

Papillary thyroid carcinoma as first and isolated neoplastic disease in a Lynch syndrome family member with a germline MLH1 mutation

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Abstract

The Lynch syndrome (LS) is an autosomal dominant disorder characterized by a strongly increased risk of developing colorectal cancer and several extra-colonic malignancies, such as carcinomas of the endometrium, ovary, ureter, stomach, and small intestine [1]. Lynch syndrome is caused by germline mutations in mismatch repair genes (MMR)[2], mainly in *MLH1* and *MSH2*, rarely in *MSH6* and *PMS2* [3,4]. Tumors usually develop at a relatively young age (<50 years). Some cancers, rare in this syndrome, can be incidentally diagnosed in one of the Lynch syndrome family members. Here we report the case of unusual presentation of papillary thyroid carcinoma in a young woman carrying the c.545+3A>G mutation (rs267607760) in *MLH1* gene.

Case History

A 33-year-old woman was referred to our endocrine unit in 2017 because of the incidental finding of anti-thyroperoxidase autoantibodies presence. Her family was affected by Lynch Syndrome: the father underwent surgery for colon cancer before 50 years of age and later for squamous cell carcinoma of the skin and duodenum cancer, two aunts underwent surgery for colorectal and endometrial cancers before age of 50 and one aunt and her sister underwent surgery for endometrial cancer. For this reason, the family was previously submitted to genetic analysis, which found the presence of the same germinal mutation c.545+3>G of *MLH1* gene in our patient and 6 first degree relatives thus confirming the Lynch syndrome diagnosis (Supplementary figure 1).

The patient was euthyroid at the first observation and the ultrasound imaging of the neck showed a hypoechoic nodule (5x6x11mm) in the left lobe of the thyroid gland. Due to the presence of the germline *MLH1* mutation, we decided to perform a fine needle aspiration biopsy of this nodule, and the cytology identified a thyroid carcinoma (TIR5).

The patient underwent total thyroidectomy and pathology confirmed the presence of a classical variant of papillary thyroid carcinoma, with the thyroid capsule infiltration (T1a, N0, Mx). The genetic analysis of the neoplastic thyroid tissue was performed by targeted next-generation sequencing (NGS) using two custom thyroid cancer-specific multi-gene panels that target single-nucleotide variants/small indels (DNA panel) and gene fusions (RNA panel), as previously described [5,6]. The analysis showed the presence of both *BRAF* p.V600E and *TERT* c.-124 C>T (also known as C228T) promoter mutations with an allelic frequency of 10% and 16%, respectively. Radioiodine ablation (1850 MBq) after 0.9 mg of rhTSH was performed and whole-body scan (WBS) post- ¹³¹I administration showed only a thyroid bed uptake, without other sites of pathological uptake. Stimulated thyroglobulin (Tg) was 0.6 ng/mL (cut-off <0.1 ng/ml), Tg autoantibodies (TgAb) were 145 UI/mL (0-60 UI/mL).

Tg is currently <0.1 ng/mL, TgAb are 15 UI/mL. Ultrasound imaging of the neck is negative for the presence of suspicious lymph nodes. No other neoplasms have been diagnosed during a four-year follow-up.

Discussion

Lynch syndrome diagnosis is based on the application of Amsterdam criteria, which applies the “3-2-1 Rule”, and the Amsterdam second classification, which includes non-colorectal cancers: endometrial, small intestine, ureteral, and kidney cancers [7]. The “3-2-1 Rule” criteria include ≥ 3 affected family members, one of whom is a first-degree relative of the other two, across at least 2 generations with at least 1 of them presenting with hereditary non polyposis colorectal cancer (HNPCC) before the age of 50. The familial adenomatous polyposis should be excluded. Lynch syndrome is caused by germline mutations in one of four DNA mismatch repair genes that lead to microsatellite instability and/or loss of mismatch repair protein expression at immunohistochemistry [1, 2]. In 90% of cases, mutations can be found in *MLH1* or *MSH2* genes, while mutations in *MSH6* and *PMS2* genes are less frequent [1, 2].

The etiology of tumors that are usually rare in the HNPCC tumor spectrum is still controversial. Here, we report the case of a 33-year-old woman with a family history of Lynch syndrome, due to a germline *MLH1* gene mutation, who developed a papillary thyroid carcinoma.

Only a few cases of thyroid cancer have been reported in patients with Lynch syndrome, mostly associated with an *MSH2* germline mutation (Table 1). Interestingly, Aswath K et al., described a family with the co-occurrence of familial non-medullary thyroid cancer (FNMTc) and HNPCC. Two out of five relatives affected by PTCs were analyzed and both carried the germline *MSH2* p.C707Y variation[8]. Although the clinical significance of the *MSH2* variant is uncertain, it may suggest a role in PTC predisposition.

To our knowledge, only one thyroid cancer associated with a germline *MLH1* mutation has been previously described in a patient with Lynch syndrome. The patient was a 43-year-old female who developed PTC a few years after the uterine and ovarian adenocarcinomas and colon cancer. Unfortunately, we have no data about the genetic analysis on thyroid cancer tissue[9].

Here, we report a second patient carrying a germline *MLH1* mutation. The *MLH1* c.545+3A>G variation is known to be pathogenic leading to the skipping of exon 6. The genetic analysis of the thyroid cancer tissue showed the coexistence of BRAF p.V600E and *TERT* C228T promoter mutations. Notably, this is the first case in whom papillary thyroid carcinoma has been diagnosed as the first and isolated neoplastic disease in a Lynch syndrome family member. During the 4 years follow-up the patient didn't develop any Lynch-associated cancers.

Genes involved in DNA repair pathways have a pivotal role in preserving genomic integrity upon exposure to genotoxic agents and the impairments in these mechanisms are related to increases in mutation frequency, genomic instability, and finally to cancer susceptibility[10,11].

Although thyroid cancer is not considered as part of Lynch syndrome, several reports have indicated an association between MMR deficiency and thyroid cancer.

Lu Y et al. found a lower expression of MLH1 protein in thyroid cancer tissues, both papillary and follicular cancers, compared with normal tissues [12]. Moreover, *MLH1* germline and somatic mutations have been described in PTCs [10, 12].

The defective DNA repair system in a condition of excessive oxidative stress that promotes the DNA oxidation and subsequent lesions may have an important role in the PTC tumorigenesis [9, 13]. It is therefore reasonable to suppose that MMR impairment, including that induced by *MLH1* mutations, may increase the chance of somatic genetic alterations. Our patient follows this hypothesis being a carrier of two somatic driver mutations in the PTC tissue. Moreover, in all analyzed PTC tissues of patients with Lynch Syndrome, including our patient, it has been found a somatic BRAF p.V600E mutation (Table 1). Interestingly, the association between the aberrant *MLH1* methylation and expression and the presence of the BRAF p.V600E mutation has been described in thyroid cancer cell lines and PTC tissues [10].

In conclusion, although we cannot exclude an incidental occurrence of PTC in Lynch syndrome patients, our study underlines the need to follow-up LS patients for the development of rare associated cancers, including thyroid cancers.

Declarations

Compliance with ethical standard

Conflict of interest

The authors declare that they have no conflict of interest

Informed consent

Patient gave her informed consent.

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Tables

Table 1: Clinical and genetic features of Lynch-syndrome patients who developed thyroid cancers

# Lynch syndrome families	Germline mutations in MMR genes	Gender	Tumor type (age at diagnosis)	Thyroid cancer TNM	Somatic mutations in thyroid cancer tissue	References
1^	MSH2 p.C707Y	F	Colon polyps (20)	T4N1bM0	BRAF p.V600E	[8]
			Papillary thyroid carcinoma (53)			
			Melanoma (63)			
		F	Papillary thyroid carcinoma (33)	T1aN0M0	BRAF p.V600E	
2	MSH2 p.Q824X	F	Anaplastic thyroid cancer (39)	na	na	[4]
3	MSH2 c.1704_1705delAG	F	Colorectal adenoma (44)	na	na	[3]
			Anaplastic thyroid cancer (44)			
4	MSH2 c.906_907insT	M	Colon cancer (21)	pT4pN1b	na	[9]
			Papillary thyroid carcinoma (34)			
5	MSH2 na	M	Colon adenocarcinoma (33)	na	BRAF p.V600E	[15]
			Salivary gland adenocarcinoma (44)			
			Papillary thyroid carcinoma (44)			
			Parotid acinic cell carcinoma (44)			
9	MSH2 c.2646delA	F	thyroid cancer ^S (37)	na	na	[16]
6	MSH6 p.G1105fs*3	F	Colorectal adenocarcinoma (54)	pT4bN1b	AKT1 p.E17K; ARID1A p.I106fs*4; ATM p.R3008H; PI3KCA p.R38C;	[17]
			Anaplastic thyroid cancer (54)			

PTEN
p.R335*;
RB1 c.265-
1G > T;
TP53
p.R196*

7	MLH1 c.1858G>T	F	Uterine adenocarcinoma (43) Ovarian adenocarcinoma (43) Colon cancer (45) Papillary thyroid carcinoma (47)	pT1aN1a	na	[9]
8	MLH1 c.545+3A>G	F	Papillary thyroid carcinoma (36)	T1aN0Mx	BRAF p.V600E + TERT C228T	this paper

na= not available

^ Family with HNPCC and FNMTC

§ Not specified

Supplementary Files

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