

Human epithelial ovarian cancer cells expressing CD105, CD44 and CD106 surface markers exhibit increased invasive capacity and drug resistance

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Abstract. The high rate of mortality associated with ovarian cancer (OC) is due in part to the development of resistance to chemotherapy, which allows the resistant tumour cells to invade and metastasise. Clarifying the mechanistic basis for drug resistance may reveal novel avenues for treatment. The present study investigated the mechanism of paclitaxel (PTX) resistance in human epithelial OC by evaluating the expression of stem cell-associated cell surface markers endoglin (CD105), CD44 antigen and vascular cell adhesion molecule 1 (CD106), in association with the malignant potential of the human OC OVCAR3 cell line and its PTX-resistant derivative OC3/TAX300. The expression of CD105, CD44 and CD106 was detected by reverse transcription quantitative polymerase chain reaction (RT-qPCR) and flow cytometry, and cell invasion was evaluated using a Transwell invasion assay. CD105, CD44 and CD106 levels were increased in OC3/TAX300 cells compared with the OVCAR3 cells, as determined by flow cytometry ($P < 0.01$) and RT-qPCR ($P < 0.05$). Additionally, the number of invading cells was increased in the OC3/TAX300 group compared with the OVCAR3 group (54.7 ± 6.65 vs. 31.8 ± 6.55 ; $P < 0.01$). A western blot analysis of cell surface marker expression in 80 clinical epithelial OC tissue samples, differing in terms of sensitivity to drug treatments, disease stage and degree of differentiation, revealed that high CD105, CD44 or CD106 expression was associated with drug resistance, advanced disease stage, poor differentiation and high rate of recurrence. These data indicated that exposure to high doses of PTX enhanced the stem-like properties of OC cells,

which are associated with drug resistance and invasion and lead to poor prognosis due to induced chemoresistance and/or metastasis. Therefore, CD105, CD44 and CD106 may serve as potential stem cell-associated cell surface and prognostic markers, and therapeutic targets, in OC.

Introduction

Ovarian cancer (OC) has one of the highest mortality rates among all gynaecological malignancies (1). Paclitaxel (PTX) combined with platinum is the standard chemotherapy regimen for OC. However, treatment failure may occur due to the development of PTX resistance, along with invasion and metastasis of tumour cells.

In cases of resistance, a subset of the tumour cell population exhibits inherited or acquired drug resistance and therefore survives chemotherapy, resulting in tumour recurrence (2). These drug-resistant cells are considered to be cancer stem cells (CSCs), which are responsible for poor prognosis in patients with cancer (3-5). CSCs have the capacity for unlimited proliferation, self-renewal and multilineage differentiation, and may also avoid the effects of chemotherapy, leading to local invasion and distant metastasis. Therefore, an ideal strategy for preventing tumour recurrence is one that targets CSCs (6). In addition, clarifying the mechanisms of action for drug resistance may reveal novel avenues for treatment.

OC is heterogeneous (7), and the tumours contain subpopulations of cells with SC characteristics (8). A number of these subpopulations have been identified, including those positive for CD44 antigen (CD44) and prominin 1 (CD133) (9-11). Cultured epithelial ovarian adenocarcinoma ascite cells have been revealed to exhibit self-renewal and long-term proliferative potential, which is associated with the overexpression of typical CSC markers, including CD44 (12). Recurrent tumours have been demonstrated to exhibit a larger fraction of CSCs expressing aldehyde dehydrogenase 1 family member A1, CD44 and CD133 compared with matched primary OC specimens. Furthermore, several genes involved in SC maintenance, including endoglin (CD105), are upregulated in residual tumour cells in samples from relapsed patients at the end of primary therapy (2), suggesting that resistant tumours

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overexpress SC genes. Cells expressing the mesenchymal SC marker CD105 isolated from human renal carcinoma samples and CD105-positive cells and clones derived from renal carcinoma samples are enriched in tumour-initiating cells with SC characteristics (13). Vascular cell adhesion molecule 1 (CD106) is a surface marker expressed by mesenchymal and neural SCs (14,15) that is associated with OC metastasis and recurrence (16). All of these factors are associated with the self-renewal, chemoresistance and metastasis of cancer cells and may be CSC surface markers. However, to the best of our knowledge, the significance of CD105, CD44 and CD106 as CSC markers in OC has not been investigated previously.

We previously established the PTX-resistant OC OC3/TAX300 cell line with a resistance index of 6.70 by exposing OVCAR3 cells to PTX (17). We hypothesised that resistance to PTX treatment leads to an enrichment of the CSC population in OC cells, with increased expression of SC surface markers including CD105. To examine this hypothesis, the present study analysed the expression of CD105 and other SC surface markers, including CD44 (2) and CD106 (16), in OC3/TAX300 cells and clinical OC tissue samples that were graded in terms of the degree of malignancy in our previous study (7). The invasiveness and metastatic potential of PTX-resistant OC3/TAX300 cells was also evaluated, and the association between the clinical features of the tumours and the expression of SC factors was examined.

Materials and methods

Ethics statement. The present study was approved by the Ethics Committee of the Beijing Shijitan Hospital of Capital Medical University (Beijing, China). Written informed consent was obtained from all participants prior to surgery. All procedures were performed in accordance with the Declaration of Helsinki.

Cell lines and culture conditions. The OVCAR3 cell line was provided by the Basic Medical Research Institute (Beijing, China) and has been described in other studies (18,19). The PTX-resistant OC3/TAX300 cell line used was previously established (17). Cells were cultured in RPMI-1640 medium (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA) supplemented with 10% bovine calf serum (BCS; Beijing Dingguo Changsheng Biotechnology Co., Ltd., Beijing, China), 0.1% penicillin and 0.1% streptomycin at 37°C in an environment containing 5% CO₂.

Reagents and antibodies. TRIzol[®] reagent and primers were obtained from Invitrogen (Thermo Fisher Scientific, Inc.). Dimethyl sulfoxide was purchased from Sigma-Aldrich; Merck KGaA (Darmstadt, Germany). Microplates were purchased from Beijing Dingguo Changsheng Biotechnology Co., Ltd. Mouse monoclonal anti-human CD105 (cat. no. 14606), CD44 (cat. no. 3570) and CD106 (cat. no. 3565) antibodies were purchased from Cell Signalling Technology, Inc. (Danvers, MA, USA).

Flow cytometry analysis of CD105, CD44 and CD106 expression. OVCAR3 and OC3/TAX300 cells in the logarithmic phase were collected and washed twice with PBS.

The cells were resuspended in PBS at a concentration of 1x10⁶/ml, and a 100- μ l suspension was incubated with 5 μ l PerCP-Cy5.5-conjugated anti-CD105, phycoerythrin-conjugated anti-CD44 or fluorescein isothiocyanate-conjugated anti-CD106 antibodies (BD Biosciences, San Jose, CA, USA) for 1 h at 37°C. The cells were washed twice with PBS and analysed with a FACSCalibur flow cytometer (FlowJo 7.6.1; BD Biosciences).

Reverse transcription quantitative polymerase chain reaction (RT-qPCR) analysis. Total RNA was extracted from cells using TRIzol reagent, according to the manufacturer's protocol and reverse transcribed using moloney murine leukemia virus reverse transcriptase, 5X first strand buffer (dithiothreitol), 10 mM deoxynucleotide triphosphates, Oligo (dT)18 (all from Tiangen Biotech Co., Ltd., Beijing, China). The temperature protocol used for RT was 42°C for 50 min and 95°C for 5 min. Primers targeting CD105, CD44 and CD106 genes were designed according to sequence data obtained from GenBank (20), with β -actin (ACTB) used as an internal control. The primer sequences were as follows: CD105, 5'-GCCAAGGGCAACTGTGTGA-3'(sense) and 5'-CCGGTT TTGGGTATGGGTACT-3' (antisense); CD44, 5'-CCTCTT GGCCTTGGCTTTG-3' (sense) and 5'-CTCCATTGCCAC TGTGATCAC-3' (antisense); CD106, 5'-TGGTCCAGCCCTT CCTCCAT-3' (sense) and 5'-AGGATTTTCGGAGCAGGA AAG-3' (antisense); and ACTB, 5'-AGGTCACCATTGGCA ATG-3' (sense) and 5'-GGTAGTTTCGTGGATGCCACA-3' (antisense). The cDNA was amplified using SYBR Green PCR Master Mix (Applied Biosystems; Thermo Fisher Scientific, Inc.) and the thermocycling conditions were as follows: 50°C for 2 min, 95°C for 10 min, then 40 cycles of 95°C for 15 sec and 60°C for 1 min for the amplification curve and 95°C for 15 sec, 60°C for 15 sec and 95°C for 15 sec for the dissociation curve. Data were analysed using Sequence Detection Software V2.2 (Applied Biosystems; Thermo Fisher Scientific, Inc.) and exported to an Excel spreadsheet (Microsoft Corporation, Redmond, WA, USA). Target gene expression levels were normalised to that of ACTB. The mRNA expression ratio of CD105, CD44 or CD106 to ACTB was calculated to obtain relative expression values using the formula: $\Delta Cq = Cq$ (target gene) - Cq (ACTB); where q is the number of cycles when the DNA concentration reached the threshold (21).

Invasion assay. The invasive capabilities of OVCAR3 and OC3/TAX300 cells were assessed using a Transwell assay. Cells in the logarithmic phase were collected and washed with PBS, and then cultured in serum-free medium for 24 h. In total, 40 μ l Matrigel was coated on the membrane of the upper chamber surface and incubated at 37°C for 30 min to solidify. The cells were then collected, and their concentration was adjusted to 1x10⁵ cells/ml; they were then seeded in the upper chamber of a 24-well Transwell insert coated with a thin layer of Matrigel (BD Biosciences). The lower chamber was filled with RPMI-1640 medium supplemented with 20% BCS. The cells were incubated at 37°C and 5% CO₂. After 24 h, cells remaining in the upper chamber were scraped off along with the Matrigel using a sterile swab, and cells that had invaded into the insert were fixed with 40 g/l paraformaldehyde for 20 min at room temperature and stained with 0.01% crystal

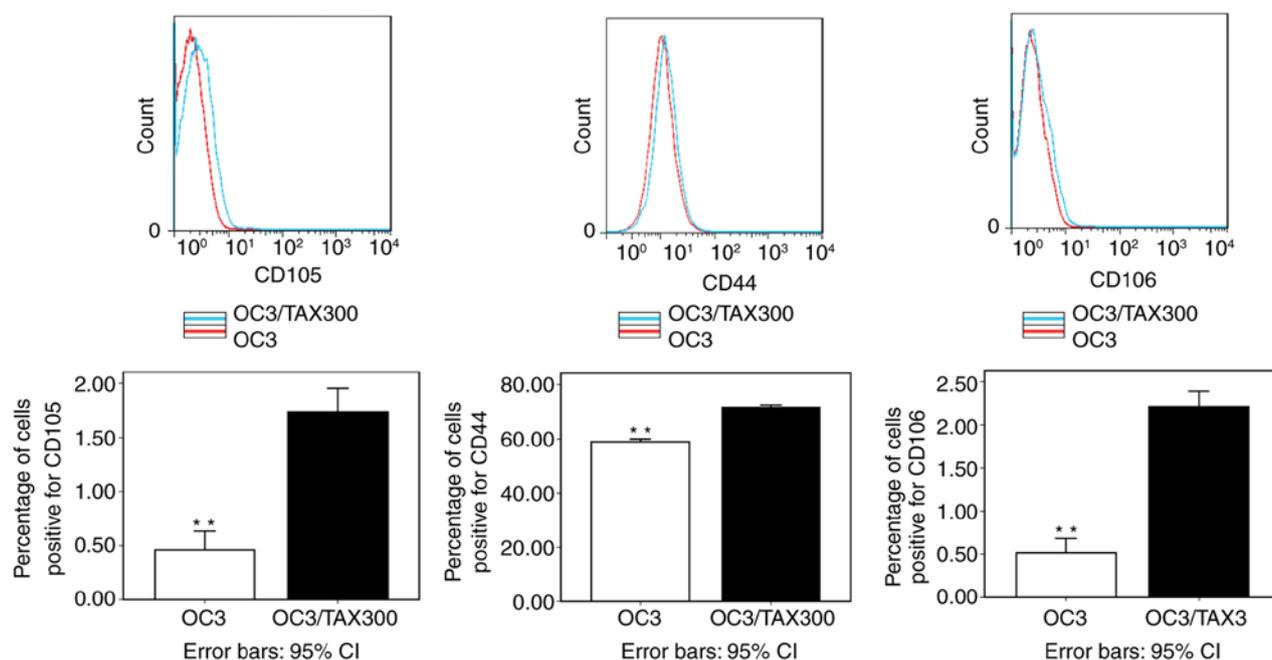


Figure 1. Quantitative analysis of OC3/TAX300 and OVCAR3 cells positive for CD105, CD44 and CD106 detected by flow cytometry. The fraction of cells positive for the 3 markers was increased in the PTX-resistant OC3/TAX300 cell line compared with the PTX-sensitive OVCAR3 cell line. ** $P < 0.01$. PTX, paclitaxel; CD44, CD44 antigen; CD105, endoglin; CD106, vascular cell adhesion molecule 1; CI, confidence interval.

violet for 20 min at room temperature for observation by light microscopy (magnification, x40). Images of cells in at least five randomly selected microscopic fields were captured and the number of cells was counted. The average number of cells was used to assess the invasive capacity of the 2 cell lines.

Clinical OC samples. A total of 80 epithelial OC tissue samples were collected from the specimen repository of Beijing Shijitan Hospital between April 2012 and February 2013 (7). The median age of the patients was 56.15 years (range, 23-79 years). There were 52 primary and 28 recurrent cases; 64 were poorly differentiated and 16 were moderately or highly differentiated OC tissue; and 63 samples were advanced-stage (III) whereas 17 were early-stage (I and II) OC. Immediately following cytoreductive surgery, all specimens were analysed with the ATP-based tumour chemosensitivity assay (ATP-TCA) as described previously (7). These specimens were graded according to the National Comprehensive Cancer Network guidelines (22). Routine histopathological analysis was performed for samples obtained from the same tissues to determine the stage and histological features of the tumour samples simultaneously with ATP-TCA testing. The sensitivities of specimens to PTX, carboplatin (CBP), topotecan, gemcitabine (GEM), docetaxel (TXT), bleomycin, etoposide and 4-hydroperoxycyclophosphamide were examined using an *in vitro* ATP-TCA procedure. Cancer recurrence was defined according to the current clinical criteria as: Return of cancer following completion of treatment following a period of time during which the cancer was not detected (23). In OC, patients with platinum-sensitive cancer were those who achieved complete remission and experienced relapse at 6 months or later following initial platinum-based chemotherapy, whereas patients with platinum-resistant cancer were those who exhibited recurrence within 6 months (24).

Western blot analysis. Total cell lysates were prepared using radioimmunoprecipitation assay lysis buffer (Beyotime Institute of Biotechnology, Shanghai, China) and the supernatant was collected via centrifugation at 4°C and 4,024 x g for 10 min. A bicinchoninic acid assay was used to determine the protein concentration. Aliquots (30-40 μ l) of cell lysates were heated at 100°C for 5 min, and 10 μ g of protein was loaded into each well of a 10% SDS-PAGE gel for electrophoresis. The proteins on the electrophoresis gel were then transferred to an Immobilon-P membrane that was incubated in blocking solution [5% bovine serum albumin (Beijing Dingguo Changsheng Biotechnology Co., Ltd.) in TBS-Tween 20] for 1-3 h at 25°C followed by overnight incubation at 4°C with mouse monoclonal anti-human CD105, CD44 and CD106 antibodies at dilutions of 1:1,000. Subsequent to washing 3 times in TBS with 0.1% Tween 20, the membrane was incubated for 1-2 h at 25°C with a fluorophore-conjugated secondary antibody (cat. no. 610-132-121; Rockland Immunochemicals, Inc., Limerick, PA, USA) at a dilution of 1:5,000. The membrane was washed and analysed using an Odyssey two-colour infrared imaging system (LICOR Odyssey, LI-COR Biosciences, Lincoln, NE, USA). The signal intensity of protein bands was calculated using Image J software (v1.8.0; National Institutes of Health, Bethesda, MD, USA).

Statistical analysis. Data are presented as the mean \pm standard deviation and were analysed using SPSS v.17.0 for Windows software (SPSS Inc., Chicago, IL, USA). Means were compared using a two-sided t-test. Linear regression analysis was used to detect the correlation between sensitivity index (SI) and expression levels of target genes. All experiments were independently repeated a minimum of three times. $P \leq 0.05$ was considered to indicate a statistically significant difference.

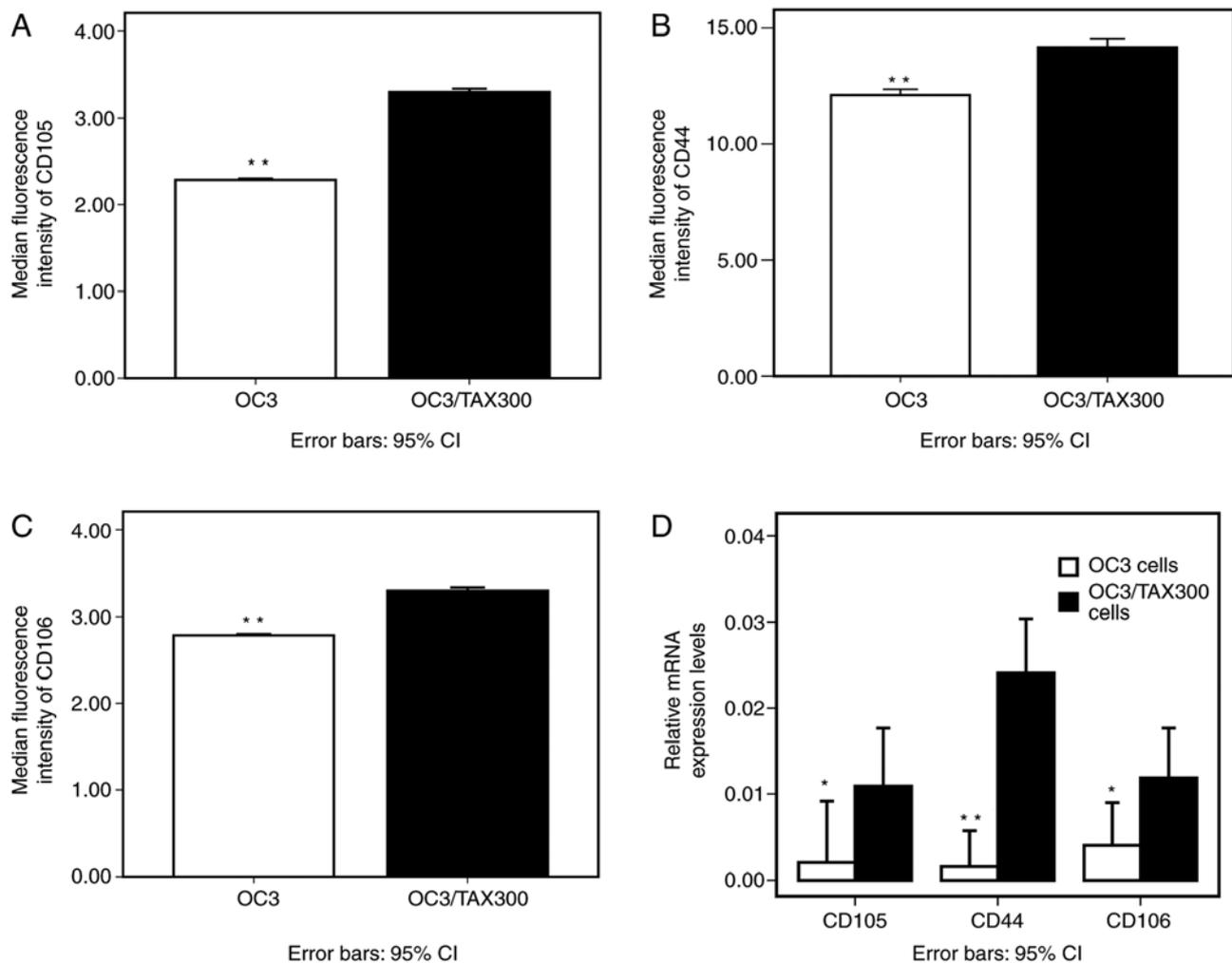


Figure 2. Median fluorescence intensity of CD105, CD44 and CD106 in PTX-resistant OC3/TAX300 and PTX-sensitive OVCAR3 cells, as determined by flow cytometry. (A-C) (A) CD105, (B) CD44 and (C) CD106 overexpression in OC3/TAX300 cells. (D) Relative mRNA expression levels of CD105, CD44 and CD106 in OC3/TAX300 cells. * $P < 0.05$ and ** $P < 0.01$. PTX, paclitaxel; CD44, CD44 antigen; CD105, endoglin; CD106, vascular cell adhesion molecule 1; PTX, paclitaxel; CI, confidence interval.

Results

CD105, CD44 and CD106 are overexpressed in PTX-resistant OC cells. The percentage of cells positive for the 3 proteins was increased in the PTX-resistant cell line compared with the PTX-sensitive cell line ($P < 0.01$; Fig. 1). Accordingly, the median fluorescence intensities of CD105, CD44 and CD106 were increased in OC3/TAX300 cells compared with the OVCAR3 cells ($P < 0.01$; Fig. 2A-C). The results from the RT-qPCR analysis demonstrated a similar trend to those obtained by flow cytometry, with increased relative expression levels of CD105, CD44 and CD106 mRNA in the OC3/TAX300 cells compared with the OVCAR3 cells ($P < 0.05$; Fig. 2D).

PTX-resistant OC cells exhibit increased invasive capabilities. The Transwell assay revealed that numerous OC cells had invaded the membrane filter (Fig. 3A and B). The numbers of invaded cells in 12 different fields of vision were counted after 24 h culture, and the quantitative analysis indicated that the number of invaded cells was increased in the OC3/TAX300 group compared with the OVCAR3 group (54.7 ± 6.65 vs. 31.8 ± 6.55 ; $P < 0.01$; Fig. 3C).

CD105, CD44 and CD106 are highly expressed in drug-resistant epithelial OC tissue samples. In the western blot analysis, the protein expression levels of CD105, CD44 and CD106 in 80 epithelial ovarian cancer tissues were markedly different. This may be associated with the heterogeneity of chemotherapy treatments in patients (7). It was identified that 66/80 were PTX-sensitive and 14/80 were PTX-resistant; and 47/80 were CBP-sensitive and 33/80 were CBP-resistant (Table I). The difference of protein expression in chemoresistant or -sensitive samples was demonstrated by the following: CD105, CD44 and CD106 were expressed at high and low levels in PTX-resistant and PTX-sensitive tissue samples, respectively (Fig. 4A). Statistical analysis of the data demonstrated the significant difference: Protein expression of CD105 was different in 80 specimens with different sensitivity to 8 drugs, and there were increased protein expression levels of CD105 in PTX/CBP/TXT resistant samples compared with sensitive specimens (Fig. 4B). There was an increased protein expression level of CD44 in PTX resistant samples compared with sensitive specimens (Fig. 4C). Among patients exhibiting chemoresistance or sensitivity to the commonly used chemotherapy drugs

Table I. Results of chemosensitivity assay in OC samples.

Drug	Sensitivity, n (%)	Weak sensitivity, n (%)	Resistance, n (%)	Sensitivity (%)
PTX	44 (55)	22 (27.5)	14 (17.5)	82.5
CBP	21 (26.4)	26 (32.4)	33 (41.2)	58.8
TPT	14 (17.5)	23 (28.7)	43 (53.8)	46.2
TXT	18 (22.5)	18 (22.5)	44 (55)	45
GEM	13 (16.3)	13 (16.3)	54 (67.5)	32.5
4-HC	3 (3.7)	18 (22.5)	59 (73.8)	26.2
VP-16	4 (6.3)	10 (11.2)	66 (82.5)	17.5
BLM	0	0	80 (100)	0

PTX, paclitaxel; CBP, carboplatin; TPT, topotecan; TXT, docetaxel; GEM, gemcitabine; 4-HC, 4-hydroperoxycyclophosphamide; VP-16, etoposide; BLM, bleomycin.

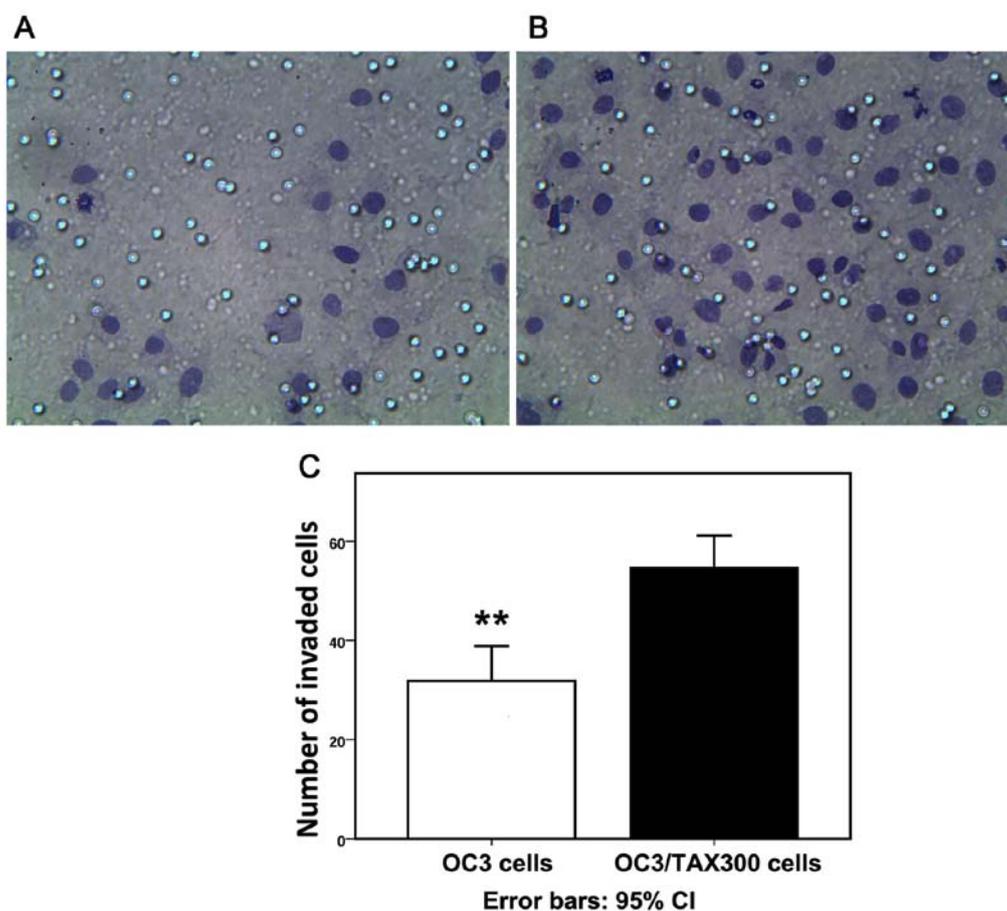


Figure 3. Detection of invasive tumour cells using the Transwell assay. A number of OC cells had invaded the membrane filter. (A) OVCAR3 cells. (B) OC3/TAX300 cells. (C) The number of invaded cells was increased in the OC3/TAX300 group compared with the OVCAR3 group. $**P<0.01$. OC, ovarian cancer; CI, confidence interval.

including PTX, CBP or TXT, CD106 levels were increased in the chemoresistance group compared with the chemosensitive group (Fig. 4D).

In our previous study, the SI was used to evaluate drug sensitivity rates and was calculated using the formula: $SI=500-\%$ tumour growth inhibition at 200, 100, 50, 25 and 12.5% + test drug concentration (7). The expression levels of these target genes in the tissue samples were associated with

the SI of a number of the drugs examined; for example, the CD105 level was correlated with the SI value of PTX and TXT (Table II).

CD105, CD44 and CD106 expression levels are associated with clinical parameters of epithelial OC. Moderately and highly differentiated OC tissue samples exhibited decreased CD105 protein expression compared with those

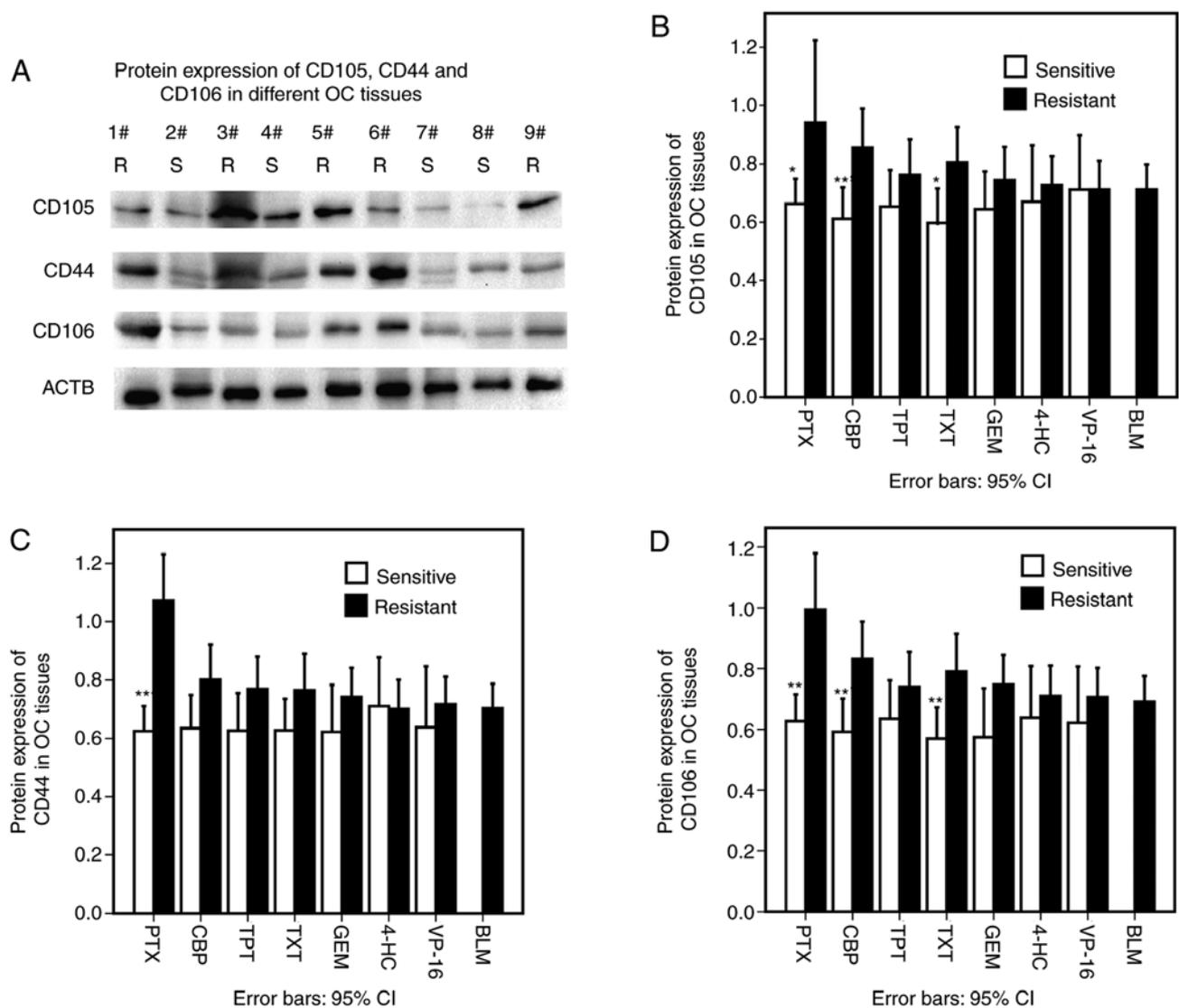


Figure 4. CD105, CD44 and CD106 are highly expressed in drug-resistant epithelial OC tissue samples. (A) Representative blots of 3 experiments of PTX-resistant and PTX-sensitive samples were subjected to western blot analysis for CD105, CD44 and CD106 protein levels. ACTB was used as a loading control. Each number corresponds to a different patient. CD105, CD44 and CD106 were expressed at high and low levels in PTX-resistant and PTX-sensitive tissue samples, respectively. (B) Difference of expression of CD105 protein in all specimens with different sensitivities to 8 drugs. There were increased protein expression levels of CD105 in PTX/CBP/TXT resistant samples. (C) Difference of expression of CD44 protein in all specimens with different sensitivity to 8 drugs. There was an increased protein expression level of CD44 in PTX resistant samples compared with the sensitive specimens. (D) Difference of expression of CD106 protein in all specimens with different sensitivity to 8 drugs. There were increased protein expression levels of CD106 in PTX/CBP/TXT resistant samples compared with the sensitive specimens. * $P < 0.05$ and ** $P < 0.01$. CD44, CD44 antigen; CD105, endoglin; CD106, vascular cell adhesion molecule 1; BLM, bleomycin; CBP, carboplatin; GEM, gemcitabine; PTX, paclitaxel; TPT, topotecan; TXT, docetaxel; VP-16, etoposide; 4-HC, 4-hydroperoxycyclophosphamide; R, PTX-resistant samples; S, PTX-sensitive samples; ACTB, β -actin; CI, confidence interval; OC, ovarian cancer.

Table II. Results of correlation analysis between expression levels of target genes and sensitivity index of tested chemotherapy drugs.

Drug combination	Correlation coefficient, <i>r</i>	P-value
CD105 and PTX	0.327	0.003
CD105 and TXT	0.285	0.010
CD44 and PTX	0.353	0.001
CD106 and PTX	0.344	0.002
CD106 and TXT	0.321	0.004

CD105, endoglin; CD44, CD44 antigen; CD106, vascular cell adhesion molecule 1; PTX, paclitaxel; TXT, docetaxel.

that were poorly differentiated ($P = 0.002$). Furthermore, CD105 expression was decreased at the early stage (I and II) compared with the advanced stage (III) tissues ($P = 0.019$) and decreased in the primary tumour samples compared with the recurrent ovarian epithelial cancer specimens ($P = 0.006$). Similar trends were observed for CD44 and CD106 expression (Fig. 5A-C).

All of the cases in the present study were followed up for at least 2 years after initial chemotherapy. A total of 44 patients (55%) were classified as clinically CBP-sensitive and 36 (45%) were clinically CBP-resistant. Clinically CBP-sensitive OC tissue samples exhibited decreased CD105 and CD106 expression compared with the resistant cases (Fig. 5D).

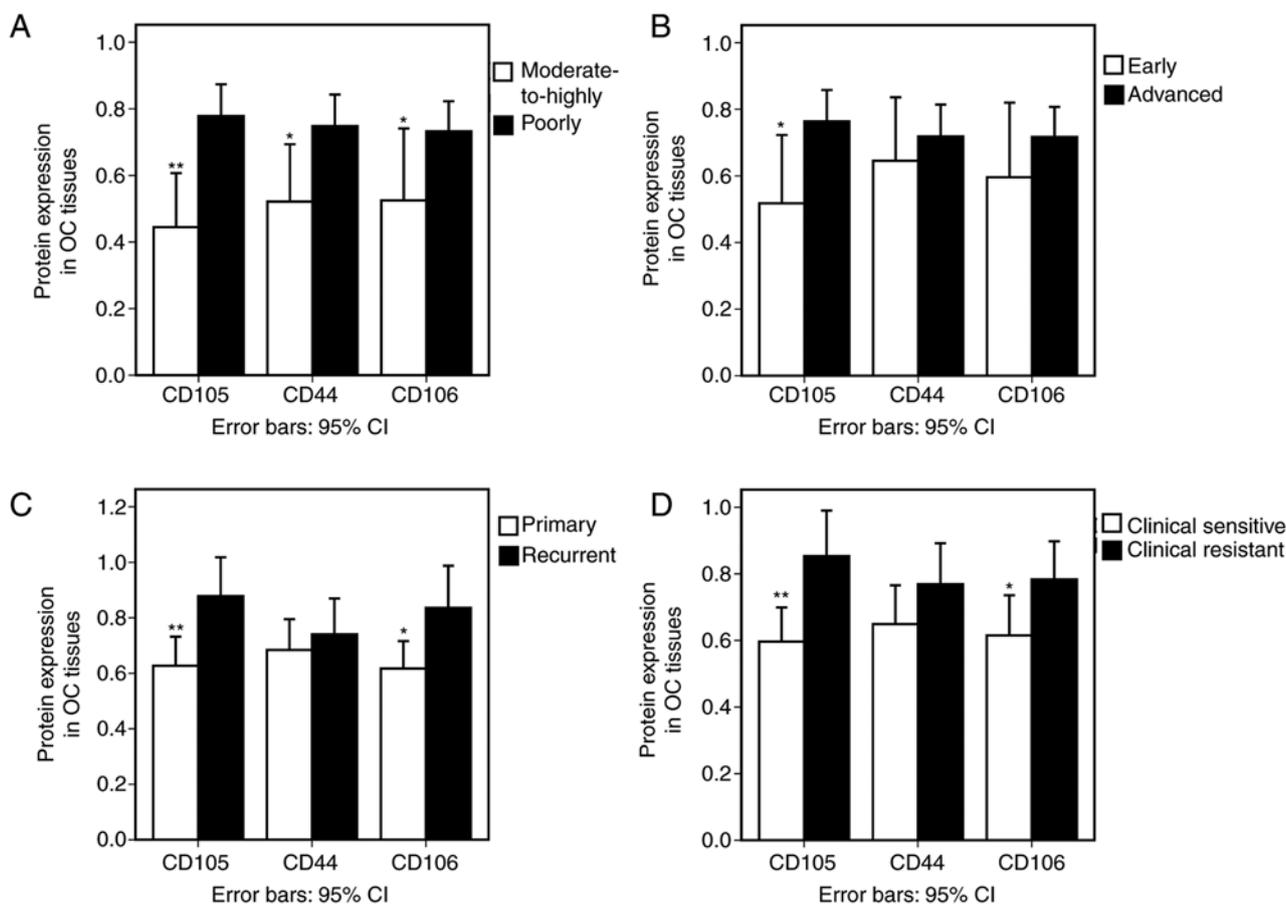


Figure 5. CD105, CD44 and CD106 expression levels are associated with clinical parameters of epithelial OC. (A) Moderately and highly differentiated OC tissue samples exhibited decreased CD105, CD44, and CD106 protein expression compared with poorly differentiated samples. (B) Early-stage (I and II) OC tissue samples exhibited decreased CD105 protein expression compared with advanced-stage (III) samples. (C) Primary OC tissue samples exhibited decreased CD105 and CD106 protein expression compared with recurrent OC tissue samples. (D) Clinically CBP-sensitive ovarian epithelial cancer tissues exhibited decreased CD105 and CD106 expression compared tissues that were clinically CBP-resistant. * $P < 0.05$ and ** $P < 0.01$. CD44, CD44 antigen; CD105, endoglin; CD106, vascular cell adhesion molecule 1; OC, ovarian cancer; CBP, carboplatin; CI, confidence interval.

Discussion

The present study investigated the association between the expression of the SC markers CD105, CD44 and CD106 and the invasive capabilities and chemotherapeutic resistance of OC cell lines. It was identified that all 3 proteins were overexpressed in PTX-resistant OC3/TAX300 cells and in chemoresistant and poorly differentiated or advanced-stage epithelial OC tissues, and that this was associated with enhanced invasive capacity. These data suggest that exposure to high doses of PTX enhances the SC properties of ovarian tumour cells (including CD105, CD44 and CD106 overexpression), leading to the development of PTX resistance, increased invasion and long-distance metastasis, and poor prognosis in OC.

CD105 is a co-receptor for transforming growth factor (TGF)- β family proteins including TGF- β 1 and - β 3, and serves a key role in development, cell proliferation, extracellular matrix synthesis, angiogenesis and the immune response (25). CD105 exhibits SC characteristics and may stimulate endothelial cell growth; its high level of expression on peri- and intratumoural vessels is associated with poor prognosis following cancer treatment (26-28). CD105 overexpression has also been identified to be associated with decreased patient

survival rates and distant metastasis (27-30); this is likely due to the angiogenesis-promoting function of CD105, which has been demonstrated to increase tumour vasculature and ultimately lead to poor prognosis (31-33).

CD105 is expressed not only in vascular endothelial cells, but it is also detected in several malignancies including gastrointestinal stromal tumours, hepatocellular carcinoma, and breast cancer (34-36), head and neck paragangliomas (37), and OC ascites (38). Furthermore, CD105-expressing cells have multi-differentiation potential: CD105-positive rhabdoid meningioma cells exhibit SC-like features and have the capacity to differentiate into adipocytes and osteocytes (39).

Various studies have demonstrated that CD105 overexpression is associated with chemoresistance. CD105 is rarely detected in primary OC cells, but is expressed at an increased level in platinum-resistant cells compared with primary untreated tumour cells (2). Notably, the protein is predominantly localised in the cytoplasm, which is consistent with the features of a CSC-like population. CD105 inhibition increases cisplatin sensitivity and decreases OC cell viability while enhancing apoptosis via induction of double-stranded DNA damage (40). It has also been suggested that chemotherapy stimulates CD90 and CD105 expression in hepatocellular carcinoma cells (41). These studies suggest that poor prognosis in cancer is not solely

due to the induction of tumour angiogenesis by CD105 (42), but that it is also caused by CD105 overexpression in tumour cells. Accordingly, CD105 has been investigated as a potential therapeutic target: One study identified that downregulation of CD105 decreased tumourigenicity and GEM resistance, suggesting that CD105 expression not only distinguishes a CSC subpopulation but also confers self-renewal capacity and contributes to chemoresistance in renal cell carcinoma (43). Additionally, the anti-CD105 antibody TRC105 inhibited tumour growth and improved survival without off-target toxicity in a mouse model of mammary carcinoma (44), with similar results demonstrated in acute leukaemia (45). A previous clinical study that enrolled 26 patients with hepatocellular carcinoma showed that TRC105 combined with sorafenib was well tolerated at the recommended single agent doses of both drugs (46), and another clinical trial that enrolled 13 patients with urothelial carcinoma also found TRC105 was well tolerated, although the benefits of extended survival of patients need further examination (47).

CD106, also known as vascular cell adhesion molecule 1, is a member of the immunoglobulin superfamily of transmembrane proteins that bind integrin (48). CD106 mediates leukocyte adhesion to endothelial cells and downstream signalling cascades (49), and serves an important role in the oncogenesis, tumour angiogenesis, tumour progression, and metastasis of human cancer (50) including colorectal carcinoma (51), non-Hodgkin lymphoma (52) and gastric carcinoma (53). CD106 is highly expressed in OC (54), has been associated with ovarian tumour growth, and may be a prognostic indicator and potential therapeutic target (16). CD106 was also demonstrated to be overexpressed in breast cancer (55) and enhances breast cancer cell metastasis to the lungs (56). It has previously been demonstrated that CD106 is highly overexpressed in lung cancer compared with normal lung tissue, and that it is associated with poor survival. Additionally, the invasive potential of lung cancer cells is significantly weakened by CD106 silencing (57).

CD44 is a classic surface marker of SCs, which promotes oncogenesis and tumour progression (58). Cells with this phenotype are more likely to form tumours compared with those with alternate phenotypes. A number of studies have suggested that CD44 is a reliable cell surface marker for CSCs in gastric (59) and breast cancer (60), glioma (61), colon cancer (62) and OC (11). High CD44 expression is associated with metastasis, recurrence, chemoresistance and survival rate in OC, whereas its downregulation suppresses tumour cell proliferation and metastasis and reverses chemoresistance (63). The present study demonstrated that OC cells expressing CD105, CD44 and CD106 on their surface exhibited greater invasive capabilities and drug resistance. Although the results are consistent with the earlier studies, additional studies are required to determine whether targeting these factors would be effective for the treatment of OC.

In conclusion, the results from the present study demonstrated that CD105, CD44 and CD106 were upregulated in PTX-resistant OC cell lines and chemoresistant epithelial OC tissues. This was associated with poor prognosis, distant metastasis and early recurrence. At present, these results have certain limitations, as they are only based on a PTX-resistant OC cell line and its primary parent cell line, and also require validation by knocking down CD105, CD44 or CD106 genes. However, these positive results suggested future avenues of study, and other PTX or platinum resistant cell lines will be examined in

subsequent experiments, and CD105 gene knockdown will be performed to study the changes of invasiveness of OC3/TAX300 cells following inhibition of the expression of CD105 gene and the tumourigenicity in nude mice. Therefore, inhibiting CD105, CD44 or CD106 expression has potential as an adjuvant therapy for OC. Additionally, as these factors confer SC characteristics, investigating other CSCs markers may provide a basis for targeted therapy and for predicting patient prognosis.

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Availability of data and materials

The datasets used and/or analysed during the present study are available from the corresponding author on reasonable request.

Authors' contributions

JZ, BY and HZ performed the experiments. JZ and HL performed the statistical analysis and wrote the manuscript.

Ethics approval and consent to participate

The present study was approved by the Ethics Committee of the Beijing Shijitan Hospital of Capital Medical University (Beijing, China). Written informed consent was obtained from all participants prior to surgery. All procedures were performed in accordance with the Declaration of Helsinki.

Patient consent for publication

Written informed consent was obtained from all participants prior to surgery.

Competing interests

The authors declare that they have no competing interests.

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