

Endoluminal tumor implant of a colorectal cancer in an anal fistula detected by FISH techniques: a case report

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Abstract: Intraluminal shedding of tumor cells is a rare infrequent sporadic colorectal cancer (sCRC) mechanism of spreading. Less than 30 cases of sCRC metastasis into anal fistula have been reported. Here, we study a 72-year-old male with an adenocarcinoma arising in an anal fistula. Subsequent studies revealed another tumor in the rectum without distant metastatic disease; therefore, a curative-intent abdominoperineal resection was performed. The histologic study showed a moderately differentiated adenocarcinoma in both locations. No perineural or lymphovascular invasion was observed, and all the lymphatic nodes resected were negative for malignancy. Both tumors showed positive CK20 and negative CK7 immunostaining, but KRAS G12D mutation was only detected in the rectal tumor. After those conventional studies, a cytogenetic profile of both tumors was performed by interphase fluorescence in situ hybridization (iFISH) techniques. The FISH study displayed an identical genetic profile in both tumors, loss of the chromosome 8 and 18q, and no alteration in chromosome 7 and 13q. Based on pathological and genetic findings, we established the same clonal origin of both tumors. Currently, the diagnosis of an intraluminal CRC metastasis relies on histologic and immunohistochemistry findings. We suggest that genetic studies at the individual cell level by FISH techniques may be useful in order to differentiate synchronous from intraluminal metastasis.

Keywords: Sporadic colorectal cancer (sCRC); endoluminal tumor implant; genetic abnormalities; fluorescence in situ hybridization (FISH)

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Introduction

Sporadic colorectal cancer (sCRC) represents the second leading cause of cancer-related mortality in Europe, being responsible for 177,400 deaths in 2018 (1). Metastatic dissemination of the primary tumor is the major cause of death among sCRC patients. The most common locations of sCRC metastasis are liver, lung and peritoneum. Hematogenous and lymphatic spreading is a well-known mechanism of dissemination. Much less infrequent is the intraluminal shedding with distant intraluminal metastasis. In these cases, the genetic characterization of each of the tumor cell clones present in the primary tumor and in the metastases could help to establish the clonal relationship between them both. In this sense, interphase fluorescence in situ hybridization (iFISH) techniques, although it does not provide specific information about each chromosomal abnormality present in a tumor, with a suitable combination of probes, is used to obtain information about the genetic diversity of a tumor at the single-cell level.

Here, we report a case of sCRC that metastasized in an anal fistula and how a detailed analysis of the chromosomal abnormalities detected by iFISH in the different tumor cell clones identified in each tumor sample (primary and in

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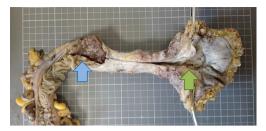


Figure 1 Surgical piece of abdominoperineal amputation in which two tumors are observed, a rectal (blue arrow), and one in the anal margin (green arrow).

anal fistula tumor) allowed us to determine the same clonal origin between both tumors

We present the following case in accordance with the CARE reporting checklist (Available at http://dx.doi. org/10.21037/jgo-20-281).

Case presentation

A 72-year-old man consulted for perineal pain and was initially diagnosed with anal fistula. A partial fistulotomy and cutting seton technique was performed in an outpatient setting, and biopsies were remitted to the pathology department. The histology study revealed a moderately differentiated adenocarcinoma; therefore, the patient was admitted completing the extension study. The colonoscopy showed a second tumor at 15 cm of the anal margin. The CT-scan did not show any distant metastases, peritoneal involvement, or abdominal suspicious lymph nodes. At this point, curative-intent surgery was offered and accepted. Abdominoperineal resection was performed and remitted to the Pathology department. The macroscopic study (Figure 1) displayed a 6-cm rectal tumor and another 3-cm tumor at the anal margin. The histologic study showed a moderately differentiated adenocarcinoma in both locations. The rectal tumor showed extension into pericolorectal fat; meanwhile, the anal tumor demonstrated invasion of the sphincter muscle. No perineural or lymphovascular invasion was observed, and all the lymphatic nodes resected were negative for malignancy. Furthermore, both tumors showed positive CK20 and negative CK7 immunostaining (Figure 2).

After conventional histological studies, a cytogenetic profile of both tumors was performed. For this, a set of 7 different probes specific for those chromosomes and chromosomal regions most frequently gained/amplified and

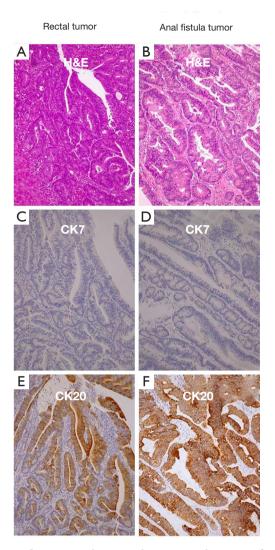


Figure 2 Comparison between the immunophenotypic features of the rectal tumor and the anal fistula tumor (20×). Rectal tumor; hematoxylin-eosin (A), cytokeratin 7 (C) and cytokeratin 20 (E). Anal fistula tumor; hematoxylin-eosin (B), cytokeratin 7 (D) and cytokeratin 20 (F). Both adenocarcinomas showed a CK7-/CK20+ staining pattern.

deleted in sCRC (2) were analyzed by iFISH techniques: 7q31 (*D7S486*), 7p11.1 (centromere), 8p22 (*LPL*), 8p11.1 (centromere) and 8q24 (*CMYC*), 13q14 (*FOXO1*) and 18q21 (*BCL2*); Vysis Inc, Chicago, IL, USA. FISH study displayed an identical genetic profile in both tumors, loss of the chromosomes 8 and 18q, and no alteration in chromosome 7 and 13q; consequently, a common clonal origin was suggested (*Figure 3*).

Eventually, KRAS, NRAS and BRAF genes were studied using PCR techniques and also mismatch repair genes

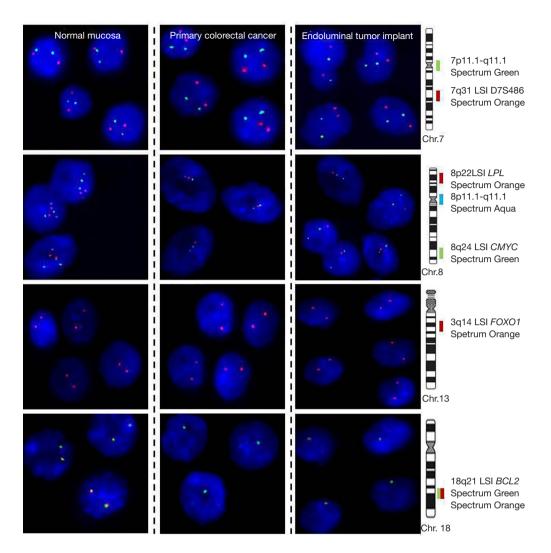


Figure 3 Representative pictures of cell nuclei from primary colorectal cancer, endoluminal implant and normal mucosa after hybridization with two probes for chromosome 7 (7p11.1 and 7q31; green and red signals, respectively), three probes for chromosome 8 (8p22, 8p11, and 8q24; red, aqua and green signals, respectively), one probe for chromosome 13q (13q14; red signals) and two probes (break apart probes) for chromosome 18q (18q21; green and red signals). All cell nuclei of both tumors showed a normal diploid number of hybridization signals for the probes of the chromosome 7 and 13q and loss of a whole chromosome 8 and of the 18q21 chromosomal region, suggesting a common clonal origin.

(MMR; *MLH1*, *MSH2*, *MSH6* and *PMS2*) were assessed by immunochemistry (IHC). *NRAS*, *BRAF* and MMR genes showed no mutation, whereas *KRAS* G12D (c.35G>A; p.Gly12Asp) mutation was only detected in the rectal tumor.

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee(s) and with the Helsinki Declaration (as revised in 2013). The study was reviewed and approved by the ethics committee of the University Hospital of Salamanca. The patient referred in the case report participant provided informed written consent.

Discussion

Intraluminal metastases of sCRC are unusual. The mechanisms underlying in the shedding of exfoliated tumor

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cells is not well understood, but it has been demonstrated experimentally that sCRC sheds viable cells capable of implanting endoluminally (3). By this mechanism, several endoluminal sCRC metastases have been suggested in different locations: colorectal anastomosis, colic ulcers, hemorrhoids, small bowel or common bile ducts (3-6). Anal metastasis has been estimated at 0.05% (95% CI, 0.006-0.08%) of patients with sCRC, and the metastatic mechanism was considered an intraluminal spread in about 80% of cases (7). Most of the intraluminal anal metastasis reported arise in a preexisting lesion: about two-thirds implanted into anal fistula and one fifth in iatrogenic scarring (7). Less than 30 cases of sCRC metastasis into anal fistula have been reported and they may be the first symptom of a left-sided sCRC, most frequently in patients with an advanced clinical stage (7-10). Since primary sCRC and anal metastasis are usually discovered at the same time, various attempts to differentiate synchronous from a metastatic anal tumor have been done, based on several criteria: similar histological findings, lack of dysplasia in the surrounding anal glands, lack of mucin and the absence of lymph node and vasculolymphatic invasion (11). Furthermore, a concordant immunochemistry cytokeratin 7 negative and cytokeratin 20 positive supports a common origin and helps rule out an anal gland adenocarcinoma. In general, the patients reported were treated with curative-intent surgery (in some cases with chemotherapy or chemoradiotherapy associated) and showed good prognosis, better than in cases with lymphatic or hematogenous spread (11).

Several studies have explored the cytogenetic alterations in sCRC, showing that gains/amplifications of chromosomes 7p, 8q, 13q and 20q together with losses of the 1p, 8p, and 17p chromosomal regions are frequent findings in these tumors (12). Although the metastatic tumor usually harbors more chromosomic abnormalities that of their paired primary tumor, it is well known that share highly similar karyotypic abnormalities. Many of these chromosomic abnormalities are acquired before the tumoral spreading (13,14). In our case, iFISH studies revealed that both tumors displayed loss of the chromosomes 8 and 18q without alteration of the chromosomes 7 and 13q. The former is an unusual finding, as Duijf et al. described loss of chromosome 8 in 0.96% of 520 sCRC tumors (15). Conversely, chromosomes 7 and 13q gains have been described in 30% in nonmetastatic disease and are significantly more frequent in metastatic tumors (P<0.05; 61% and 75%, respectively) (16). Those shared infrequent cytogenetic alterations among both tumors support the existence of a close genetic relationship between primary tumors and their intraluminal metastasis, suggesting a common clonal origin instead of synchronous sCRC. Given that some authors propose a more aggressive approach for synchronous sCRC—suggesting subtotal or total colectomy in that cases—differentiate between synchronous and metastatic tumors is important to choose the most adequate therapeutic option (17,18). The generalized use of iFISH techniques in pathology departments makes it a useful and accessible option in daily clinical practice to make an accurate diagnosis in these patients.

Conventional cytogenetics or Spectral Karyotyping (SKY) to define the chromosomal and molecular abnormalities of sCRC has important limitations, mainly related to the need for obtaining tumor metaphases, after an in vitro cell culture period (2); even more, when normal metaphases are obtained it is difficult to confirm that they correspond to the tumor cells and not to the normal cells present in the sample. These include the potential existence of clonal selection during cell culture, the analysis of a small fraction (typically $\leq 1\%$) of all the cells present in the tumor sample, and the difficulties in obtaining tumor cell metaphases in a significant proportion of cases. In addition, other molecular approaches such as array comparative genomic hybridization (aCGH) or single nucleotide polymorphism arrays (aSNPs) also has several limitations: the detection of gains or losses of genetic material present in the majority (e.g., more than 50%) of cells in the tumor sample from which DNA was extracted (12). These approaches do not provide detailed information about the clonal heterogeneity and genetic diversity of a given tumor and the potential co-existence of different abnormalities in the same tumor sample, particularly when tumor cell clones are present at relatively low frequencies (2). Conversely, iFISH techniques, with an adequate combination of probes, we can to obtain information about the genetic heterogeneity of a tumor at the single-cell level.

On the other hand, the mutation *KRAS* discordance observed between the primary tumor and the endoluminal implant may be explained by two mechanisms. The first hypothesis explains that it is a consequence of the divergent evolution of the tumors once the metastases are established. The presence of WT *KRAS*-metastasis with KRAS-mutated primary tumor has been reported in 3 of 99 cases (3%) by Santini *et al.* (19). and in 5 cases of 305 patients (2%) by Knijn *et al.* (20). Furthermore, Miranda *et al.* described a lower rate of mutated *KRAS* in lymph node metastasis than in primary tumor or visceral metastasis, suggesting that those colonizing neoplastic cells spread through lymphatics before *KRAS* mutation occurs (21). They hypothesize that native *KRAS* confers an advantage in the ability to migrate to lymph nodes. A similar mechanism may underlie in intraluminal metastasis; nevertheless, the data available is scant, and further research is necessary. The second possibility is that because of the primary tumor heterogeneity (a well-known circumstance in sCRC), tissue sampling has not included *KRAS*-mutated areas (22).

Conclusions

In conclusion, the use of an accessible technique as iFISH may be an additional useful instrument for the assessment of anal tumors in order to establish their metastatic nature if there is a CRC coexisting tumor.

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Footnote

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Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at http://dx.doi. org/10.21037/jgo-20-281). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee(s) and with the Helsinki

Declaration (as revised in 2013). The study was reviewed and approved by the ethics committee of the University Hospital of Salamanca. The patient referred in the case report participant provided informed written consent

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