

Case Report

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Colon cancer in a 12-year-old girl with hypertriglyceridemia

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Abstract

Colorectal cancer is usually considered a disease of the elderly; however, in a small fraction of patients (2%-3% of all affected individuals), colorectal malignancies may develop earlier. The reasons whereby some individuals develop colorectal cancer at a young age are poorly understood. In a 12-year-old girl, a malignancy was diagnosed in the ascending colon. There was no familial history of Lynch syndrome or familial adenomatous polyposis. The metabolic profile of the patient revealed hypertriglyceridemia and low high-density lipoprotein cholesterol levels at nine years, then diagnosed as familial hypertriglyceridemia due to a constitutional mutation in the *APOA5* gene (c.427delC). Moreover, variants possibly increasing the risk of cancer were detected in *MSH6* (c.3438+11_3438+14delCTTA, intron 5) and *APC* (I1307K). The patient showed a rather unusual dietary pattern, since her basic alimentation from weaning consisted almost exclusively of meat homogenates and, subsequently, roasted meat or cutlets. Other foods, including fish, vegetables, sweets, and pasta, were refused. In this case, genetic and environmental factors could have acted in a particularly accelerated manner. Indeed, the genetic background of the patient (familial hypertriglyceridemia and polymorphisms predisposing to colorectal cancer) may have favored a dietary-driven colorectal carcinogenesis, resulting in an extremely early onset development of



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malignancy.

Keywords: Colorectal cancer, hypertriglyceridemia, *APOA5*, *MSH6*, *APC*, case report

INTRODUCTION

Colorectal cancer is common in elderly individuals but rather infrequent under the age of 50 years, although recent reports show that the incidence is increasing in young people in many countries^[1]; the disease may be considered a rarity in the first and second decades of life^[2]. While approximately 3% of all colorectal malignancies show a genetic origin, the fraction raises to 20%-30% among people who develop the tumor before the age of 35-40 years^[3,4]. It follows that in early or very early onset colorectal cancer a possible genetic background, in particular Lynch syndrome or familial adenomatous polyposis, should always be taken into consideration^[5,6]. However, the large majority of colorectal neoplasms developing in young individuals are not clearly associated with hereditary factors - at least within the limits of our present knowledge - and their etiology remains undetermined.

We describe the unusual case of a 12-year-old girl in whom a Dukes C (T4N1M0) carcinoma of the ascending colon developed in the absence of a documented genetic syndrome or a background of polyposis. Rather unexpectedly, the metabolic profile of the patient (familial hypertriglyceridemia with evidence of a known heterozygous frameshift mutation in the *APOA5* gene), and possibly predisposing variants in the *MSH6* and *APC* genes, together with an unusual dietary pattern (excessive consumption of meat) may have been the basis of an advanced malignant tumor before adolescence.

CASE REPORT

The index case was born in 2004 with a caesarean delivery (weight at birth 3150 g). Since then, physical and mental development appeared in the normal range; she entered school at age 5 and attended regular courses. At age 9, she was diagnosed with hypertriglyceridemia and low high-density lipoprotein (HDL) cholesterol levels [Table 1], which was treated irregularly with polyunsaturated fatty acids. The youngest sister, the father, and the paternal grandfather showed a similar lipid profile. At age 10, she underwent appendectomy (2014).

In October 2015, she started to complain of fatigue, palpitations, and mild abdominal pain. After almost one year, blood tests revealed severe anemia (Hb: 6.0 g/dL), with low serum iron levels, and a fecal occult blood test was positive. In September 2016, at age 12 and 7 months, a colonoscopy showed a large and substenosing tumor of the ascending colon. A PET (positron emission tomography) scan revealed accumulation of fluorodeoxyglucose in the right colon and in two small pelvic nodules. Two cycles of chemotherapy (FOLFOXIRI, folinic acid, 5-fluorouracil, oxaliplatin, and irinotecan) were executed before surgery. The patient underwent a right hemicolectomy in November 2016. At anatomical examination, a substenosing lesion of 1.5 cm was observed in the right colon. Histologic diagnosis was poorly differentiated adenocarcinoma - with aspects of signet ring cell carcinoma - with tumor budding, neural and vascular invasion, and metastasis in 3 of 57 lymph nodes (T4N1Mx). Two 8-10 mm peritoneal nodules resected with the colon showed fibrotic tissue free of malignant cells. The patient recovered promptly and had her menarche two months after surgery; she underwent eight more cycles of chemotherapy (FOLFOX, folinic acid, 5-fluorouracil, and oxaliplatin) in 2017. In July 2017, the patient was in good clinical condition and well developed. Her height was 149 cm and weight was 48 kg. She had a disease relapse in November 2017 (a 13 cm mass of the ovary and four peritoneal masses), so she underwent ovariectomy, peritonectomy, and intraperitoneal hyperthermic antineoplastic perfusion with cisplatin and mitomycin. In November 2020, she

Table 1. Serum lipid pattern over time of the four family members with hypertriglyceridemia

	Calendar year						
	2007	2011	2012	2013	2014	2015	2017
Grandfather I-1							
Triglycerides (mg/dL)	439						
Total cholesterol (mg/dL)	185						
LDL cholesterol (mg/dL)	-						
HDL cholesterol (mg/dL)	23						
Father II-1							
Triglycerides (mg/dL)	405	711					232
Total cholesterol (mg/dL)	157	247					194
LDL cholesterol (mg/dL)	-	-					134
HDL cholesterol (mg/dL)	23	25					34
Patient (proband) III-1							
Triglycerides (mg/dL)			114	198	333	179	332
Total cholesterol (mg/dL)			205	195	208	-	241
LDL cholesterol (mg/dL)			147	-	151	-	154
HDL cholesterol (mg/dL)			33	24	31	-	38
Sister III-3							
Triglycerides (mg/dL)			284		123	214	66
Total cholesterol (mg/dL)			169		232	211	190
LDL cholesterol (mg/dL)			-		164	147	130
HDL cholesterol (mg/dL)			19		43	-	48

had a new relapse near the colonic anastomosis and a suprapubic nodule in the rectious muscles of the abdomen, which were surgically removed in January 2021, along with a new peritonectomy with intraperitoneal hyperthermic perfusion of cisplatin and mitomycin. Then, she underwent six more cycles of FOLFOXIRI (folinic acid, 5-fluorouracil, oxaliplatin, and irinotecan) plus bevacizumab. As of November 2021, she is in a stable condition, and she will be clinically re-evaluated in February 2022.

Colorectal cancer molecular studies

From the genealogic tree of the family [Figure 1], no pattern of vertical transmission of cancer was clearly evident, and the proband was the only case in the family. As a matter of fact, in rare cases, Lynch syndrome may initiate with a new mutation, especially in a young patient, with neoplasms of the proximal colon and histologic pattern of mucinous or signed ring cell carcinoma^[7]. Moreover, by endoscopy and anatomical examination of the resected colon, no polyps or background of diffuse polyposis could be observed. Similarly, features of Peutz-Jeghers or Cowden disease^[8,9] were absent in the proband and in her first-degree relatives.

However, because of the unusual case, the possible genetic origin of the tumor was studied using next generation sequencing (NGS) techniques. In particular, a TruSeq Custom Amplicon workflow was applied in searching for constitutional sequence variants in genes which increase the risk of colorectal cancer. The panel included all exons and ± 20 padding intronic regions of the following genes: *APC*, *MUYH*, *MLH1*, *MSH2*, *MSH6*, *PMS2*, *EPCAM*, *STK11*, *PTEN*, *SMAD4*, and *BMP1A*. Variants with global minor allele frequency of < 1% were confirmed by Sanger sequencing^[10,11].

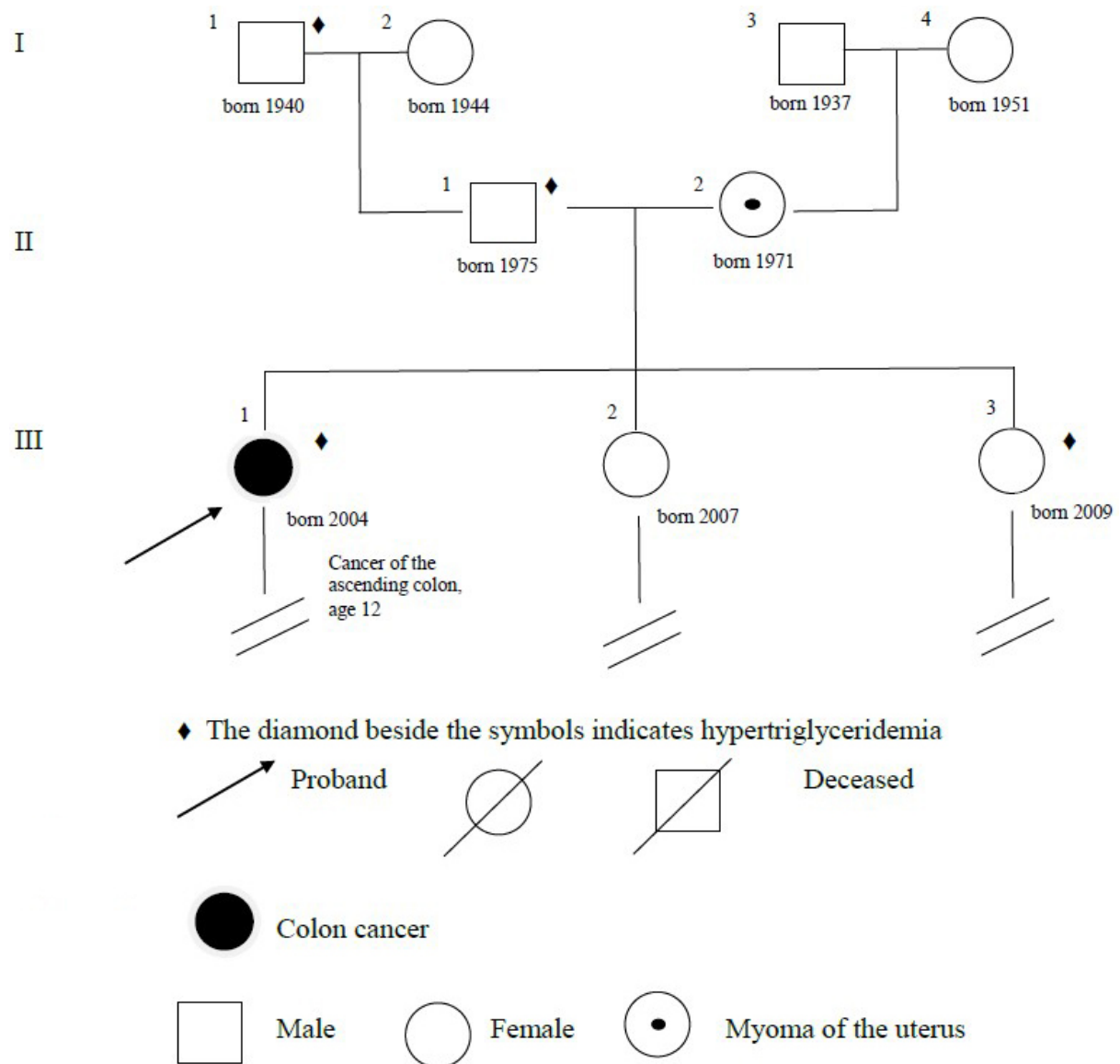


Figure 1. Genealogical tree of the family.

The NGS study revealed wild-type sequences and no large gene deletion/duplications. A four-base deletion at intron 5 in the *MSH6* gene was detected (c.3438+11_3438+14delCTTA). This constitutional alteration is considered as a variant of uncertain significance, according to the international database (InSight Variant Database and Universal Mutation Database), not clearly associated with Lynch families or segregating in kindreds. Moreover, the missense variant c.3920 T>A (p.Ile1307Lys, also known as I1307K) was detected in the *APC* gene [Figure 2]; this is a low-penetrance, relatively frequent (especially in Ashkenazi Jews) mutation associated with an increased risk of colorectal cancer, but with no definite role in hereditary colorectal cancer and familial adenomatous polyposis^[12,13]. In addition, loss of heterozygosity studies in the tumor tissue showed that the variant maintained its heterozygous state. The immunohistochemical evaluation of the mismatch repair proteins showed normal expression of the four main proteins (MLH1, MSH2, MSH6, and PMS2) in the resected tumor [Figure 3]^[14].

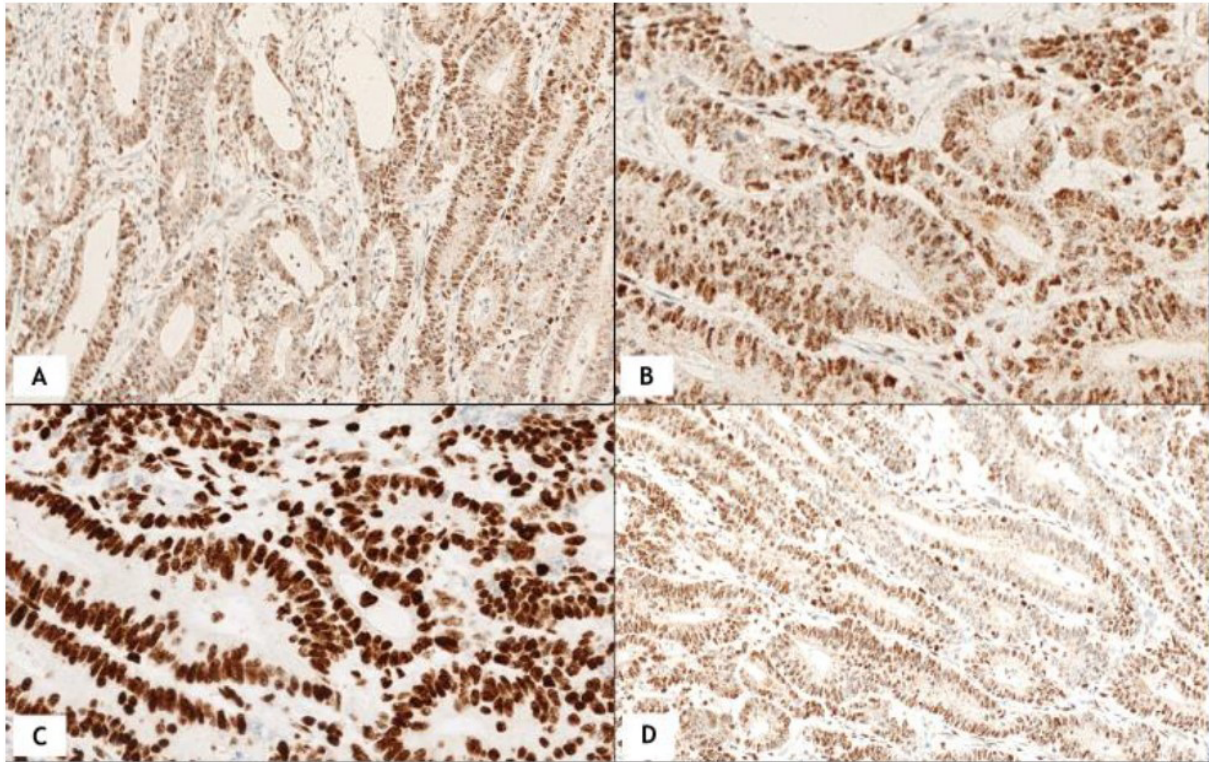


Figure 3. Typical immunohistochemical nuclear staining of the four mismatch-repair proteins in tumor cells of the carcinoma of the patient: MLH1 (A); PMS2 (B); MSH2 (C); and MSH6 (D). Magnification is 40 \times .

refused by the baby; when the mother insisted on forcing the proband to eat these foods, they were regularly vomited. Fish was never eaten because it was refused; similarly, at variance with other children, sweets and ice cream were not accepted. Meat was always cooked in the same way: roasted with some olive oil or dipped in bread crumbs, with some salt.

This meat-based dietary pattern was maintained up to the end of 2016 (and, thus, to the development of symptoms). In the last years, after the first surgical intervention, the continuous efforts of the mother and the suggestions of the medical personnel led to a change of the dietary habits, increasing the consumption of pasta and vegetables. The other members of the family, including those with high triglyceride levels, regularly consumed any kind of food, without restriction of dietary fat.

Family history and lipid disorders

The genealogical tree of the nuclear family is depicted in [Figure 1](#). No other malignancy was reported among first-degree relatives and grandparents. The mother of the proband (II-2) underwent hysterectomy at age 38 because of myoma of the uterus. The father, born in 1975, had acute pancreatitis in 2011, at age 36; on that occasion, overweight-associated non-alcoholic fatty liver and hypertension were also detected. Four years earlier, in 2007, hypertriglyceridemia was diagnosed; blood tests were executed because of the known high lipid levels of his father (the paternal grandfather of the proband), who referred triglycerides values in the order of 500-1000 mg/dL. Hypertriglyceridemia was also diagnosed in the proband (age 9) and in one of her sisters (III-3 in the family tree, age 3). Thus, familial high triglyceride levels were present in four patients and in three consecutive generations. Moreover, a lifelong history of overweight was present in the father and in the younger sister of the proband.

Table 1 summarizes the serum lipid levels that we could document in the four members of the family affected by hypertriglyceridemia. Fluctuations of values could depend on dietary recommendations and therapy (fibrates and/or fish oil). Total and LDL cholesterol were also moderately elevated in most samples, while HDL levels were constantly below the normal range.

NGS exome analysis using the Ion Torrent Platform^[15] revealed the presence of a heterozygous constitutional mutation in the *APOA5* gene in the proband and her relatives (I-1, II-1, and III-3) affected by hypertriglyceridemia. The variant (c.427delC) was a frameshift mutation causing a stop codon downstream and, thus, a truncated protein (p.Arg143Alafs*57)^[16]. The deleterious *APOA5* mutation was described previously by Evans *et al.*^[16] and was associated with moderate-to-severe hypertriglyceridemia including in heterozygous carriers.

DISCUSSION AND CONCLUSIONS

We describe the case of a girl who had cancer of the large bowel very early in her life. Our initial hypothesis was that inheritance could play a determinant role in inducing the development of malignancy at 12 years of age; however, family history and molecular studies apparently excluded, or rendered extremely improbable, the role of genetics in this case. As a matter of fact, there was no family history of colorectal cancer or polyposis, immunohistochemistry of the main DNA mismatch repair proteins (*MLH1*, *MSH2*, *MSH6*, and *PMS2*) revealed a normal expression, and no truncating mutation in 11 cancer-related genes could be detected. The variants observed in *MSH6* and *APC*, although possibly linked to an increased risk of tumors, were not associated with very early colorectal cancer occurrence, and they can hardly be invoked as the main factors for explaining the observed case.

We therefore turned our attention to hypertriglyceridemia, which ran in the family for at least three generations (including the proband), and to the peculiar alimentation of the index case, which consisted from birth almost exclusively of meat. High triglycerides levels and meat consumption have been associated with the development of colorectal cancer.

Evidence linking triglyceride levels to tumors in general is rather consistent. McKeown-Eyssen^[17] gathered epidemiological evidence that hypertriglyceridemia was associated with the risk of colorectal cancer. Among possible explanations, the author suggested an association of triglycerides with fecal bile acids (which promote colorectal carcinogenesis), increased insulin levels, and the fact that triglycerides might be indicators of energy available through the circulation for tumor cells. Other studies reported that high serum triglycerides levels were associated with an increased risk of colonic or colorectal adenomas^[18-20]. In a large group of patients with prostate cancer, Hayashi *et al.*^[21] reported a correlation between triglyceride levels and incidence of the disease, especially in patients over the age of 60 years, and concluded that individuals with high triglyceride levels may be more vulnerable to prostate cancer. Among possible explanations, the authors suggested insulin resistance - which is closely associated with obesity and high lipid levels - inflammatory state, and oxidative stress. Similar results were reported in a recent study by Allott *et al.*^[22], in which the authors evaluated recurrence of prostate cancer after surgery. In a large study including 31,000 women recruited between 1995 and 1997, Lindemann *et al.*^[23] found a significant positive association between serum triglyceride levels with risk of endometrial cancer; the authors suggested that some of the effects related to triglycerides could be mediated by body mass. Since adipose tissue is a source of endogenous estrogens, unopposed estrogen may play a relevant role in the occurrence of endometrial cancer. Moreover, a recent meta-analysis reported that serum triglycerides levels are higher in premenopausal women with breast cancer^[24]. To summarize, high triglyceride levels are closely associated with obesity and insulin resistance, which in turn induce a proinflammatory state that favors cell

replication, activation of mutagens (products of oxidative stress), and ultimately cancer^[25]. In accordance with this hypothesis and sequence of events, Otani *et al.*^[26] found that plasma levels of C-reactive protein - a marker of inflammation - were associated with a subsequent risk of colorectal cancer.

In our study, the proband and three of the relatives affected by hypertriglyceridemia were heterozygous carriers of a frameshift mutation in the *APOA5* gene, which is associated with moderately to severely increased triglycerides levels^[16]. Indeed, several genetic studies have shown a correlation between *APOA5* variants and triglycerides^[27,28]. In addition, *APOA5* was associated with insulin resistance and metabolic syndrome^[29], two conditions whose association with systemic inflammation - potentially leading to cardiovascular diseases and neoplasm - has been documented.

The ingestion of meat has been associated with colorectal cancer in epidemiological studies; however, the strength of the association and types of meat involved were not always consistent. Studies on red meat consumption and tumors of the large bowel showed odd ratios in the order of 1.2-1.3^[30,31]. Cotterchio *et al.*^[32], however, reported odd ratios of 1.4-1.7 and suggested that the cancer risk may be further elevated in individuals with some variants (*CYP1B1* and *SULT1A1*) of the enzymes involved in carcinogen metabolism. Moreover, Ferrucci *et al.*^[33] found that red meat and pan-fried meat were associated with increased risks (1.7-2.0) of colorectal adenoma. In addition, Pan *et al.*^[34] found evidence that consumption of red meat was significantly linked with an increased risk of total, cardiovascular, and especially cancer-related mortality.

The reasons whereby meat consumption increases the risk of some types of cancer, in particular large bowel cancer, remain unclear, at the point that Forman^[35], in 1999, entitled his editorial on this topic as "Meat and cancer: a relation in search of mechanisms". Among the hypothetical mechanisms which might be invoked, we can mention: (1) the risk due to the carcinogenic polycyclic aromatic hydrocarbons produced when meat is cooked at high temperature^[36]; (2) the fact that cooked, pan-fried, salted, and processed meat might induce the formation of carcinogenic substances such as heterocyclic amines and other nitroso compounds^[37]; (3) a meat-based alimentation may induce changes of the intestinal bacterial flora, as well as the (possible) formation of bacterial metabolites with a known carcinogenic effect^[38]; and (4) iron and especially the heme iron present in red meat might promote colorectal cancer through the ability of iron to gain and lose electrons and, thus, to participate to potentially deleterious free radical-generating reactions^[39,40].

In conclusion, it is likely that both triglycerides and meat consumption play some role in colorectal cancer development, although controversies do exist in many investigations and reproducibility of results has been rather poor. If we turn our attention to the young girl described in the report, we can imagine that two common putative etiological factors - triglycerides and meat - which are usually diluted with many other factors in the general population in this specific case could exert their oncologic effect to the maximum level, starting from the birth - or very early childhood - and continuing for 12 consecutive years. In addition, the genetic variants detected in *MSH6* and *APC* colorectal cancer-associated genes might have rendered the patient more susceptible to cancer development.

This case emphasizes the possible role of metabolic factors and unbalanced diets in colorectal cancer occurrence. However, other factors and agents might have been involved or facilitated the role of triglycerides and meat, such as common polymorphisms associated with colorectal cancer risk^[41] or mutation of unknown (or unexplored) colorectal cancer-associated genes^[42].

DECLARATIONS

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Authors' contributions

Made a substantial contribution to acquisition of clinical data, analysis, drafting and review of the manuscript: Ponz de Leon M, Noto D, Roncucci L

Made a substantial contribution to genetic and clinical chemistry analyses, drafting, and review of the manuscript: Pedroni M, Viel A, Nascimbeni F, Sena P

Made a substantial contribution to pathological analyses: Reggiani Bonetti L

All authors have read and approved the final version of the manuscript.

Availability of data and materials

Data sharing is not applicable to this article as no datasets were generated or analysed during the current study. Personal, genetic, and clinical data cannot be disclosed, of course; however further de-identified information may be requested to the corresponding author.

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Conflicts of interest

All authors declared that there are no conflicts of interest.

Ethics approval and consent to participate

Written informed consent to undergo genetic testing, and to participate in this study was obtained from the parents of the child (under 16) described in this study. The research involved genetic testing of individuals as part of the routine genetic approach of subjects at increased risk for inherited oncologic diseases. Hence, the research does not necessitate approval from the ethics committee.

Consent for publication

Although no personally identifiable information is reported in this study, informed consent for publication was obtained from the parents of the child (under 16) reported in this study.

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