Peritoneal Carcinosarcoma and Ovarian Papillary Serous Carcinoma Are the Same Origin: Analysis of *TP53* Mutation and Microsatellite Suggests a Monoclonal Origin

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Carcinosarcoma is an aggressive malignant neoplasm of the female genital tract that is biphasic and composed of carcinomatous and sarcomatous components. It was previously considered as a subtype of uterine sarcoma but is currently regarded as a type of metaplastic carcinoma.¹ Here, we report a case in which a carcinosarcoma forming a discrete mass in the peritoneal cavity was diagnosed concurrently with smaller bilateral ovarian masses showing unequivocal histologic features of high-grade serous carcinoma. Although there seems to be a close relationship between the two tumors in this case, firm evidence for the histopathologic determination of the primary site is lacking because primary carcinosarcomas have been described in many extragenital organs, including the peritoneum, in the literature. We demonstrated two tumors with different histologic findings at distant tumor locations but with a common origin in their pathogenetic mechanism, as suggested by their similar immunohistochemical overexpression of p53 protein and identical pattern TP53 mutation and microsatellites (the reference panel, the Bethesda markers) by molecular analysis.

CASE REPORT

The patient, a 57-year-old postmenopausal woman, was found to have a palpable mass in the right lower quadrant without other clinical symptoms. Her past medical history was unremarkable. On computerized tomographic scan, multiple heter-

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ogeneously enhanced masses of variable size with lobulating contours were identified in the abdominal and pelvic cavities. Whole body ¹⁸F-fluorodeoxyglucose positron emission tomography scan revealed multiple hypermetabolic masses in the bilateral paracolic gutter, mesentery, and pelvic cavity. Of the tumor markers, the serum level of cancer antigen (CA) 72-4 was increased (5.7 U/mL; normal range, 0 to 4 U/mL), but CA 19-9 (11.0 U/mL) and carcinoembyonic antigen (1.5 ng/mL) were within normal range.

Presuming a sarcoma of an unknown primary site, a needle biopsy was performed on the paracolic mass for pathological diagnosis and determination of the primary site. The mass was determined to be carcinosarcoma (a malignant mixed Müllerian tumor). Total abdominal hysterectomy, bilateral salpingo-oophorectomy, right hemicolectomy, and low anterior resection were performed. Currently, the patient received six cycles of postoperative chemotherapy composed of carboplatin and paclitaxel, and she has survived without tumor recurrence for eight months postoperatively. However, a vesicovaginal fistula developed during the treatment as a complication of the surgery.

Grossly, the largest mass (12 cm in diameter) was located in the pericolic soft tissue of the ascending colon. The mass protruded into the peritoneal cavity and displaced the colon to the anterior side, but it was well delineated from the colonic wall without invasion. The cut surface of the pericolic mass was bluish pink and firm, and it also had focal necrotic areas. The ovaries were slightly enlarged bilaterally, with the right ovary measuring $5 \times 4 \times 3.8$ cm and the left ovary measuring $2 \times 1.5 \times 1.5$ cm. Both ovaries had been entirely replaced by solid, well-defined and lobulated tumor tissue. The cut surfaces of both ovaries were yellowish white, firm, and granular. A separate, welldefined, multilobulated mass was also identified within the peri-

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rectal soft tissue. This mass measured 3.8 cm in diameter and did not invade the rectal wall.

Microscopically, the tumor in the pericolic soft tissue showed characteristic features of carcinosarcoma and contained both carcinomatous and sarcomatous components. The carcinomatous portion showed histopathologic features of a high-grade papillary serous carcinoma, with complex and interconnecting papillary structures lined by stratified malignant epithelial cells. The sarcomatous section was composed of malignant spindle cells showing nuclear atypia, increased cellularity, a high mitotic index, and occasional malignant-appearing cartilage (Fig. 1A). The bilateral ovarian and perirectal tumors were histologically characterized by complex and interconnecting papillary structures lined by stratified malignant epithelial cells showing marked cytologic atypia, a high nuclear-to-cytoplasmic ratio and frequent mitotic structures (Fig. 1B). A sarcomatous component was not found despite a thorough examination of the bilateral ovarian tumors and the perirectal mass. A metastatic mass identified in a pelvic lymph node was entirely carcinomatous.

On immunohistochemical staining, both the carcinomatous and the sarcomatous components of the pericolic mass and the carcinomatous component of the ovarian and perirectal masses showed strong immunopositivity for p53 (1:1,500, clone DO-7, Dako, Glostrup, Denmark) (Fig. 1C, D). For the molecular genetic analysis, formalin-fixed, paraffin-embedded tissue was obtained from the pericolic and right ovarian masses, which showed characteristic features of carcinosarcoma and high-grade papillary serous carcinoma, respectively. An area of the tumor in which the percentage of tumor cells exceeded a minimum of 50% of all cells was marked and scraped with a knife blade. Genomic DNA was prepared by proteinase K digestion and followed by phenol/chloroform extraction. Polymerase chain reac-

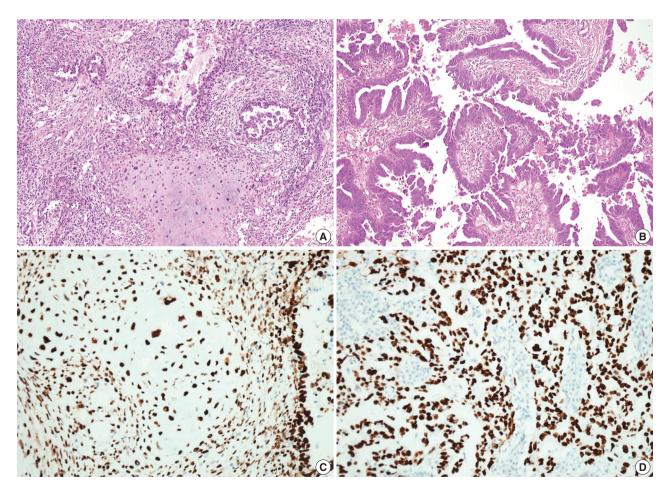


Fig. 1. The pericolic mass shows both carcinomatous and sarcomatous components composed of complex and interconnecting papillary structures lined by stratified malignant epithelial cells and a malignant cartilaginous element (A), but the ovarian mass contains only a carcinomatous component (B). On immunohistochemical staining, both the carcinomatous and the sarcomatous components in the pericolic mass (C) and the carcinomatous component of the ovarian tumor (D) show strong immunopositivity for p53.

tion covering the coding region of exons 5, 6, 7, and 8 of *TP53* was performed, and the amplified DNA was subjected to direct sequencing by the Sanger method (Applied Biosystems, Foster City, CA, USA) and compared with genomic repository data to determine the mutational status of *TP53*. An identical *TP53* somatic point mutation (c.G713A; p.C238Y) in exon 7 was detected in both tumors (Fig. 2). Additionally, microsatellite in-

stability (MSI) was assessed at the loci BAT25, BAT26, D2S123, D5S346, and D17S250. We selected a consensus panel of five microsatellite markers, BAT25, BAT26, D5S346, D17S250, and D2S123, which have been used to diagnose the MSI status of hereditary non-polyposis colorectal cancer (HNPCC) based on the Bethesda guidelines. All of these markers are highly polymorphic microsatellite markers (Fig. 3). All tested microsatel-

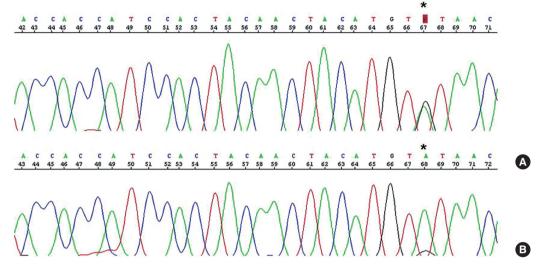


Fig. 2. An identical somatic point mutation (TGT to TAT; p.C238Y) is detected in exon 7 of *TP53* in both the carcinomatous and the sarcomatous component of the pericolic tumor mass (A) and in the epithelial component of the ovarian tumor (B).

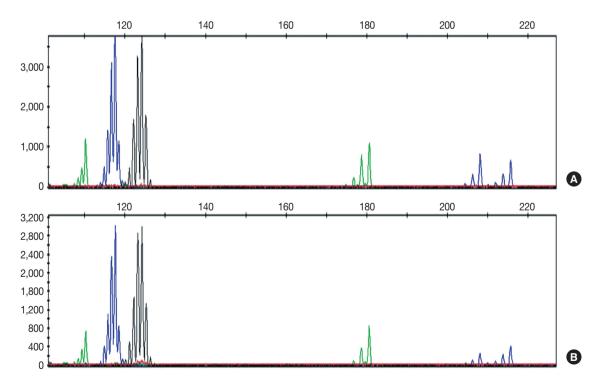


Fig. 3. Several peaks show the same migration pattern, and each main peak of the five markers is seen at the same migration length in the pericolic tumor mass (A) and the epithelial component of the ovarian tumor (B).

lite markers had identical DNA profile patterns with the same allelic sizes between the tumors of the two areas. The overall probability that all five microsatellite markers were identical between the two different tumors was extremely low.

DISCUSSION

A malignant tumor showing a mixed histology with both carcinomatous and sarcomatous components can arise in almost every organ in the body. In the pelvic organs, carcinosarcoma usually occurs in the uterus, but it can also arise in the ovary, fallopian tube, broad ligament, mesentery, cul-de-sac, and pelvic peritoneum, and it comprises approximately 1%–4% of all ovarian malignancies. Currently, it is thought to be a form of metaplastic carcinoma¹ because 1) pure carcinomatous foci are often identified on extensive sampling of the carcinosarcoma, 2) identical genetic changes are identified in both the carcinomatous and the sarcomatous components, 3) carcinosarcoma is often preceded by or associated with endometrioid or serous carcinoma, and 4) the response to adjuvant therapy against high-grade carcinoma is more effective than sarcoma-based therapy.²

The pathogenetic mechanism of peritoneal carcinosarcoma has not been completely clarified. Since carcinosarcoma is often associated with synchronous or metachronous gynecologic tumors of Mullerian origin, including serous and endometrioid carcinoma,^{3,4} and some extrauterine carcinosarcomas were found to be associated with endometriosis or Wolffian duct remnants. carcinosarcoma is thought to arise from extraovarian endometriosis,⁵ immature multipotential cells of the surface epithelium or subcapsular connective tissue of the ovary. The peritoneal lining epithelium in the broad ligament, mesentery, cul-de-sac, and pelvic peritoneum also have the potential to give rise to Mullerian tumors because the Mullerian ducts are derived from the longitudinal invaginations of the mesothelium, which gives rise to the peritoneal lining. Therefore, theoretically, extragenital carcinosarcoma may occur as a primary neoplasm from the pelvic peritoneum.

Since there has been no effective treatment modality for carcinosarcoma, efforts have been made to identify underlying molecular mechanisms to make use of targeted therapies.⁶ Carcinosarcomas in different organs are associated with different oncogenic mutations, including mutations in the epidermal growth factor receptor (*EGFR*) gene, the *PIK3CA* gene, and the p53 gene in tumors of the lung, uterus, and ovary, respectively, indicating distinct oncogenic mechanisms for carcinosarcomas in these different organs. Loss of nuclear WT1 expression in the mesenchymal component might be involved in sarcomatous differentiation. Our molecular analysis revealed an identical mutation in *TP53* occurring in both the carcinomatous and sarcomatous components. All five microsatellites of two tumors retained very similar migration patterns, and a main peak of each microsatellite marker was seen at the same length. These findings of molecular studies suggest that these components share a monoclonal origin.⁷ Moreover, one case of extragenital carcinosarcoma in the literature had the same mutation as a pre-existing ovarian epithelial carcinoma.⁸ The results of our molecular studies in this case and in previous reports support that the ovarian carcinoma and peritoneal carcinosarcoma had the same origin.

Although the carcinosarcoma and the pure carcinoma share a common origin, however, the clinical significance of carcinosarcoma is significantly different from that of pure carcinoma. Patients with ovarian carcinosarcoma had worse five year progression-free and disease-specific survival rates than ovarian papillary serous carcinoma patients at the same International Federation of Gynecologists and Oncologists (FIGO) stages. Patients with a heterologous sarcomatous component also had a worse prognosis than those with a homologous component, although the difference was not statistically significant. The objective response rate to platinum-based chemotherapy was found to be significantly lower in patients with carcinosarcoma compared to those with serous carcinoma.

In conclusion, in cases having two separate tumors in the same patient at distant locations with different histologic findings, meticulous histological examination is necessary, and molecular studies can be helpful to identify the origin of the tumors.

Conflicts of Interest

No potential conflict of interest relevant to this article was reported.

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