

Thyroid cancer in a patient with Lynch syndrome – case report and literature review

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Abstract: Lynch syndrome describes a familial cancer syndrome comprising germline mutations in one of four DNA mismatch repair genes, *MLH1*, *MSH2*, *MSH6*, and *PMS2* and is characterized by colorectal, endometrial, and other epithelial malignancies. Thyroid cancer is not usually considered to be part of the constellation of Lynch syndrome cancers nor have Lynch syndrome tumor gene mutations been reported in thyroid malignancies. This study reports a woman with Lynch syndrome (colonic cancer and a DNA mismatch repair mutation in the *MSH2* gene) with a synchronous papillary thyroid cancer. Six years later, she developed metachronous breast cancer. Metastatic bone disease developed after 3 years, and the disease burden was due to both breast and thyroid diseases. Despite multiple interventions for both metastatic breast and thyroid diseases, the patient's metastatic burden progressed and she died of leptomeningeal metastatic disease. Two prior case reports suggested thyroid cancer may be an extraintestinal malignancy of the Lynch syndrome cancer group. Hence, this study examined the genetic relationship between the patient's known Lynch syndrome and her thyroid cancer. The thyroid cancer tissue showed normal expression of *MSH2*, suggesting that the tumor was not due to the oncogenic mutation of Lynch syndrome, and molecular analysis confirmed *BRAF V600E* mutation. Although in this case the thyroid cancer was sporadic, it raises the importance of considering cancer genetics in familial cancer syndromes when other cancers do not fit the criteria of the syndrome. Careful documentation of other malignancies in patients with thyroid cancer and their families would assist in better understanding of any potential association. Appropriate genetic testing will clarify whether a common pathogenic mechanism links seemingly unrelated cancers.

Keywords: cancer genetics, familial cancer syndromes, thyroid and hereditary non-polyposis colorectal cancer

Introduction

Lynch syndrome (hereditary non-polyposis cancer) is an autosomal dominant disorder associated with germline mutations in one of four DNA mismatch repair genes. Approximately 90% of mutations are located in *MLH1* or *MSH2*, the remainder in *MSH6* and *PMS2*.^{1,2} Lynch syndrome carries increased risk of colorectal cancer, endometrial cancer and other epithelial malignancies.¹

Lynch syndrome classifications do not include thyroid cancer. The Amsterdam criteria apply the "3-2-1 Rule": three relatives with colorectal cancer, one a first degree relative of the other two; two successive generations affected; one diagnosed before the age of 50 years. The expanded Amsterdam II classification includes non-colorectal cancers: endometrial, small intestine, ureteric, or renal cancers.³

This study reports a female with Lynch syndrome-related colon cancer that developed synchronous thyroid cancer and 6 years later breast cancer with progression to metastatic disease due to thyroid and breast cancer burden. The aim of this study was

As some of the metastatic burden was thyroid cancer related, 5.6 GBq I¹³¹ was administered; the unstimulated thyroglobulin level after 4 weeks was 221 µg/L (Figure 1A). The tyrosine kinase inhibitor sorafenib was then commenced. The patient died after 3 months from disseminated leptomeningeal metastases.

Discussion

This study presents a fascinating case of a young woman with Lynch syndrome-related colonic cancer with synchronous papillary thyroid cancer and metachronous breast cancer with progression to metastatic disease related to both thyroid and breast cancers.

Thyroid cancer generally has an excellent long-term prognosis. Mutations in *BRAF*, *RET/PTC*, *NTRK1*, or *RAS* have been reported in up to 70% of differentiated thyroid cancers.⁵ *BRAF* mutation confers a more aggressive clinical course, as evident in this case.

This is the third case of thyroid cancer described in Lynch syndrome. The first case was a female from a Lynch syndrome pedigree with a germline *MSH2* mutation who developed an undifferentiated thyroid cancer as the index cancer. Immunohistochemistry showed that *MSH2* expression was absent.¹ The second case was a female from a Lynch syndrome pedigree with anaplastic thyroid cancer as the index cancer. Immunohistochemistry also showed that *MSH2* expression was absent.² Genetic analysis confirmed a germline mutation in case 1 and in family members of case 2.

The present case demonstrated loss of *MSH2* and *MSH6* staining in her colonic cancer only. Lynch syndrome was confirmed by DNA sequencing of *MSH2* that revealed a c.998G>A missense change in exon 6. In this case, the *PTC* displayed normal mismatch repair protein expression and showed a *BRAF* mutation, making it unlikely for mismatch repair defects due to the cancer genetics of Lynch syndrome. *MSH2* promoter hypermethylation has been shown to occur in colorectal cancers related to Lynch syndrome, with somatic inactivation of the remaining wild-type allele.⁶

Second, primary tumors, synchronous or metachronous, have been reported in literature.⁷ It is hypothesized that the synchronous occurrence of two tumors may be a result of a mutagenic trigger that causes independent genetic events⁷ and that metachronous tumors may be due to persisting effects of genetic and behavioral risk factors.⁷ It is possible the thyroid gland may be activated by prior hormone-secreting cancers with overstimulation in a reactive autoimmune environment, as hypothesized by Sisti et al.⁸ Furthermore, a history of thyroid cancer is associated with the increased

risk of developing breast cancer, and several studies have evaluated the thyroid hormone and proposed mechanisms for activation of the associated oncogene.⁹ The most likely explanation for synchronous cancers in the present case seems to be a sporadic thyroid cancer on a background of genetic cancer predisposition.

It should be noted that normal immunohistochemistry does not completely exclude the mismatch repair defects that characterize Lynch syndrome. False negatives may occur due to the following reasons: First, the effectiveness of immunohistochemistry in revealing germline mismatch repair mutations depends on the number of antibodies used. Mismatch repair proteins form heterodimers to maintain function: *MSH2* with *MSH6*; *MHL2* with *PMS2*. Immunohistochemistry with *MLH1/MSH2* antibodies has a sensitivity of 85% in detecting a mismatch repair mutation; sensitivity is increased to 92% using *PMS2* and *MSH6* antibodies.¹⁰ Furthermore, if immunohistochemistry is performed with *MLH1/MSH2* antibodies only, some *MLH1* and *MSH2* abnormalities may be missed, as some pathogenic mutations retain protein expression. Second, not all pathogenic mutations result in loss of protein by immunohistochemistry. Immunohistochemistry is effective in screening for mutations due to a truncated or degraded protein, but performs poorly in differentiating abnormal protein due to a missense mutation from normal protein. Greater than 30% of *MLH1* mutations are missense mutations that may not be detected by immunohistochemistry. Third, there may be a misleadingly false normal staining pattern due to a second hit that inactivates the wild-type allele resulting in non-functional protein but allowing normal antibody binding.¹⁰ Immunohistochemistry with a four antibody panel and microsatellite instability (MSI) testing have similar sensitivity for detecting mismatch repair mutations; however, each may miss cases detectable by the other method. Therefore, if high clinical suspicion synchronous cancers may be linked to the same pathogenic process, normal immunohistochemistry result should be verified with MSI testing.

These few cases raise the importance of considering individual cancer genetics in familial cancer syndromes, particularly when other cancers do not fit the described syndrome. Careful documentation of other malignancies in patients with thyroid cancer and their families would assist in better understanding any potential association of thyroid cancer to known familial cancer syndromes. Appropriate genetic testing of the cancers will clarify whether common genetics explain synchronous or subsequent malignancies in individuals and their families and add to greater

understanding of the genetic mechanisms that may explain apparently unrelated cancers.

Disclosure

The authors report no conflicts of interest in this work.

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