

An Effective Hormonal Therapy for a Patient with Estrogen Receptor I (ESRI)-Amplified Metastatic Ovarian Cancer: A Case Report

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Abstract: Hormonal therapy is an important treatment option for estrogen receptor (ER)-positive patients with advanced ovarian cancer. Although ER overexpression has been previously used as an indicator for hormonal therapy, the clinical outcomes of advanced ovarian cancer patients receiving hormonal therapy remain unsatisfactory. Additional biomarkers for screening patients are needed to improve its efficacy. In this study, we reported a metastatic ovarian cancer case with estrogen receptor 1 (*ESR1*) gene amplification and protein overexpression, which showed sustained partial response to hormonal therapy, including letrozole and tamoxifen, and displayed an overall survival of 47 months. The response to the therapy was evaluated by imageological examinations, cancer antigen-125 (CA-125) tests, and circulating tumor DNA (ctDNA) sequencing using capture-based hybrid next-generation sequencing. Our clinical data suggested that *ESR1* amplification might be a potential predictor of response to hormonal therapy in ovarian cancer. The combination of tumor detection techniques including imaging, CA-125 and ctDNA would enable confirmation of tumor response with high confidence.

Keywords: ESR1 amplification, ovarian cancer, hormonal therapy, letrozole, tamoxifen, circulating tumor DNA

Background

Ovarian cancer is the second leading cause of gynecologic cancer death.¹ It accounts for 3.6% and 4.4% of female new cancer cases and deaths per year worldwide.² In China, its incidence and mortality among all female cancer cases are 2.9% and 2.2%, slightly lower than the global level.³

At present, tumor cytoreductive surgery combined with platinum-based chemotherapy is the primary treatment for advanced ovarian cancer. Although the surgery in combination with chemotherapy is effective for most patients, its efficacy for ovarian cancer is not satisfactory and the recurrent rate of ovarian cancer remains high. Hormonal therapy is one of the treatment options for patients with estrogen receptor (ER)-positive advanced ovarian cancer. Studies have shown that the objective response rate and disease stability duration of high-grade serous ovarian cancer patients receiving hormonal therapy were ~15% and 9.6 months, respectively. The insufficient efficacy of hormonal therapy in ER-positive patient cohort limits its clinical application in advanced ovarian cancer and the identification of additional biomarkers for rational patient selection to improve its efficacy is paramount.

There are two subtypes of ER proteins, ER α and ER β , which are encoded by estrogen receptor 1 (ESR1) and ESR2 genes, respectively. It has been reported that 25%–86% of ovarian cancers have ER expression. The success of hormonal therapy in breast cancer and the high frequency of ER expression in ovarian cancer have spurred great effort in evaluating hormonal therapy's efficacy in ovarian cancer. Albeit its low frequency in ovarian cancer, ERS1 gene

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amplification has been shown 100% associated with ER expression in ovarian cancer, indicating it is one of the mechanisms for upregulating ER expression. To our knowledge, we reported the first case of ESR1-amplified metastatic ovarian cancer that showed a durable response to both tamoxifen- and letrozole-based hormonal therapy. Our case report encouraged further exploration of hormonal therapy biomarkers for precision medicine in ovarian cancer.

Case Presentation

In March 2016, a 78-year-old female patient diagnosed with stage IIIC ovarian cancer underwent cytoreductive surgery. Tumor nodules were found in the right ovary, right fallopian tube, left ovary, appendix, omentum, and umbilicus. Postoperative pathology of tumor samples from the ovary revealed high-grade serous adenocarcinoma (Figure 1A). Cancer antigen 125 (CA-125) decreased from 441.82 U/mL to 71.01 U/mL after cytoreductive surgery (Figure 1B).

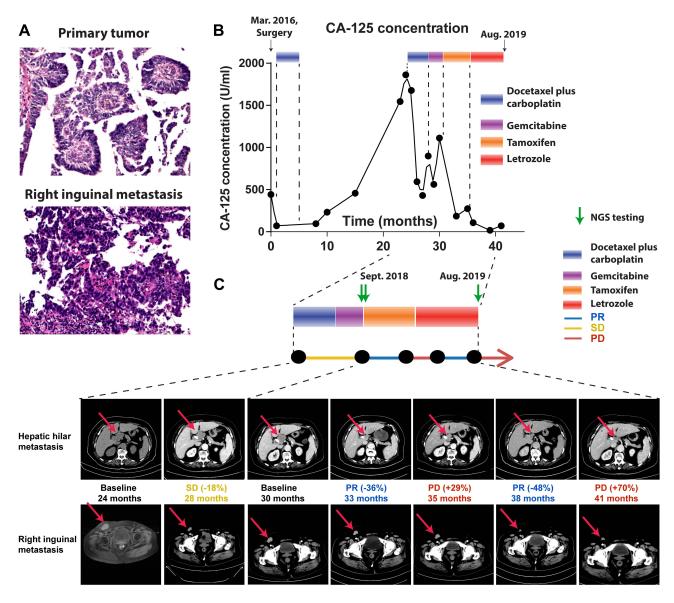


Figure I Patient's tumor response to corresponding chemo and hormonal therapy. (A) Hematoxylin and eosin stain of the primary tumor and right inguinal metastasis. (B) Patient's CA-125 level as a function of time. Mar. 2016 when the patient received cytoreductive surgery, was marked as zero time point. The duration of chemo and hormonal therapies this patient received were indicated as colored boxes at the top of the graph. (C) The CT/MRI images of hepatic hilar metastasis (top) and right inguinal metastasis (bottom) pointed by the red arrows at each time point, along with evaluated tumor response to therapy were shown. The time of next generation sequencing of tumor samples was indicated by a green arrow.

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Table I Systemic Therapy Received by the Patient

#	Regimen	Schedule	Start to End (Month Since Surgery†)
ı	Docetaxel 60 mg/m ² + carboplatin AUC4 [§]	IV [‡] every 21 days × 6 cycles	Apr 2016 to Sep 2016 (1–6)
2	Docetaxel 60 mg/m ² + carboplatin AUC4	IV every 21 days × 4 cycles	Apr 2018 to Jun 2018 (25–27)
3	Gemcitabine 1000 mg/m² d1 d8	IV every 21 days × 2 cycles	Jul 2018 to Sep 2018 (28–30)
4	Tamoxifen	20 mg oral daily	Sep 2018 to Feb 2019 (30–35)
5	Letrozole	2.5 mg oral daily	Mar 2019 to Aug 2019 (36–41)

Notes: †Cytoreductive surgery on Mar. 2016. ‡IV, intravenous. §AUC4, area under curve 4.

Subsequently, she was treated with six cycles of docetaxel plus carboplatin chemotherapy. After postoperative chemotherapy, regular follow-up with monitoring of CA-125 was recommended and performed.

The information of systematic therapies is summarized in Table 1, and as shown in Figure 1B, the level of CA-125 remained at a relatively low level within nine months after surgery. While an enlarged lymph node in the right-groin area was noted after ten months of surgery, the size of which increased at the next follow-up with an elevated CA-125 level. At that time, the patient refused the recommended chemotherapy and continued the regular follow-ups. Chemotherapy was recommended to the patient; however, she refused the treatment. Unfortunately, multiple metastatic lesions were found by abdominal enhanced computed tomography (CT) in the hepatic hilum, intraperitoneal cavity, and right cardiophrenic angle (36*47 mm) after 24 months of surgery with a dramatic increase in CA-125 level (1862.18 U/mL). Eastern Cooperative Oncology Group (ECOG) performance status of the patient at the time was grade 1. Four cycles of docetaxel plus carboplatin were administered as the second-line chemotherapy, which led to a stable disease (SD) according to Response Evaluation Criteria in Solid Tumors (RECIST) version 1.1 criteria (Figure 1C). Side effects of grade 2 white blood cell decrease and grade 1 platelet decrease were reported. Then, she received two cycles of gemcitabine, which only resulted in a temporary decrease in CA-125 level.

In September 2018, the level CA-125 increased to 1113.17 U/mL and an aspiration biopsy was performed on the right inguinal lymph node metastasis. The hematoxylin and eosin stains of the biopsy sample confirmed highgrade serous adenocarcinoma (Figure 1A). Both primary and metastatic tumor specimens underwent capture-based hybrid next-generation sequencing (NGS) by targeting 425 cancer-related genes, which revealed 8.7- and 5.3-fold gene amplification in the primary tumor and metastatic lesion, respectively (Figure 2A). Immunohistochemical (IHC) staining (VENTANA anti-ER (SP1) Rabbit Monoclonal Primary Antibody, Roche) results showed a positive expression of ER in both primary tumor (>80%) and metastatic inguinal lymph node (>90%) (Figure 2B). Based on the NGS-detected ESR1 amplification and ER overexpression, the patient received tamoxifen (20 mg oral daily), a hormonal therapy targeting the ER, starting in September 2018. After three months of tamoxifen treatment, both the right groin and the hepatic hilar tumor nodules significantly shrank (Figure 1C), which reached a partial response (PR) according to RECIST 1.1, and the level of CA-125 decreased to 186.38 U/mL. However, the size of hepatic hilar lymph nodes significantly increased (Figure 1C) after another two months of tamoxifen treatment, suggesting a progressive disease (PD). Aromatase inhibitor letrozole (2.5 mg oral daily) was subsequently given, which led to the shrinkage of metastatic hepatic hilar and right-groin nodules as well as the decrease in CA-125 (Figure 1C). However, after five months of letrozole treatment, CA-125 increased and hepatic hilar lymph node metastasis enlarged again. A re-biopsy of the right inguinal lymph node metastasis was performed for NGS testing, and a 3.2-fold amplification of ESR1 was detected (Figure 2A). The patient then tried hormonal and targeted therapy and chemotherapy but responded poorly and died in February 2020.

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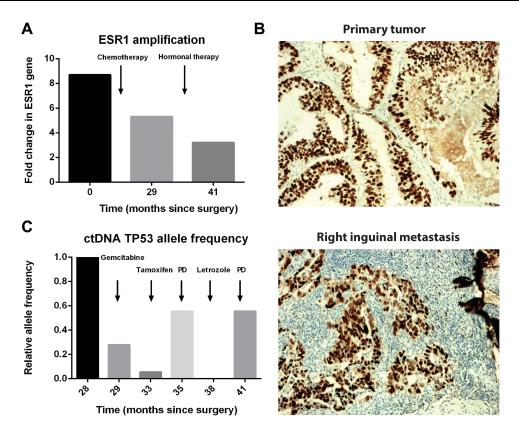


Figure 2 Amplification and overexpression of ESR1 in tumor samples from hepatic hilum and right groin. (A) The amplification of the ESR1 gene in tumor tissue samples at the indicated time. A decrease of ESRI amplification in tumor tissue was observed after chemotherapy and hormonal therapy. Primary tumor and two right inguinal metastasis samples for next-generation sequencing were collected on Feb. 2016, Aug. 2018 and Aug. 2019 respectively. (B) Immunohistochemical staining of ESR1 in samples from the primary tumor (Mar. 2016) and right inguinal metastasis (Aug. 2018). (C) Change of allele frequency of TP53 in ctDNA of the patient. Patient's ctDNA was sequenced at indicated time points, which were calculated as months since surgery on Mar. 2016. The allele frequency of TP53 gene was normalized to the sample at 27 months and compared with each other. PD represents the progressive disease.

Discussion and Conclusions

Although estrogen has a well-defined clinical value in breast cancer, its role in ovarian cancer is less well established. Significant research interest has been seen in evaluating the clinical value of ER expression in ovarian cancer, especially regarding the sensitivity to hormonal therapy.^{8–12} However, mixed clinical outcomes of hormonal therapy were reported in several clinical trials assessing their correlation with ER expression in ovarian cancer. Such diverse results might result from the different methods used to determine ER positivity and the lack of additional predictive biomarkers. In our study, we incorporated targeted NGS with IHC examination, which demonstrated the correlation between ESR1 gene amplification and positive ER expression.

Tamoxifen is one of the first anti-estrogens to enter clinical trials before 2000. As early as 1981, Myers et al reported a favorable response to tamoxifen in three ovarian cancer patients. 13 A systematic review of 648 ovarian cancer patients treated with tamoxifen in clinical trials reported an overall response rate of 13% and a disease control rate of 38%. 14 Since 2002, aromatase inhibitors have been evaluated in a number of clinical trials for ovarian cancer treatment.^{8,15,16} Letrozole is one of the aromatase inhibitors that can block the conversion of androgen to estrogen by inhibiting the key enzyme aromatase in the estrogen synthesis pathway.¹⁷ Two early studies of letrozole have shown that the CA-125 response rate of ovarian cancer patients was improved by restricting ER expression status.^{8,10} A recent clinical study showed that letrozole as a maintenance therapy for patients with advanced ER-positive serous ovarian cancer significantly increased the 24-month relapse-free survival rate from 38.5% to 60%. 18

In this case report, we presented a patient who was sensitive to hormonal therapy and obtained an overall survival of 47 months. The 18-month platinum-free interval after the first-line chemotherapy prompted the use of second-line

Dovepress Wang et al

platinum-based chemotherapy in the hope of a good tumor response. However, the side effect, suboptimal response (stable disease) and the rapid increase in CA-125 level during the second-line chemotherapy promoted a search for alternative therapies. The targeted NGS and IHC tests were performed for primary and metastatic tumor samples, which showed an *ESR1* gene amplification and ER protein overexpression, respectively. These findings led to the subsequent use of hormonal therapies, including tamoxifen and letrozole. The best response to tamoxifen and letrozole was PR with a five-month progression-free survival. However, this single *ESR1*-amplified case with optimal response to hormonal therapy is not sufficient for concluding the predictive role of *ESR1* gene amplification; thus, more comprehensive studies in larger cohorts are needed. A Phase III trial of chemotherapy with or without bevacizumab in patients with platinum-resistant recurrent ovarian cancer has shown that patients' median overall survival is 16.6 months and 13.3 months, respectively. However, the presented patient obtained a survival of 47 months (from Mar. 2016 to Feb. 2020) by the combination of cytoreductive surgery, chemotherapy and hormonal therapy, which was better than the median overall survival of 33.6 months for the platinum-sensitive recurrent ovarian cancer treated with chemotherapy plus bevacizumab in another phase III clinical trial. ²⁰

Circulating tumor DNA (ctDNA) sequencing with liquid biopsy has become a reliable tool for comprehensive genomic assessment and resistance mechanism exploration during treatment. In this case, the changes in ctDNA levels, represented by the allele frequency of *TP53* R273C mutation, were associated with CA-125 levels and treatment efficacy (Figure 2C). The R273C mutation of *TP53* was observed in several cancer types and related to cancer cell proliferation and invasion.²¹ Previous studies also showed that *TP53* mutation in breast cancer could inhibit ER expression and increase tamoxifen resistance.²² Thus, the accumulation of *TP53* mutation detected by NGS might explain the progression upon hormonal therapies.

In conclusion, we reported a metastatic ovarian cancer case with high level of *ESR1* gene amplification, which was associated with strong ER protein expression and prolonged the favorable response to hormonal therapies including tamoxifen and letrozole. Our study promoted the discovery of additional biomarkers for hormonal therapy in ovarian cancer and NGS-detected *ESR1* gene amplification might be a strong candidate in addition to IHC-validated ER expression.

Abbreviations

ER, estrogen receptor; *ESR1*, estrogen receptor 1; CA-125, cancer antigen 125; CT, computed tomography; ECOG, Eastern Cooperative Oncology Group; RECIST, Response Evaluation Criteria in Solid Tumors; SD, stable disease; NGS, next-generation sequencing; IHC, immunohistochemical; PD, progressive disease; ctDNA, circulating tumor DNA.

Data Sharing Statement

The datasets used and/or analyzed during the current study are available from the corresponding authors on reasonable requests.

Ethics Approval and Consent to Participate

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 ethics committee of the Second Affiliated Hospital of Dalian Medical University (reference number 2020059) and with the Helsinki Declaration (as revised in 2013).

Consent for Publication

Written informed consent was obtained from the patient for publication of this case report and accompanying images.

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Wang et al **Dove**press

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Disclosure

Evenki Pan, Yutong Ma, and Xue Wu are employees of Nanjing Geneseeq Technology Inc., China. The other authors declare no conflicts of interest related to this case report.

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