2023;2023:9537832

FIGURE 1 | Schematic representation of the method used to obtain results.

DICER1-Mutated tumors and DICER1 Syndrome

Congenital nodular

Nodular goitre

Table 2 Clinical and genetic characteristics of all *DICER1*-mutated cases

Case ID	Diagnosis	Sex	Age, years	Nodule size, cm	Architectural pattern	Papillary formations	Atrophic changes	<i>DICER1</i> hotspot mutation	Constitutional <i>DICER1</i> mutation (A/E)	Other mutations apart from <i>DICER1</i> in thyroid lesions (A/E)
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Should patients with *DICER1* mutated thyroid nodules discovered incidentally, undergo clinical genetics evaluation to rule out *DICER1* syndrome?

- ✓ As more and more cases undergo molecular profiling, more and more *DICER1* mutated thyroid nodules are being discovered...
- **Ideally yes** (germline mutations can be transmitted to progeny!), although germline mutations are uncommon in patients with unremarkable medical history
- Germline testing and genetic counselling [Altaraihi M et al. Prevalence of Pathogenic Germline *DICER1* Variants in Young Individuals Thyroidectomised Due to Goitre - A National Danish Cohort. *Front Endocrinol (Lausanne)*. 2021;12:727970. doi: 10.3389/fendo.2021.727970]
 - Patients thyroidectomised for goiter aged <21 years
 - Patients thyroidectomised for goitre aged <25 years + family history of goiter
 - Patients of all ages thyroidectomised for goitre, who are affected by another *DICER1* manifestation

nodular goitre

N=111

N=24

N=2 (8.3%)

Low-specificity (less likely to have *DICER1*)

Wilms tumor
Pineoblastoma
Cerebral sarcoma
Lung cysts (<18 yrs)

Hatton JN et al. Specifications of the ACMG/AMP Variant Classification Guidelines for Germline *DICER1* Variant Curation. *Hum Mutat*. 2023;2023:9537832

FIGURE 1 | Schematic representation of the method used to obtain results.