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Vorstand: Prof. Dr. Jens Werner

New Prognostic and Predictive Biomarkers in Primary Ovarian Cancer

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Katharina Anna Dötzer

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Berichterstatter:	PD Dr. rer. nat. Barbara Mayer
Mitberichterstatter:	Prof. Dr. med. Fabian Trillsch
	PD Dr. med. Julia Jückstock
	Prof. Dr. Robert Pernenczky
Mitbetreuung durch den	
promovierten Mitarbeiter:	Prof. Dr. med. Nina Ditsch
Dekan:	Prof. Dr. med. Thomas Gudermann
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List of Abbreviations

BRCA: Breast cancer gene CAM-DR: Cell-adhesion mediated drug resistance CD: Cluster of differentiation DCR: Disease Control Rate ECM: Extracellular matrix EGFR: Epidermal growth factor receptor ERα: Estrogen receptor alpha FIGO: Fédération Internationale de Gynécologie et d'Obstétrique HGFR: Hepatocyte growth factor receptor NSCLC: non-small-cell lung cancer **ORR: Overall Response Rate OS: Overall Survival** PD-1: Programmed cell-death protein 1 PD-L1: Programmed cell-death ligand 1 PFI: Platinum Free Interval PFS: Progression Free Survival QTiS: Quantification of the tumor immune stroma SCLC: small-cell lung cancer TAM: Tumor associated macrophage **TIL:** Tumor infiltrating Lymphocyte VEGF: Vascular Endothelial Growth Factor WHO: World Health Organization

List of Publications

Paper 1:

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Integrin α2β1 Represents a Prognostic and Predictive Biomarker in Primary Ovarian Cancer.

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1. Introduction

1.1. Ovarian Cancer

1.1.1.Epidemiology

In 2020, worldwide ovarian cancer incidence was 313,959 and 207,252 deaths as a result were counted. In Germany there have been 7,162 new cases and 5,328 deaths [1]. This makes ovarian cancer to the second leading death cause among gynecologic cancers. As the incidence rates increase with age, the average age at time of diagnosis is 68 years. One in 76 women will suffer from ovarian carcinoma during their lifetime, which results in a lifetime prevalence of 1.3% [2].

1.1.2. Clinical Manifestation and Classification

Currently, there are no established screening methods that can detect ovarian cancer in early stages [3,4]. Usually, unspecific symptoms occur in advanced stages, leading to late diagnosis for the majority of patients [5]. Around 75% of the patients receive diagnosis in stage III and IV (see Table 1), in which tumor mass has usually spread in the whole peritoneal cavity.

FIGO-Stage	Definition	Distribution at initial diagnosis
I	Tumor confined to ovaries or fallopian tubes	22%
Ш	Tumor with pelvic extension (below pelvic brim)	7%
111	Tumor with spread outside the pelvis	43%
IV	Distant metastasis (excluding peritoneal metastases)	29%
Table 1: FIGO	staging classification for ovarian cancer (according to [2,6])	

1.1.3. Therapy (according to German Guidelines)

Primary surgery combines staging and radical tumor debulking in advanced stages. An open laparotomy allows exploration of the complete peritoneal cavity. Peritoneal cytology, multiple peritoneal biopsies, bilateral salpingo-oophorectomy, total abdominal hysterectomy, appendectomy, omentectomy and pelvic and paraaortal lymph node dissection are conducted to define tumor stage. Further, macroscopic tumor should be completely removed by radical debulking [4]. Despite rigorous surgical resection around 50 to 60% of patients suffer from macroscopic residual tumor burden, which is associated with poor prognosis [7-9].

After primary surgery in nearly all cases (except stage IA/IB and grading G1/G2) an adjuvant first-line systemic chemotherapy is recommended. It should consist of six cycles carboplatin combined with paclitaxel. Due to high toxicity and severe side effects of this drug combination, chemotherapy has to be terminated early for more than 10% of patients, which is associated with poor prognosis. Further

dose modification is required even more frequently [8,10,11]. To improve prognosis, a large number of studies was conducted, evaluating the effect of targeted therapies in ovarian cancer. So far, only a small number of candidates could be included in the guideline recommendations [12-14].

In advanced stages, Bevacizumab, a VEGF-inhibitor, can be administered simultaneously to chemotherapy and as maintenance therapy for up to 15 months. For patients with BRCA-mutation the PARP-inhibitor Olaparib can be applied as single maintenance therapy or in combination with Bevacizumab. Recently, the PARP-inhibitor Niraparib is recommended for all patients in advanced stages independent of BRCA-mutation status [4]. Currently many phase-III-trials are ongoing, evaluating the efficacy of various PARP-inhibitors in combination with chemotherapy or other targeted therapies, like checkpoint-inhibitors (NCT03642132, NCT03602859, NCT03737643, NCT03522246).

In case of recurrence, further therapy options depend on the interval from completion of chemotherapy to relapse. There are four categories: Platinum-refractory (no response to chemotherapy, relapse < 4 weeks), platinum-resistant (relapse < 6 months), partial platinum-sensitive (relapse < 6 months), partial platinum-refractory and platinum-resistant disease platinum-free monotherapy is recommended for second-line therapy. Recently, the definition of 'platinum-resistant' has begun to change. Instead of strictly using a cut-off of six months for selection of second-line therapy, more factors should be included e.g., age and performance status of the patient, patient's preference, histology, and BRCA-mutation. Nevertheless, there are no established markers, to predict response to platinum-based first- or second-line therapy [15]. Due to frequent relapse, predictive markers could help to stratify patients for effective therapies and thus improve poor prognosis.

1.1.4. Prognosis and Prognostic Factors

The 5-year-survival rate of ovarian cancer is stage-dependent. For patients in stage III at the time of diagnosis its 39%, in stage IV it is 20% [2]. Low survival rates originate from frequent and early relapses after chemotherapy. 70 to 80% of patients suffer from a relapse. In most patients it occurs within the first three years after chemotherapy [16,17].

There are several proven prognostic clinicopathological factors. High age and a low performance status of the patient, high tumor grading and advanced tumor stage are associated with poor prognosis [4,18,19]. The presence of macroscopic residual tumor after surgery is another important prognostic factor. Best prognosis can be expected after optimal cytoreduction. If the amount of macroscopic residual tumor is less than 10mm, prognosis is still better than with more residual tumor [7,20]. Consequently, prognosis also depends on the quality of primary surgery.

Additional to clinicopathological factors recently factors concerning tumorbiology have proven their prognostic value. The new WHO classification defined the histologic subtypes by molecular patterns, differing in survival rates. Certain gene signatures and cell cycle proteins were found to correlate with prognosis [21,22]. As it has been already proven for other cancer entities, high counts of TILs are associated with better prognosis [23,24].

In summary the poor prognosis of ovarian cancer and the lack of biomarkers for patient stratification indicates the high medical need for new prognostic and predictive factors. Identification of those was the purpose of the present work and is summarized in two publications. In the paper "Immune Heterogeneity Between Primary Tumors and Corresponding Metastatic Lesions and Response to Platinum Therapy in Primary Ovarian Cancer" the immune infiltrate in the microenvironment of primary ovarian cancer was analyzed. The protein expression pattern of drugable targets was investigated in the paper "Integrin $\alpha 2\beta 1$ Represents a Prognostic and Predictive Biomarker in Primary Ovarian Cancer".

1.2. Immune Infiltrate in Ovarian Cancer

1.2.1. Microenvironment in Ovarian Cancer

Many characteristics and features of ovarian cancer are connected to its tumor microenvironment. Single tumor cells and aggregates separate from the primary tumor, resulting in mainly peritoneal and omental metastases [25]. Components of the ovarian cancer microenvironment are (1) the ECM, (2) residential cells at the site of metastasis, like mesothelial cells or omental adipocytes, or (3) recruited cells, like fibroblasts, macrophages, and other immune cells. By interaction with ovarian cancer cells, they can be reprogrammed and therefore enable chemotaxis, adhesion, tumor growth, angiogenesis, metastasis, immune modulation and chemoresistance [26-28].

1.2.2. Tumor-Infiltrating Lymphocytes in Ovarian Cancer

As a part of the tumor microenvironment immune cells can enable, but also inhibit tumor growth. While TAMs or regulating T-cells induce immunosuppression, TILs contribute to antitumoral activity [29]. Meta-analyses have proven the prognostic value of TILs. A high amount of intraepithelial CD3⁺ or CD8⁺-cells is associated with better PFS and OS in ovarian cancer [23,24].

1.2.3. Relevance of Checkpoint-Inhibitors in Ovarian Cancer

PD-1 and its ligand PD-L1 play an important role in regulation of the immune system and tolerance of T-cells. Binding of the two ligands leads to suppression of immune response in the PD-1 expressing T-cell. Usual PD-L1-expression on antigen-presenting cells or tissue in sites of immune privilege, like placenta or testes, leads to immune regulation by the PD-1 pathway. But expression on tumor cells of various entities is frequent and therefore decreases antitumoral activity of the peritumoral immune infiltrate [30-32].

Monoclonal antibodies, targeting PD-1 and PD-L1, were developed to restore antitumoral activity of T-cells. In recent years, the development of checkpoint-inhibitors led to an improvement of cancer therapy in many entities. They are part of the first-line therapy for NSCLC, SCLC, renal cell carcinoma and malignant melanoma [33-35]. Due to the discovery of the role of TILs for antitumoral activity in ovarian cancer, checkpoint inhibitors have become promising therapeutic options.

Initial studies of checkpoint inhibitors in recurrent ovarian cancer showed ORRs of 10 to 15% with DCRs of around 40 to 50% [36-38]. KEYNOTE-100 was the first large, phase-II-trial. Pembrolizumab, a

PD-1 inhibitor, was administered to patients with recurrent ovarian cancer, showing an ORR of 8% and DCR of 37%. In this study PD-L1 expression was evaluated as a possible biomarker and correlated with a higher response rate [39]. Nevertheless, monotherapy with checkpoint-inhibitors showed only modest anti-tumoral activity in recurrent ovarian cancer. More promising results were reported in the phase-II-trials MEDIOLA and TOPACIO. Combination of checkpoint- and PARP-inhibitors in recurrent ovarian cancer resulted in ORR of 18 to 72% and DCR of 65% [40,41]. Consequently, ongoing phase-III-trials frequently combine checkpoint-inhibitors with standard chemotherapy or PARP-inhibitors. Currently (accessed on 23 February 2022) there are fourteen phase-III-trials registered on clinicaltrials.gov which are evaluating checkpoint-inhibitors in ovarian cancer (six in primary and eight in recurrent disease). In Phase I and II more than 100 trials can be found. This demonstrates the future potential and importance of understanding the immune infiltrate and checkpoint-inhibitor based therapy in ovarian cancer.

1.2.4. Intratumoral Heterogeneity of Microenvironment and Immune Infiltrate

As it can be seen in Table 1, more than 70% of the patients suffer from tumor spread outside the pelvis at initial diagnosis, mostly in form of peritoneal and omental metastases. Due to the different immunological structure of peritoneal and omental tissues, differences in the tumor microenvironment can be expected. Intratumoral heterogeneity has already been observed on genomic, transcriptomic, and proteomic level [42,43]. Considering the different composition of peritoneal and omental tissues [27,44], various tumor microenvironments can be expected. Heindl et al. showed a correlation of high intratumoral diversity of the tumor microenvironment composition and poor prognosis [45]. Jimenez-Sanchez et al. could prove heterogeneity between the immune infiltrate of multiple tumor lesions of one patient. Further different phenotypes were associated with stable or progressing metastases [46]. Considering the increasing relevance of immunotherapy, immune heterogeneity must be considered as a further potential prognostic and predictive factor in ovarian cancer.

1.3. Biomarker Expression in Ovarian Cancer

1.3.1. Importance of Biomarkers for Patient Stratification

Biomarkers are defined as a "characteristic that is measured as an indicator of normal biological processes, pathogenic processes, or biological responses to an exposure or intervention, including therapeutic interventions." They can "include molecular, histologic, radiographic, or physiologic characteristics". [47] Defined by the process that is measured, there are various categories of biomarkers.

Category	Description
Susceptibility/ risk	Indicates potential for developing a disease
Diagnostic	Detects or confirms presence of a disease
Monitoring	Detects a change in the degree or extent of disease over time
Prognostic	Identifies likelihood of a clinical event, disease recurrence or progression
Predictive	Identifies individuals, who will experience effect from exposure to treatment

Table 2: Categories of Biomarkers (according to [47,48])

As already mentioned above, due to poor prognosis and early relapse, there is an urgent need for prognostic and predictive biomarkers in primary ovarian cancer. They could improve patient stratification and therefore enable treatment planning for individual patients. There are some features of tumor biology, which have already proven to be prognostic biomarkers (see 1.1.4) and help to assess the course of disease in individual patients. Identification of new negative prognostic biomarkers, which can be addressed by targeted therapy, could expand treatment options.

High rates of relapse and platinum-resistance in ovarian cancer have triggered a search for reliable predictive biomarkers. Many candidates for prediction of platinum-sensitivity have been found in the past [49,50], but none has proven its predictive value in large trials and been established in guidelines. Once platinum-resistance is proven, there are only a few therapy options with small chance of long-term response. Consequently, more targeted therapies must be established for patients with platinum-resistance.

The aim of the following publications was the identification of prognostic and predictive biomarkers, that can be detected by immunohistochemistry and addressed by existing targeted therapies.

1.3.2. Integrins in Ovarian Cancer

Integrins are transmembrane cell-adhesion receptors. They are heterodimers built from one α subunit and one β -subunit. Currently, there are 18 α -subunits and 8 β -subunit identified in mammals, which can form 24 different heterodimers. As they are transmembrane proteins, they connect the cytoskeleton to the ECM and are capable of bidirectional signaling. Changes in the extracellular milieu can be detected and influence intracellular processes. Otherwise, intracellular signaling can change allosteric conformation of integrins and therefore their affinity to ligands [51,52]. Due to these features, various steps of cancer progression are mediated by integrins. As main cell-adhesion-receptors they are crucial for migration, invasion, and metastasis [53,54]. As already mentioned above, ovarian cancer metastasis predominantly takes part in the peritoneal cavity. Cells detach from the primary tumor and spread in the peritoneal fluid. Final attachment to mesothelial cells of peritoneal and omental tissue, local invasion and proliferation is mediated by integrins [25,55,56]. Consequently, they are crucial for interactions with the tumor microenvironment [26,27]. Previous studies indicate the ability of integrins in promoting chemotherapy resistance in ovarian cancer. These cell-adhesion mediated drug resistance (CAM-DR) is based on intracellular signaling initiated by integrin-binding to the ECM [57,58].

Especially β 1-integrins are involved in the above-mentioned processes in ovarian cancer. Integrin α 4 β 1 and α 5 β 1 are known to play a major role in attachment of cancer cells to the mesothelial cells of peritoneal and omental tissue [56,59]. Less is known about the heterodimer α 2 β 1, but previous studies suggest a contribution to chemotherapy resistance in primary ovarian cancer [60,61].

Being an essential part of progression and metastasis in ovarian cancer, integrins are promising candidates as prognostic and predictive biomarkers, but also as targets for development of further therapeutic strategies in ovarian cancer.

1.3.3.Integrin-targeted Therapy

Due to their involvement in a wide range of cellular processes and functions integrins are promising targets for new treatment strategies concerning different pathologies. Several drugs have already been approved: α IIb β 3-antagonists are used for the treatment of myocardial infarction, Natalizumab and Vedolizumab (antibodies against integrin α 4 β 1 and α 4 β 7) are approved for treating inflammatory bowel diseases and multiple sclerosis [62]. Although integrin-targeted therapy in cancer has been subject of extensive research in the past two decades, there has been no approval until now. Especially inhibitors for integrin α v β 3, which is known to play an important role in tumor angiogenesis, showed promising results in preclinical data. Cilengitide, a cyclic pentapeptide, was administered to glioblastoma patients even in a phase-III-trial, but finally failed for a positive outcome [63,64].

Failure due to intensive research in the last decades can be explained by the complex biology of integrins. After binding to a ligand, integrins perform conformational changes, which affect the affinity to the ligand or the intracellular activity. Therefore, antibodies or peptides, which are actually designed for inhibition of integrin activity, can have the opposite effect. Another point is the reduced selectivity of designed ligands. Mostly they are directed towards one subunit, which results in binding to all heterodimers including this subunit [62].

There are new approaches, directing these problems. Allosteric ligands which stabilize the inactive state of an integrin are designed. Further, especially tumor entities or single patients with high expression of a single integrin can be targeted by an integrin-binding-partner, which is conjugated with a cytotoxic molecule [62,65]. Therefore, identifying expression levels of integrin in ovarian cancer might specify the suitability as a possible therapeutic target.

1.4. Objectives

As shown above, patients with ovarian cancer suffer from poor prognosis. Despite radical surgical resection and adjuvant chemotherapy, 70 to 80% of patients experience relapse. There is an urgent need for prognostic and predictive biomarkers, which enable patient stratification. They could help to identify patients, who benefit from platinum-based chemotherapy and further represent possible drugable targets for new therapy options in ovarian cancer.

The aim of the following publications was to find prognostic and predictive biomarkers in the tumor microenvironment of ovarian cancer. Expression of possible candidates was analyzed via immunohistochemical staining and correlated with clinicopathological data.

As one crucial component of the tumor microenvironment, the immune infiltrate represented the main topic of the first publication. It evaluates the composition of the immune infiltrate, its predictive value and intratumoral heterogeneity between corresponding lesions within the peritoneal cavity.

The second publication focused on the cell adhesion molecule integrin $\alpha 2\beta 1$. The prognostic and predictive value of the single integrin expression, but also of dual expression with other biomarkers has been evaluated. In addition to the first publication, integrin $\alpha 2\beta 1$ expression was correlated with the immune infiltrate.

1.5. Contribution

For both publications, patients and their clinicopathological data from the SpheroID-study were included. This study was initiated by PD Barbara Mayer and approved by the institutional review board of the Ludwig-Maximilians-Universität, Munich, Germany at 07th August 2012 (No.278/04).

Experimental design of both studies was developed by the authors Katharina Dötzer and PD Barbara Mayer.

In the first publication the author contributed to the development of a new immune cell counting method (QTiS algorithm) [66]. She conducted the immunohistochemical staining, photodocumentation of the results and the application of the new-established QTiS algorithm. The statistical analysis, including the correlation of experimental data with clinicopathological data and survival analysis as well as the comparison of primary tumor and corresponding lesions, was performed by the author under the supervision of Alexander Crispin. She prepared the final manuscript, including all tables and figures under the supervision of Barbara Mayer.

For the second publication Katharina Dötzer conducted the immunohistochemical staining with the help of Michael Pohr, Anton Stolp and Frank Arnold. She evaluated the biomarker expression by semiquantitative scoring. The statistical analysis, including the correlation of experimental and clinicopathological data as well as uni- and multivariate survival analysis was performed by the author. The manuscript, including tables and figures, was prepared by Katharina Dötzer under the supervision of Barbara Mayer.

2. Summary

2.1. Abstract (English)

Despite intensive research in recent years, diagnosis of ovarian cancer is still connected to a poor prognosis. Due to advanced tumor stage at the time of diagnosis, most of patients must undergo radical surgery followed by platinum-based chemotherapy. 70 to 80% of patients suffer from relapse in spite of this intensive treatment. To identify patients, who do not respond to platinum-based therapy, additional prognostic and predictive biomarkers must be found. They can enable patient stratification and further represent targets for new therapy options. Due to fast tumor spread within the peritoneal cavity, ovarian cancer cells are interacting with different cell types of the tumor microenvironment from early stages on. Addressing this key feature, prognostic and predictive markers were searched in the tumor microenvironment of ovarian cancer. For this purpose, tumor samples were analyzed immunohistochemically and correlated with clinicopathological data of patients.

The publication "Immune Heterogeneity Between Primary Tumors and Corresponding Metastatic Lesions and Response to Platinum Therapy in Primary Ovarian Cancer" analyzed differences in the immune infiltrate of primary tumors and corresponding omental and peritoneal lesions, as well as the associated predictive impact.

Immune heterogeneity was observed between omental lesions and corresponding primary tumors. A higher count of stromal CD45⁺ (p = 0.007), CD3⁺ (p = 0.005), CD8⁺ (p = 0.012) und PD-1⁺ (p = 0.013) cells was found in omental lesions. Furthermore, lymph node metastasis correlated with a higher stromal infiltrate of CD45⁺ (p = 0.018) and CD3⁺ (p = 0.037) cells in omental lesions.

Intratumoral heterogeneity was also found between primary tumors and corresponding peritoneal lesions. While more PD-1⁺ immune cells were detected in primary tumors (p = 0.054), PD-L1 expression was trending to be higher in peritoneal lesions (p = 0.078). Additionally, tumor heterogeneity between primary tumors and their peritoneal lesions was identified as a predictive marker for platinum-sensitivity. In case of a higher amount of intratumoral CD8⁺ cells in the peritoneal lesion compared to the primary tumor, platinum-sensitivity was more frequent (p = 0.045). On the contrary, a higher stromal infiltrate of PD-1⁺ cells in the peritoneal lesion was associated with reduced platinum-sensitivity (p = 0.045).

In addition to the composition of the tumor microenvironment, mediators of interaction between cancer cells and the tumor microenvironment were analyzed in the publication "Integrin $\alpha 2\beta 1$ Represents a Prognostic and Predictive Biomarker in Primary Ovarian Cancer".

High expression of the cell-adhesion-molecule integrin $\alpha 2\beta 1$ was an independent prognostic factor for shorter PFS (p = 0.021) and PFI (p = 0.022). Relations to other therapy-relevant biomarkers have been evaluated. A significant correlation was found between high expression of integrin $\alpha 2\beta 1$ with EGFR (p = 0.027) and ER α (p = 0.035). Further, dual expression of Integrin $\alpha 2\beta 1$ with HGFR (PFS/PFI: p = 0.004) or CD44v6 (PFS: p = 0.000; PFI: p = 0.001; OS: p = 0.025) is associated with poor survival. A reference to the first publication was made by evaluation of the immune infiltrate in correlation to integrin $\alpha 2\beta 1$ expression. Low counts of various cell types of the immune infiltrate were associated with a high expression of Integrin $\alpha 2\beta 1$ (CD3 intratumoral: p = 0.017; CD3 stromal: p = 0.034; PD-1 intratumoral: p = 0.002; PD-1 stromal: p = 0.049). This confirms integrin $\alpha 2\beta 1$ as a poor prognostic factor and outlines its possible immunosuppressive role.

In search for additional therapeutic strategies to treat ovarian cancer more attention must be paid to the tumor microenvironment. The expression of integrin $\alpha 2\beta 1$ and the the heterogeneity of the immune infiltrate have proven to be predictive markers for platinum sensitivity. Therefore, they may represent possible stratification markers for future therapy choices. In addition, the findings can contribute to a closer understanding of platinum resistance in ovarian cancer.

The above identified prognostic and predictive biomarkers can help to personalize treatment choices and identify patients fitting for platinum-based chemotherapy, immunotherapy, like checkpointinhibitors, or new targeted therapies. Due to frequent co-expression and to evade therapy resistance dual-targeting must be considered as promising approach.

2.2. Zusammenfassung (Deutsch)

Trotz intensiver Forschung in den letzten Jahren ist die Prognose für Patienten mit der Diagnose Ovarialkarzinom weiterhin schlecht. Aufgrund meist später Diagnosestellung in einem fortgeschrittenen Tumorstadium müssen sich die meisten Patienten einer ausgedehnten Operation mit anschließender platinhaltiger Chemotherapie unterziehen. Bei 70 bis 80% der Patienten kommt es trotz dieser intensiven Therapie zum Rezidiv. Daher besteht ein hoher Bedarf an prognostischen und prädiktiven Biomarkern, welche Patienten erfassen, die schlecht auf eine platin-basierte Chemotherapie ansprechen. Diese Biomarker können sowohl bei der Einteilung der Patienten nach Prognose helfen als auch potenzielle Ziele für neue Therapieoptionen darstellen. Da sich das Ovarialkarzinom meist früh in der Peritonealhöhle ausbreitet, interagieren die Tumorzellen der Metastasen mit sehr unterschiedlichen Formen von Mikromilieu. Hier kann auf der Suche nach neuen prognostischen und prädiktiven Markern angesetzt werden. Zu diesem Zweck erfolgte die immunhistochemische Analyse verschiedener Tumorproben und die anschließende Korrelation mit klinischen und pathologischen Daten der entsprechenden Patienten.

Die Veröffentlichung "Immune Heterogeneity Between Primary Tumors and Corresponding Metastatic Lesions and Response to Platinum Therapy in Primary Ovarian Cancer" beschreibt die Unterschiede des Immuninfiltrats im Primärtumor und in den zugehörigen Metastasen des Omentums und Peritoneums. Diese Unterschiede erweisen sich als mögliche prädiktive Faktoren.

Es zeigten sich deutliche Unterschiede in der Zusammensetzung der Immunzellen zwischen dem Primärtumor und den Metastasen des Omentums. Im Omentum konnten deutlich mehr CD45⁺ (p = 0.007), CD3⁺ (p = 0.005), CD8⁺ (p = 0.012) und PD-1⁺ (p = 0.013) Zellen im Stroma beobachtet werden. Zudem fanden sich bei Patientinnen mit bereits bestehender Lymphknotenmetastasierung mehr stromale CD45⁺ (p = 0.018) and CD3⁺ (p = 0.037) Zellen im Omentum.

Unterschiede zeigten sich auch zwischen dem Primärtumor und den entsprechenden peritonealen Metastasen. Es fanden sich mehr PD-1⁺ Immunzellen im Primärtumor (p = 0.054), dafür war die PD-L1 Expression der Tumorzellen in den Metastasen des Peritoneums höher (p = 0.078). Diese Heterogenität zwischen Primärturmor und peritonealen Metastasen stellte sich als möglicher prädiktiver Marker für Platinsensitivität heraus. Falls sich intratumoral in den peritonealen Metastasen mehr CD8⁺ Zellen als im Primärtumor fanden, zeigten sich die Karzinome häufiger platinsensitiv (p = 0.045). Ein ausgeprägteres stromales PD-1⁺ Infiltrat der peritonealen Metastase war hingehen mit einer reduzierten Platinsensitivität assoziiert (p = 0.045).

Neben der Zusammensetzung des Mikromilieus wurden in der Veröffentlichung "Integrin α2β1 Represents a Prognostic and Predictive Biomarker in Primary Ovarian Cancer" auch das Zusammenspiel zwischen Mikromilieu und Krebszellen untersucht.

Eine hohe Expression des Zelladhäsionsmolekül Integrin $\alpha 2\beta 1$ zeigte sich als unabhängiger prognostischer Faktor für ein kürzeres PFS (p = 0.021) und PFI (p = 0.022). Zudem ergaben sich auch Zusammenhänge bezüglich anderer therapierelevanter Biomarker. Es zeigte sich eine hohe Expression von EGFR (p = 0.027) oder ER α (p = 0.035) bei gleichzeitiger hoher Expression von Integrin $\alpha 2\beta 1$. Eine gleichzeitige hohe Expression von Integrin $\alpha 2\beta 1$ und HGFR (PFS/PFI: p = 0.004) oder CD44v6 (PFS: p = 0.000; PFI: p = 0.001; OS: p = 0.025) war mit schlechterem Überleben assoziiert.

Ein Bezug zur ersten Veröffentlichung wurde hergestellt, indem ein möglicher Zusammenhang zwischen der Expression von Integrin $\alpha 2\beta 1$ und Immunzellen untersucht wurde. Tumoren mit geringerer Anzahl verschiedener Leukozyten (CD3 intratumoral: p = 0.017; CD3 stromal: p = 0.034; PD-1 intratumoral: p = 0.002; PD-1 stromal: p = 0.049) zeigten gleichzeitig häufig eine hohe Integrin $\alpha 2\beta 1$ Expression. Dies bestätigt nochmals Integrin $\alpha 2\beta 1$ als negativen prognostischen Faktor und unterstreicht seine immunsuppressive Wirkung.

Auf der Suche nach neuen Therapiemöglichkeiten für das Ovarialkarzinom muss der Fokus in Zukunft auf das Mikromilieu des Tumors gelegt werden. Sowohl die Expression von Integrin $\alpha 2\beta 1$ als auch die Heterogenität der Immunzellen zeigten sich als prädiktive Faktoren zur Vorhersage von Platinsensitivität. Somit eignen sie sich sowohl als Marker für weitere Therapieentscheidungen, als auch als möglicher Ausgangspunkt um die Mechanismen der Platinresistenz beim Ovarialkarzinom zu erforschen.

Die genannten neuen prognostischen und prädiktiven Biomarker können in Zukunft helfen, Therapieentscheidungen zu personalisieren und die richtige Wahl aus platinbasierter Chemotherapie, Immuntherapie oder neuen zielgerichteten Therapien für jeden Patienten zu treffen. Um Therapieresistenzen zu vermeiden, müssen aufgrund von häufig gleichzeitiger Expression der entsprechenden Marker auch Kombinationstherapien in Erwägung gezogen werden.



Article

Immune Heterogeneity Between Primary Tumors and Corresponding Metastatic Lesions and Response to Platinum Therapy in Primary Ovarian Cancer

Katharina Dötzer¹, Friederike Schlüter¹, Markus Bo Schoenberg¹, Alexandr V. Bazhin^{1,2}, Franz Edler von Koch³, Andreas Schnelzer⁴, Sabine Anthuber⁵, Dieter Grab⁶, Bastian Czogalla⁷, Alexander Burges⁷, Jens Werner^{1,2}, Sven Mahner⁷ and Barbara Mayer^{1,2,*}

- ¹ Department of General, Visceral and Transplant Surgery, Ludwig-Maximilians-University Munich, Marchioninistraße 15, 81377 Munich, Germany
- ² German Cancer Consortium (DKTK), Partner Site Munich, Pettenkoferstraße 8a, 80336 Munich, Germany
- ³ Department of Obstetrics and Gynecology, Klinikum Dritter Orden, Menzinger Straße 44, 80638 Munich, Germany
- ⁴ Department of Obstetrics and Gynecology, Klinikum rechts der Isar, Technical University Munich, Ismaninger Straße 22, 81675 Munich, Germany
- ⁵ Department of Obstetrics and Gynecology, Clinic Starnberg, Oßwaldstraße 1, 82319 Starnberg, Germany
- ⁶ Department of Obstetrics and Gynecology, Clinic Harlaching, Sanatoriumsplatz 2, 81545 Munich, Germany
- ⁷ Department of Obstetrics and Gynecology, University Hospital, Ludwig-Maximilians-University Munich, Marchioninistraße 15, 81377 Munich, Germany
- * Correspondence: barbara.mayer@med.uni-muenchen.de; Tel.: +49-89-4400-76438

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Abstract: CD3⁺ and CD8⁺ lymphocytes are well known prognostic markers in primary ovarian cancer. In contrast, the predictive value of the immune infiltrate concerning treatment response and the involvement of immune heterogeneity between primary and metastatic lesions are poorly understood. In this study, the immune infiltrate of 49 primary tumors and 38 corresponding lesions in the omentum (n = 23) and the peritoneum (n = 15) was immunohistochemically analyzed and correlated with clinicopathological factors and platinum-sensitivity. Immune heterogeneity was observed between paired primary and metastatic lesions for all immune cell phenotypes. The stromal immune infiltrate was higher in the omental lesions than in the primary tumors, which was reflected by CD45 (p = 0.007), CD3 (p = 0.005), CD8 (p = 0.012), and PD-1 (programmed cell-death protein 1) (p = 0.013). A higher stromal infiltrate of both CD45⁺ and CD3⁺ cells in the omental lesions was associated with the detection of lymph node metastasis (CD45, p = 0.018; CD3, p = 0.037). Platinum-sensitive ovarian cancers revealed a higher intratumoral CD8⁺ infiltrate in the peritoneal lesions compared to the primary tumors (p = 0.045). In contrast, higher counts of stromal PD-1⁺ cells in the peritoneal lesions have been associated with reduced platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with reduced platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with platinum-sensitivity (p = 0.045). Immune heterogeneity was associated with p

Keywords: ovarian cancer; metastatic lesions; tumor microenvironment; TILs; immune heterogeneity; platinum-sensitivity; immune checkpoints

1. Introduction

High cell counts of various immune cell markers in ovarian cancer have been identified as positive prognostic factors. An especially high intratumoral infiltrate of CD3⁺ and CD8⁺ cells is associated with increased progression free survival (PFS) and overall survival (OS). This has been investigated in several studies [1,2]. Whereas the prognostic value of immune infiltrate in ovarian cancer has



been evaluated in detail, there have been less studies investigating the predictive value for treatment response. More than two-thirds of the patients experience relapse within the first three years, despite optimal surgery and adjuvant chemotherapy with carboplatin and paclitaxel [3].

Incorporation of biological drugs may improve PFS [4,5]. Tumor biology has an increasing impact in the treatment of recurrent ovarian cancer, extending the classification according to time to relapse after chemotherapy: platinum-resistant (<6 months), partially platinum-sensitive (6–12 months), and platinum-sensitive (>12 months) [6,7]. This clinical classification has been used when choosing second-line chemotherapy [8]. Further stratification factors are required for specifying first-line therapy [9]. Evaluation of the impact of the immune infiltrate on chemosensitivity might help to select the most appropriate patients for treatment with chemotherapy. In addition, the immune infiltrate might be a predictive marker for immunotherapy.

As a result of the discovery of lymphocytes as an important factor in the antitumoral defence of ovarian cancer, checkpoint inhibitors have rapidly emerged in past years in new therapeutic approaches [10–13]. Recently, various studies about treatment with checkpoint inhibitors in combination with platinum-based chemotherapy have started, investigating both primary ovarian cancer (NCT02718417, NCT02520154, NCT02766582) and the recurrent situation (NCT02891824) [14]. As none of these trials considers specific biomarkers to preselect appropriate patients, there is an obvious need to investigate the immune infiltrate in primary tumors. Because most ovarian cancer patients are diagnosed in an advanced stage [15], metastatic lesions in the peritoneum and omentum need to be analyzed in addition. While genomic, transcriptomic, and proteomic intratumoral heterogeneity are well published in ovarian cancer [16–19], immune heterogeneity has not been systematically investigated.

In the present study, various immune cell phenotypes were analyzed in the primary tumor and compared with metastatic lesions in the peritoneum and the omentum. In detail, the density of CD45⁺, CD3⁺, CD8⁺, PD-1⁺ (programmed cell-death protein 1), and PD-L1⁺ (programmed cell-death ligand 1) cells was evaluated in the stromal and intratumoral areas of the different lesions and correlated with a number of clinicopathological factors and the platinum-sensitivity. Heterogenous distribution of the immune infiltrate might impact treatment management.

2. Results

2.1. Patient Characteristics

Clinicopathological characteristics are shown in Table 1. Most patients presented with high-grade, serous ovarian carcinoma and have been diagnosed in an advanced FIGO (International Federation of Gynaecology and Obstetrics) stage, with the presence of ascites and lymphatic vessel invasion. Seventy-one percent of all patients had a complete surgical resection of all macroscopic visible tumor. In accordance with the advanced tumor stage, 82% of the patients received a carboplatin–paclitaxel-based chemotherapy. The median PFS was 19 months (range 9–42). Concerning time to relapse after chemotherapy, 14 patients were defined as reduced platinum-sensitive (tumor relapsed \leq 12 months after chemotherapy) and 28 patients as full platinum-sensitive (tumor relapsed >12 months after chemotherapy).

The presence of distant metastases at time of diagnosis was significantly associated with reduced platinum-sensitivity (p = 0.015, Table S1). The presence of ascites before surgery (p = 0.083) and macroscopic residual tumor after surgery (p = 0.067) showed a trend to reduced platinum-sensitivity.

In accordance with these results, the presence of metastases ($p_{log-rank} = 0.031$, $p_{Breslow} = 0.035$), macroscopic residual tumor after surgery ($p_{log-rank} = 0.01$, $p_{Breslow} = 0.005$), and vascular invasion ($p_{log-rank} = 0.006$, $p_{Breslow} = 0.03$) correlated significantly with shorter PFS (Figure S1).

		n or Value	%
Age	mean/median range	62/66 years 24–83 years	
FIGO Stage	I/II	0	0.0
-	III	35	71.4
	IV	14	28.6
рТ	pT2	5	10.2
	pT3	44	89.8
pN	pN0	6	12.2
-	pN1	32	65.3
	Nx	11	22.4
cM	cM0	35	71.4
	cM1	14	28.6
Primary Tumor Site	Ovarian	39	79.6
	Fallopian Tube	7	14.3
	Peritoneal	3	6.1
Histological Subtype	Serous	44	89.8
0 11	Other	5	10.2
Grading	G1/G2	1	4.0
C C	G3	47	95.9
Ascites	yes	41	83.7
	no	8	16.3
Macroscopic Residual Tumor	None	35	71.4
after Surgery	<1 cm	8	16.3
	>1 cm	6	12.2
Lymphatic Vessel Invasion	yes	26	53.1
	no	21	42.9
	missing	2	4.1
Vascular Invasion	yes	9	18.4
	no	38	77.6
	missing	2	4.1
First-Line-Treatment	С	5	10.2
	C+P	15	30.6
	C+P+B	25	51.0
	None	4	8.2
Relapse after Chemotherapy	< 6 months	2	4.1
	6–12 months	12	24.5
	>12 months	28	57.1
	none or non-sufficient chemotherapy	7	14.3

Table 1. Patient characteristics.

n: number of patients, FIGO: International Federation of Gynaecology and Obstetrics, p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, Nx: no evaluation of lymph node status, M: distant metastasis, C: Carboplatin, P: Paclitaxel, B: Bevacizumab.

2.2. Immune Infiltrate in Primary Tumor

All immune cell phenotypes were detected in the stromal area of the primary tumor in a higher fraction compared to the intratumoral area. This finding was independent from the method of evaluation. The highest density was observed for CD45⁺ cells, followed by CD3⁺ cells, CD8⁺ cells, and PD-1⁺ cells (Table 2).

-	-			
		РТ	ОМ	PE
	CD45 stromal			
Rating	Mode (Range)	3 (1–5)	3 (2–5)	3;4 (2–5)
Natilig	CD45 intratumoral			
	Mode (Range)	1 (0–3)	1 (0–3)	1 (0–3)
	CD3 stromal			
	Mean (Range)	626 (5-2491)	1241 (202–4157)	851 (117–2766
	CD3 intratumoral			
	Mean (Range)	201 (0-1134)	212 (0-569)	272 (0–985)
	CD8 stromal			
Call Count	Mean (Range)	318 (0-1049)	623 (83–1910)	364 (0-843)
Cell Coulit	CD8 intratumoral			
	Mean (Range)	88 (0-716)	104 (0-636)	130 (0–494)
	PD-1 stromal			
	Mean (Range)	73 (0-404)	91 (0-335)	130 (0-601)
	PD-1 intratumoral			
	Mean (Range)	26 (0–191)	21 (0-83)	33 (0–145)
Expression	PD-L1			
LAPICSSION	Median (Range)	1% (0–20%)	0.5% (0–20%)	3% (0–20%)

Table 2. Density and spatial distribution of immune cell phenotypes in different lesions of ovarian cancer.

n: number of analyzed tumor samples, PT: Primary tumor, OM: Omental lesion, PE: Peritoneal lesion, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. To compare primary tumor and corresponding lesions, a Wilcoxon signed-rank test was performed. Results are given in Section 2.3.

Patients who showed characteristics of tumor progression, i.e., vascular invasion, the detection of distant metastasis and the presence of ascites, at the time of diagnosis showed a strong leukocyte infiltrate. A high density (rating \geq 3) of CD45⁺ leukocytes was frequently observed in the stromal area of primary tumors with distant metastasis (cM1, 93%, Table S2). Contrary, stromal CD45⁺ cells were less frequently found in cancers without distant metastasis (cM0, 63%, *p* = 0.042). All patients with a strong (rating \geq 2) CD45⁺ intratumoral infiltrate in primary tumor were suffering from ascites (*p* = 0.006). Vascular invasion significantly correlated with a high (>73 counts/mm²) density of PD-1⁺ cells in the stromal area of the primary tumor (*p* = 0.013). PD-L1 positivity was found more often in primary tumors with distant metastasis (86%) compared to cancers without distant metastasis (51%, *p* = 0.049). A high intratumoral density (>88 counts/mm²) of CD8⁺ cells was predominantly observed in older patients (>62 years, 78%, *p* = 0.037).

Primary tumors with a high (>201 counts/mm²) intratumoral CD3⁺ cell density showed a trend to full platinum-sensitivity (p = 0.057, Table 3).

			РТ				C	OM/PT		PE/PT					
			Platin	um-Sensi	tivity *			Platir	num-Sensit	tivity *			Platin	um-Sensit	ivity *
		n	Red	Full	p #		n	Red	Full	p #		n	Red	Full	p #
		42			1		19			0.350		12			0.250
CD45 stromal	Low		4	8		OM ≤ PT		2	7		PE ≤ PT		2	9	
	High		10	20		OM > PT		5	5		PE > PT		1	0	
		42			1		19			1		12			1
CD45 intratumoral	Low		8	15		OM ≤ PT		6	11		PE ≤ PT		3	8	
	High		6	13		OM > PT		1	1		PE > PT		0	1	
		42			1		19			1		12			0.523
CD3 stromal	Low		8	16		OM ≤ PT		1	3		PE ≤ PT		2	3	
	High		6	12		OM > PT		6	9		PE > PT		1	6	
		42			0.057		19			0.633		12			1
CD3 intratumoral	Low		11	13		OM ≤ PT		2	6		PE ≤ PT		2	4	
	High		3	15		OM > PT		5	6		PE > PT		1	5	
		42			0.748		19			1		12			0.523
CD8 stromal	Low		7	16		OM ≤ PT		2	3		PE ≤ PT		1	6	
	High		7	12		OM > PT		5	9		PE > PT		2	3	
		42			1		19			0.656		12			0.045
CD8 intratumoral	Low		9	18		OM ≤ PT		2	5		PE ≤ PT		3	2	
	High		5	10		OM > PT		5	7		PE > PT		0	7	
		42			0.283		19			1		12			0.045
PD-1 stromal	Low		12	19		OM ≤ PT		2	4		PE ≤ PT		0	7	
	High		2	9		OM > PT		5	8		PE > PT		3	2	
		42			0.738		19			1		12			1
PD-1 intratumoral	Low		10	18		OM ≤ PT		5	8		PE ≤ PT		3	7	
	High		4	10		OM > PT		2	4		PE > PT		0	2	
		42			1		19			0.603		12			1
PD-L1 Positivity	No		5	10		OM ≤ PT		5	10		PE ≤ PT		2	7	
	Yes		9	18		OM > PT		2	2		PE > PT		1	2	

Table 3. Immune cell	phenotypes o	f primary tu	umor and correspo	nding lesion	ns in relation to	platinum-sensitivity.

n: number of patients, red: reduced, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. * Platinum-sensitivity was defined as follows: reduced (relapse \leq 12 months after chemotherapy) and full (relapse > 12 months after chemotherapy), # *p*-value calculated by Fisher's exact two-tailed test.

2.3. Immune Infiltrate in Metastatic Lesions

Direct comparison of primary tumors and omental lesions revealed a higher rating for stromal CD45⁺ cells in the omental lesion in more than half of the patients (52%, p = 0.007, Table S3). Furthermore, in 16 cases (70%), the omental lesion showed a higher density of stromal CD3⁺ and CD8⁺ cells compared to the primary tumor (p = 0.005 and p = 0.012, Figure 1). Consequently, the mean count in omental lesions of stromal CD3⁺ and CD8⁺ cells was nearly two times higher. In addition, the majority of omental lesions (65%) revealed a higher infiltrate of stromal PD-1⁺ cells (p = 0.013). There was no significant difference in intratumoral counts comparing primary tumors and omental lesions.



Figure 1. Scatter plots comparing immune cell phenotypes between primary tumor and the corresponding omental lesion. Counts of (**A**) CD3⁺, (**B**) CD8⁺, and (**C**) PD-1⁺ (programmed cell-death protein 1) stromal cells. Counts of CD3⁺, CD8⁺, and PD-1⁺ cells have been significantly higher in the omental lesions. *p*-value calculated by Wilcoxon signed-rank test.

In the stromal area of primary tumors and corresponding peritoneal lesions, no significant differences in the immune infiltrate could be found. In contrast, in ten cases (67%), the counts of intratumoral PD-1⁺ cells were higher in the primary tumor compared to the corresponding peritoneal lesion (p = 0.054, Figure 2). Conversely, only one case (7%) showed a higher expression of PD-L1 in the primary tumor, while it was equal or lower in most patients (93%, p = 0.074).

2.4. Associations of Immune Heterogeneity

Patients with lymph node metastases revealed a strong stromal immune infiltrate in the omental lesion. Lymph node metastases have been found at the time of diagnosis in all patients with a higher immune infiltrate of stromal CD45⁺ cells in the omental lesion compared to the primary tumor (n = 11, p = 0.018, Table S4). Most tumors with lymph node metastases (87%) revealed more stromal CD3⁺ cells in the omental lesion than in the primary tumor (p = 0.037). No significant correlations have been found for peritoneal lesions (Table S5). Interestingly, immune heterogeneity between peritoneal lesions and primary tumors could be identified as a predictive marker. All patients with reduced platinum-sensitivity showed a stronger stromal infiltrate of PD-1⁺ cells in the peritoneal lesion than in the primary tumor (p = 0.045, Table 3). In contrast, higher counts of intratumoral CD8⁺ cells in the peritoneal lesions were associated with full platinum-sensitivity (p = 0.045).



Figure 2. Scatter plots comparing immune cell phenotypes between primary tumor and the corresponding peritoneal lesion. (**A**) Counts of intratumoral PD-1⁺ (programmed cell-death protein 1) cells have been in tendencies higher in primary tumor. (**B**) PD-L1 (programmed cell-death ligand 1) expression has been slightly higher in peritoneum. *p*-value calculated by Wilcoxon signed-rank test. Cases with the value '0' in primary tumor and peritoneum (A) n = 3 (B) n = 2 have been excluded from graph, but not from calculation.

3. Discussion

The urgent need for sustainable therapeutic strategies in ovarian cancer has triggered a comprehensive search for predictive biological markers. A number of new candidates have been identified in primary tumors [20]. In contrast, corresponding metastatic lesions are rarely analyzed although it is well known that the microenvironment has a profound impact on tumor cell biology and therapeutic response [21,22]. For the first time, in the present study, a systematic side-by-side comparison of the quantitative immune infiltrate in the primary tumor and metastatic lesions located in the omentum and peritoneum was performed and correlated with clinicopathological factors. In primary tumors, the presence of a strong CD45⁺ infiltrate in both the stromal and intratumoral compartments correlated with parameters of advanced disease at time of initial diagnosis. In addition, strong PD-L1 expression in primary tumors correlated with metastatic disease. The presence of a strong stromal PD-1⁺ infiltrate was frequently observed in primary tumors characterized by vascular invasion, suggesting an immunosuppressive role in advanced primary ovarian cancer. Indeed, a high expression of PD-1 and PD-L1 was reported to be associated with poor prognosis [23–25]. An increased intratumoral CD3⁺ infiltrate in chemo-naive primary tumors was associated with response to platinum-based chemotherapy. This supports the 2011 study from Bösmüller et al., which indicated intratumoral lymphocyte density superior to ERCC-1 (excision repair cross-complementation group 1) expression to predict platinum response [26]. A number of different mechanisms is known to be involved in an increased antitumoral activity of lymphocytes induced by platinum-based chemotherapy [27,28]. Consequently, a pre-therapeutic increased number of CD3⁺ cells might be associated with a higher efficacy. Methods to enhance this effect could be the application of dose-dense chemotherapy [29]. Stratification of primary tumors according to a strong CD3⁺ infiltrate and a high PD-1/PD-L1 expression might identify ovarian cancer patients who are responsive to a combination of platinum-based chemotherapy with checkpoint-inhibitors [30,31].

In more than 70% of the patients, disease has already spread in peritoneal and omental tissue at the time of diagnosis, which is associated with an unfavorable prognosis [15,32]. A higher fraction of CD45⁺, CD3⁺, CD8⁺, and PD-1⁺ cells was observed in the stroma of omental lesions compared to the corresponding primary tumors. Higher counts of stromal CD45⁺ and CD3⁺ cells in the omental lesions were associated with the presence of lymph node metastases. Indeed, aggregates of leukocytes named milky spots are characteristic for the omental tissue and were found to promote immune suppression and, in consequence, facilitate tumor cell implantation [33,34]. They are discussed as a potential target for immunotherapy [35]. Combining strategies that enforce intratumoral recruitment of CD8⁺ cells might, therefore, increase immunotherapeutic effects [36–38].

Similar to omental lesions, comparison between peritoneal lesions and their corresponding primary tumors identified changes in the immune cell infiltrate, i.e., less intratumoral PD-1⁺ cells and a higher PD-L1 expression in the peritoneal metastases. Moreover, peritoneal lesions were characterized by a higher infiltration of intratumoral CD8⁺ cells, which correlated with the response to the platinum-based first-line treatment. For example, in patient 5082, a high density (30/mm²) of intratumoral CD8⁺ cells was found in the peritoneal lesion, while there was a low intratumoral CD8⁺ cells density (7/mm²) in the corresponding primary tumor. In this individual patient platinum-based chemotherapy was successful, resulting in a relapse-free time of 26 months after chemotherapy. Conversely, in the peritoneal lesion of patient 5072, a low number (2/mm²) of intratumoral CD8⁺ cells was found, while a high infiltrate (46/mm²) was found in the primary tumor. This patient suffered from a relapse within six months after platinum-based chemotherapy. Thus, for patients with a low peritoneal infiltrate, other therapeutic options than platinum-based chemotherapy should be considered right from the start. Treatment options might be non-platinum chemotherapeutic drugs or targeted therapies.

Conversely to the findings concerning intratumoral CD8⁺ cells, a higher density of stromal PD-1⁺ cells in the peritoneal lesions compared to the primary tumors was associated with a reduced platinumsensitivity, which might be explained by impairment of these cells [23,24]. This further supports the close correlation between immune heterogeneity and platinum response. Comparing the immune contexture, defined as density, composition, and functionality of the immune infiltrate [39], in omental, peritoneal, and primary lesions, profound differences were observed. These changes might go along with dynamic variations in the tumor microenvironment at the different locations. In fact, the peritoneal cavity is characterized by complex multicellular interactions regulated by a network of multiple soluble factors [22,40,41].

Although the cohort of the present study can be described as representative and is comparable to cohorts of other clinical trials [42–44], the main limitation is the small number of patients. In particular, the number of samples of omental and peritoneal tissue was low. To confirm the results of the current pilot study, further analysis of immune heterogeneity should be pursued in a larger cohort. This study indicates that immune heterogeneity represents an important characteristic of ovarian cancer. Location-dependent changes in the immune cell infiltrate were identified as potential predictive markers for standard chemotherapy and immunotherapy. Thus, the systematic analysis of the immune contexture in both primary tumors and the corresponding metastatic lesions could have a relevant impact on treatment planning for individual cancer patients.

4. Patients and Methods

4.1. Study Population

Forty-nine patients diagnosed with a primary, chemo-naive ovarian, fallopian tube, or peritoneal cancer from the SpheroID-Study were included. Patients suffering from another neoplasia within the last five years were excluded. Informed consent was obtained from all patients. The study was approved by the institutional review board of the Ludwig-Maximilians-Universität Munich, Munich, Germany (No.278/04). Patients were recruited between September 2012 and January 2015 in five ovarian cancer centers, namely University Hospital, LMU Munich (n = 16), Klinikum Dritter Orden

(*n* = 16), Klinikum rechts der Isar, Technical University Munich (*n* = 7), Clinic Harlaching (*n* = 5), and Clinic Starnberg (*n* = 5). Standardized surgical resection and pathological analysis was conducted by the recruiting clinic. In 33 cases, in addition to primary tumor samples, corresponding omental or peritoneal lesions were resected. A detailed description of the tumor samples analyzed is given in Table S6. Patient-, tumor-, and treatment-related data for correlations was given in the routine reports and delivered in a pseudonymized form. Subsequently, for each clinicopathological parameter, patients were divided according to clinical relevance into two groups. For the parameter 'age', the mean value was chosen as cutoff. Analysis of platinum-sensitivity was performed after completion of chemotherapy [7,45]. Seven patients with no chemotherapy or a reduced number of treatment cycles (≤ 2) had to be excluded. In the present cohort, patients with platinum-resistant disease (*n* = 2) and patients with partially platinum-resistant disease (*n* = 12) were grouped and defined as reduced platinum-sensitive. This group was compared to the full platinum-sensitive group (*n* = 28) [6,7]. PFS was defined as the time from surgical treatment to the time of relapse or progression. As the median follow-up time was 26 months, OS has not been analyzed. Data from patients who did not die and had no relapse or progression was censored at the date of their last visit.

4.2. Immunohistochemistry

After surgical removal, tumor samples were snap frozen in liquid nitrogen. Serial cryosections (5 µm) were performed, fixed in acetone, and stained immunohistochemically using the avidin-biotinperoxidase complex method [46]. Briefly, nonspecific binding Fc-regions were saturated with 10%-AB-Serum in phosphate-buffered-saline (Bio-Rad, Hercules, CA, USA, 805135) for 20 min. Endogenous biotin was blocked by the Avidin-/Biotin-Blocking-Kit (Vector Laboratories, Burlingame, CA, USA, SP-2001) according to the manufacturer. Primary antibodies were incubated for 60 min at room temperature, namely clone 2B11 + PD7/26 directed against CD45 (Dako, Santa Clara, CA, USA, M0701, working concentration [wc] 4.5 µg/mL), clone UCHT1 directed against CD3 (BD Biosciences, San Jose, CA, USA, 550368, wc 1.25 µg/mL), clone C8/144B directed against CD8 (Dako, M7103, wc 3 µg/mL), clone MIH4 directed against PD-1 (affymetrix eBiosciences, San Diego, CA, USA, 14-9969, wc 10 µg/mL) and clone MIH1 directed against PD-L1 (affymetrix eBiosciences, 14-5983, wc 10 µg/mL). The antibody Ber-EP4 directed against EpCAM (Dako, M0804, wc 2.5 µg/mL) was used to define tumor areas. The antibody MOPC21 (Sigma-Aldrich, St. Louis, MO, USA, M9269) was used as Immunoglobulin G1 isotype control in the relevant wc. Antibodies directed against CD45, CD3, and EpCAM were incubated with the secondary biotinylated antibody (Jackson ImmunoResearch, Cambridgeshire, U.K., 315-065-048, wc 0.75 µg/mL) for 30 min and, subsequently, with peroxidase-conjugated streptavidin (Jackson ImmunoResearch, 016-030-084, wc 1 µg/mL) for 30 min. Antibodies directed against CD8, PD-1, and PD-L1 were incubated with the ZytoChem Plus HRP-Kit (Zytomed Systems, Berlin, Germany, HRP060), as recommended by the manufacturer. Antigen–antibody reaction was visualized by 3-amin-9-ethylcarabazol (AEC, Sigma-Aldrich, A5754)-peroxide-solution for eight min in darkness. Sections were counterstained with Mayers hemalum solution (Merck, Darmstadt, Germany, 109249).

4.3. Semiquantitative Analysis of the CD45⁺ Infiltrate

CD45⁺ cells were detected as single cells and often in cell clusters of different size. This staining pattern could not be dissected precisely by the software ImageJ (Version 1.51h, National Institutes of Health, Bethesda, MD, U.S). Therefore, the CD45 staining was semiquantitatively rated in two different spatial areas (Table S7). In the intratumoral area, CD45⁺ cells were found in direct contact with the cancer cells. In the stromal area adjacent to the tumor compartment, CD45⁺ cells did not touch the tumor cells. Analysis was performed with the microscope (BX41, Olympus Corporation, Tokio, Japan).

4.4. Quantitative Analysis of Immune Cells

Quantitative analysis was performed for CD3, CD8, and PD-1. For each tissue section and each immune cell antigen, three separate regions characterized by an enriched infiltrate were selected,

separately for the stromal and the intratumoral area (defined previously and demonstrated in Figure 3). CD8⁺ cells were counted in the same region as CD3⁺ cells. For PD-1, known to be expressed on different cell types [47], enriched regions were evaluated independently from the CD3/CD8 area. A picture of each enriched region was captured (×200 magnification) using Zen 2.0 lite software (Carl Zeiss Inc., Oberkochen, Germany) with an AxioCam MRc5 camera (Carl Zeiss Inc., Oberkochen, Germany).



Figure 3. Immunohistochemical analysis demonstrating intratumoral (**A**) and stromal (**B**) $CD3^+$ cells. The black borders mark the enriched regions, in which $CD3^+$ cells were counted. Tumor and stroma areas are indicated. Magnification 200×.

Counting was done according to the QTiS algorithm by Miksch et al. [48–51] with the ImageJ software (Version 1.51h, National Institutes of Health, Bethesda, MD, USA). Briefly, after substraction of background, color deconvolution was performed, selecting example regions for the following spectra: background, hematoxylin, and AEC. In the AEC-spectrum an automatic threshold was applied to get a binary image. Watershed method was used to separate cell clusters. Cells were counted with the 'Analyze-Particle-Tool', considering the values of '700-infitiy' for size and '0.2–1.0' for circularity. The 'ROI-Manager' was used to distinguish between intratumoral and stromal area. For each tissue section, an average count of the three images that were taken was calculated. Cell counts were normalized to an average number of cells/mm².

The reliability of the semiquantitative rating system for CD45⁺ cells was checked by correlation of ratings and counts of CD3⁺ cells intratumoral and stromal. Spearman correlation coefficient was 0.803 (p = 0.000).

4.5. Semiquantitative Analysis of PD-L1

The fraction of all PD-L1⁺ cells, including both cancer and immune cells in the intratumoral area, was estimated. PD-L1 positivity was defined as \geq 1%. This approach has already been used in former clinical trials [52,53].

4.6. Statistical Analysis

Clinicopathological factors were grouped by clinical relevance. For the immune cell phenotypes CD3, CD8, and PD-1 identified in the respective spatial area the mean cell count was defined as cutoff to distinguish low infiltrates from high infiltrates. The cutoff for CD45⁺ cells was chosen considering balanced group sizes (intratumoral: ≤ 1 ; >1; stromal: ≤ 2 ; >2). For comparison between corresponding location and primary tumor, the ratio between the cell counts was calculated for each immune cell phenotype in both spatial areas. To form two groups—'corresponding location lower/equal than primary tumor' and 'corresponding location higher than primary tumor'— ≤ 1 ; >1 was chosen as the cutoff. The defined cutoffs were correlated with various clinicopathological data and platinum-sensitivity using the Fisher's exact two-tailed test. Cumulative survival probabilities were calculated by the Kaplan–Meier

method. Log-rank and Breslow tests were used to compare clinicopathological factors regarding PFS. Immune cell phenotypes between the primary tumor and different corresponding locations were compared by Wilcoxon signed-rank test. *p*-values < 0.050 were considered to be statistically significant; *p*-values < 0.100 have been reported. All statistical analyses were performed using SPSS Statistics (Version 23.0, IBM, Armonk, NY, USA).

5. Conclusions

Immunohistochemical comparison in ovarian cancer between primary tumors and the corresponding omental and peritoneal lesions demonstrated profound differences in the expression pattern of CD45, CD3, CD8, PD-1 and PD-L1. This immune heterogeneity made an impact on platinum-sensitivity. Variations in the immunologic tumor microenvironment might influence treatment selection for the individual ovarian cancer patient.

Supplementary Materials: The following are available online at http://www.mdpi.com/2072-6694/11/9/1250/s1, Figure S1: Kaplan-Mayer-Curves for PFS of different clinicopathological parameters, Table S1: Clinicopathological factors in relation to platinum-sensitivity. Table S2: Immune cell phenotypes in primary tumor in relation to clinicopathological factors. Table S3: Comparison of the immune cell phenotypes between primary tumor and corresponding lesions. Table S4: Ratios of immune cell phenotypes between primary tumor and omental lesion in relation to clinicopathological factors. Table S5: Ratios of immune cell phenotypes between primary tumor and omental lesion in relation to clinicopathological factors. Table S5: Ratios of immune cell phenotypes between primary tumor and corresponding lesions in relation to clinicopathological factors. Table S5: Ratios of immune cell phenotypes between primary tumor and corresponding lesions in relation to clinicopathological factors. Table S5: Ratios of immune cell phenotypes between primary tumor and corresponding lesion in relation to clinicopathological factors. Table S5: Ratios of immune cell phenotypes between primary tumor and corresponding lesions per patient., Table S7: Description of rating method.

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Supplementary materials



Figure S1. Kaplan-Mayer-Curves for progression free interval (PFS) of different clinicopathological parameters. (A) Age (B) pT (C) pN (D) cM (E) Primary Tumor Site (F) Histological Subtype (G) Grading (H) Ascites (I) Macroscopic Residual Tumor (J) Lymphatic Vessel Invasion (K) Vascular Invasion. p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, M: distant metastasis. *p*-values calculated by log-rank-test and Breslow-test.

		Platinum-S	ensitivi	ty*
	n	reduced	full	p^{*}
Age	42			0.191
≤ 62 years		4	15	
> 62 years		10	13	
рТ	42			0.668
<pt3c< td=""><td></td><td>3</td><td>4</td><td></td></pt3c<>		3	4	
рТ3с		11	24	
pN	33			0.559
pN0		0	5	
pN1		7	21	
cM	42			0.015
cM0		6	23	
cM1		8	5	
Primary Tumor Site	42			0.668
Ovarian		11	24	
Other		3	4	
Histological Subtype	42			0.1
Serous		11	27	
Other		3	1	
Grading	42			1
G1/G2		1	1	
G3		13	27	
Ascites	42			0.083
No		0	6	
Yes		14	22	
Macroscopic Residual Tumor after Surgery	42			0.067
Absent		7	23	
Present		7	5	
Lymphatic Vessel Invasion	40			0.521
No		7	10	
Yes		7	16	
Vascular Invasion	40			0.159
No		10	24	
Yes		4	2	

Table S1. Clinicopathological factors in relation to platinum-sensitivity.

n: number of patients, p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, M: distant metastasis. *Platinum-sensitivity was defined as follows: reduced (relapse ≤ 12 months after chemotherapy) and full (relapse >12 months after chemotherapy), **p*-value calculated by Fisher's exact two-tailed test.

Table S2. Immune cell phenotypes in primary tumor in relation to clinicopathological factors.

	Age ≤ >6				Age pT				pT			рN		(сM		Primar S	y Tur Site	nor	Η	listolo Subt	ogica type	ıl		Grac	ling		Asc	cites		Macro	scopic Res Tumor	idual	Ly	mph/ In	iytic V vasioi	/essel n	V Iı	ascu Ivasi	lar ion
	n 6	≤ >6 2 2	<i>p</i> * n	<pt 3c</pt 	pT3 c	p [≠] r	۳ ^{рN} 0	pN 1	<i>p</i> * n	сМ 0	cM 1	p [≠] n	Ovari an	Oth er	<i>p</i> *	n S	ero (us	Oth er	<i>p</i> [#]	n	G1/ G2	G 3	<i>p</i> * n	n y o	ye s	p#	n abse	nt presen	t <i>p</i> [#]	n	no	yes	<i>p</i> *	n n	ye s	<i>p</i> *				
CD45 stromal	4 9	().54 4 2 9			0.41 8	3 3		0.15 4 4 9			0.04 4 2 9			0.70 2	4 9			0.61 6	4 9		().49 4 4 9	ŧ)		1	49		1	47			0.528	4 7		0.24				
Low	5	77		1	13		0	11		13	1		12	2			12	2			1	13		2	12		10	4			5	9		13	3 1					
High	1	4 21		7	28		6	21		22	13		27	8		3	32	3			1	34		6 2	<u>2</u> 9		25	10			16	17		2	58					
CD45	4	().77 4			1 3	3		0.66 4			1 4			1	4			1	4			1 4	ł	0	.00	10		1	47			0 220	4		0.48				
intratumoral	9		99			1 8	3		39			¹ 9			1	9			1	9			1 9)		6	±7		1	47			0.239	7		6				
Low	1	1 16		4	23		4	16		19	8		21	6		2	24	3			1	26		8	19		19	8			14	12		22	2 4					
High	1	0 12		4	18		2	16		16	6		18	4		2	20	2			1	21		0 2	22		16	6			7	14		1	55					
CD3 stromal	4 9	().14 4 4 9			0.43 3	3 3		0.68 4 2 9			0.22 4 2 9			1	4 9			0.37 6	4 9			0.5 g	1)	0	.71	49		0.222	47			0.374	4 7		0.46 5				
Low	1	5 13		6	22		3	19		22	6		22	6		2	24	4			2	26		4 2	24		22	6			14	13		23	3 4					
High	e	5 15		2	19		3	13		13	8		17	4		2	20	1			0	21		4	17		13	8			7	13		1	55					
CD3	4	().24 4			0.71 3	3		0.39 4			0.52 4			0.15	4			0.05	4			1 4	ł		1	40		0.207	477			0.07(4		0.26				
intratumoral	9		69			5 8	3		59			0.53 9			2	9			6	9			1 9)		1	49		0.207	47			0.076	7		3				
Low	1	4 13		5	22		4	14		18	9		19	8		2	22	5			1	26		4 2	23		17	10			15	11		23	3 3					
High	5	7 15		3	19		2	18		17	5		20	2		2	22	0			1	21		4	18		18	4			6	15		15	56					
CD8 stromal	4 9	().77 4 1 9			0.43 3	3 3		0.39 4 5 9			0.22 4 2 9			0.72 6	4 9			0.37 6	4 9			1 4	1)	0	.26 3	49		0.542	47			1	4 7		0.46 5				
Low	1	3 15		6	22		2	18		22	6		23	5		2	24	4			1	27		3 2	25		21	7			12	15		2	34					
High	8	3 13		2	19		4	14		13	8		16	5		2	20	1			1	20		5	16		14	7			9	11		1	55					
CD8	4	(0.03 4			0.69 3	3		. 4			0.52 4			0.72	4			0.14	4			. 4	ł			10			4.5			0 = (0	4		0.27				
intratumoral	9		79			3 8	3		1 9			79			6	9			3	9			1 9)		1	49		0.744	47			0.562	7		4				
Low	1	7 14		6	25		4	20		21	10		24	7		2	26	5			1	30		5 2	26		23	8			14	15		2	54					
High	4	4 14		2	16		2	12		14	4		15	3		-	18	0			1	17		3	15		12	6			7	11		13	35					
PD-1 stromal	4 9		$1 \frac{4}{9}$			0.20 3	3 3		0.15 4 4 9			$1 \frac{4}{9}$			0.70	4 9			1	4 9			1 9	ł)		1	49		0.294	47			0.528	4 7		0.01				
Low	1	5 20		4	31		2	23		25	10		27	8		3	31	4			2	33		6 2	29		23	12			16	17		30) 3					
High	6	58		4	10		4	9		10	4		12	2			13	1			0	14		2	12		12	2			5	9		8	6					
PD-1	4	,	4			0.41 3	3		0.39 4			0.50 4			0.46	4				4			. 4	ł			10			4.5				4		0.12				
intratumoral	9	().76 9			1 8	3		29			19			4	9			1	9			1 9)		1	49		1	47			0.355	7		1				
Low	1	5 18		4	29		3	22		25	8		25	8		2	29	4			2	31		6 2	27		23	10			16	16		28	3 4					
High	6	5 10		4	12		3	10		10	6		14	2			15	1			0	16		2	14		12	4			5	10		1) 5					
PD-L1	4	().37 4			1 3	3		o o 7 4			0.04 4			1	4			4	4		().14 4	ł		4	10		1	477			0 885	4		0.27				
Positivity	9		69			1 8	3		0.37 9			9 9			1	9			1	9			5 9)		1	49		1	47			0.775	7		8				
No	1	09		3	16		1	15		17	2		15	4			17	2			2	17		3	16		14	5			9	10		12	72					
Yes	1	1 19		5	25		5	17		18	12		24	6		2	27	3			0	30		5 2	25		21	9			12	16		2	l 7					

n: number of patients, p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, M: distant metastasis, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. * *p*-value calculated by Fisher's exact two-tailed test.

	n PT > OM	n PT = OM	n PT < OM	$p^{\#}$	n PT > PE	n PT = PE	n PT < PE	p^{*}
CD45 stromal	2	9	12	0.007	3	9	3	0.739
CD45 intratumoral	3	18	2	0.655	4	10	1	0.48
CD3 stromal	7	0	16	0.005	4	1	10	0.331
CD3 intratumoral	8	1	14	0.221	8	0	7	0.82
CD8 stromal	7	0	16	0.012	8	0	7	0.82
CD8 intratumoral	7	1	15	0.131	4	2	9	0.124
PD-1 stromal	7	1	15	0.013	8	1	6	0.95
PD-1 intratumoral	9	7	7	0.569	10	3	2	0.054
PD-L1 Expression	4	15	4	0.944	1	10	4	0.078

Table S3. Comparison of the immune cell phenotypes between primary tumor and corresponding lesions.

n: number of analyzed tumor samples, PT: Primary tumor, OM: Omental lesion, PE: Peritoneal lesion, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. *p*-value calculated by Wilcoxon signed-rank test.

	Ag n ≤ ≻				р	т			рN			c	м		Prin	nary T	umor	Site		Histol Subt	ogical type		G	radin	ıg		Asci	tes		Macrosco T	opic Resid Tumor	lual	I	.ympł In	ytic V vasion	essel	Va Inv	scular vasion
	n	≤ >6 52 2	5 p*	n [^]	c c	pT3 c	p^*	n ph 0	V pN 1	p^*	n	сМ 0	cM 1	p^* 1	n O	varia n	Othe r	p^{*}	n	Serous	Other	r <i>p*</i> r	n G1/ 2	G G 3	p^{*}	n	nyo os	е <i>p</i> [*]	n	absent	present	p^*	n	no	yes	p^*	n no	yes p*
CD45 stromal	2 3		0.19 3	2 3			1	1 9		0.01 8	2 3			0.59	2 3			0.06 9	23			1 3	<u>2</u> 3		0.4 8	72 3		0.59 0	23			1	22			0.395	22	0.476
OM ≤ PT OM > PT		29 66			1 1	10 11		4 0	4 11			9 11	2 1			6 11	5 1			10 10	1 2		0 2	11 10	1)		2 9 1 1	L		8 8	3 4			7 4	4 7		11 9	0 2
CD45 intratumoral	2 3		1	2 3			1	1 9		1	2 3			0.24 2	2 3			1	23			1 3	2 3		0.1	72 3		1	23			1	22			1	22	1
OM≤PT OM>PT		$\begin{array}{ccc} 7 & 14 \\ 1 & 1 \end{array}$	L		2 0	19 2		4 0	13 2			19 1	2 1			15 2	6 0			18 2	3 0		1 1	20 1)		3 18 0 2	3		14 2	7 0			10 1	11 0		19 1	2 0
CD3 stromal	2 3			2 3				1 9		0.03 7	2 3			2	2 3				23			1 2	2 3		1	2 3		1	23			1	22			1	22	1
OM≤PT OM>PT		16 79			0 2	7 14		3 1	2 13			6 14	1 2			4 13	3 3			6 14	1 2		0 2	7 14	, 1		1 6 2 14	1		5 11	2 5			4 7	3 8		6 14	1 1
CD3 intratumoral	2 3		0.08 6	2 3			0.14 2	1 9		0.60 3	2 3			1	2 3			0.64 3	23			1 3	<u>2</u> 3		0.5 2	0 2 3		0.50 8	3 23			0.176	22			0.659	22	0.515
OM ≤ PT OM > PT		18 77			2 0	7 14		2 2	5 10			8 12	1 2			6 11	3 3			8 12	1 2		0 2	9 12	2		2 7 1 13	3		8 8	1 6			5 6	3 8		8 12	0 2
CD8 stromal	2 3		1	2 3			1	1 9		0.27 2	23			0.52	2 3			0.31 8	23			1 3	2 3		1	2 3		0.20 9	23			0.626	22			1	22	1
OM≤PT OM>PT		25 61()		0 2	7 14		2 2	3 12			7 13	0 3			4 13	3 3			6 14	1 2		0 2	7 14	, 1		2 5 1 1	5		4 12	3 4			4 7	3 8		6 14	1 1
CD8 intratumoral	2 3		0.17 6	2 3			0.11 1	1 9		0.55 7	2 3			1	2 3			0.62 1	23			1 3	2 3		1	2 3		0.2 9	⁵ 23			0.345	22			1	22	0.515
OM ≤ PT OM > PT		17 78			2 0	6 15		2 2	4 11			7 13	1 2			5 12	3 3			7 13	1 2		1 1	7 14	, 1		2 6 1 14	1		7 9	1 6			4 7	4 7		8 12	0 2
PD-1 stromal	2 3		0.65 7	2 3			1	1 9		0.07 1	2 3			1	2 3			0.62 1	23			1 3	2 3		0.5 6	22 3		1	23			0.657	22			1	22	0.121
OM≤PT OM>PT		26 69			1 1	7 14		3 1	3 12			7 13	1 2			5 12	3 3			7 13	1 2		0 2	8 13	3		1 7 2 13	3		5 11	3 4			4 7	4 7		6 14	2 0
PD-1 intratumoral	2 3		1	2 3			1	1 9		0.25 5	2 3			0.52	2 3			0.62 1	23			1 2	<u>2</u> 3		0.5 6	22 3		1	23			1	22			1	22	1
OM≤PT OM>PT		6 10 2 5)		2 0	14 7		4 0	9 6			13 7	3 0			11 6	5 1			14 6	2 1		1 1	15 6	5		2 14 1 6	1		11 5	5 2			8 3	7 4		13 7	2 0
PD-L1 Positivity	2 3		0.25 7	2 3			1	1 9			2 3			0.45	2 3			0.53 9	23			1 3	<u>2</u> 3		0.3 4	22 3		1	23			1	22			0.586	22	1
OM≤PT		8 11			2	17		3	13	0.53 0		17	2			13	6			16	3		1	18	3		3 10	5		13	6			10	8		16	2
OM > PT		0 4			0	4		1	2			3	1			4	0			4	0		1	3			0 4			3	1			1	3		4	0

Table S4. Ratios of immune cell phenotypes between primary tumor and omental lesion in relation to clinicopathological factors.

n: number of patients, PT: Primary tumor, OM: Omental lesion, p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, M: distant metastasis, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. * *p*-value calculated by Fisher's exact two-tailed test.

		Ag	e			рТ			pN			сM		P	rimary	Tumor	Site		Histol Subt	ogical type	Grad	ling		A	scite	s	N	Macrosco T	opic Resid 'umor	ual	I	.ympl Ir	nytic V Ivasion	essel	V Ir	ascul vasio	ar on
	n	≤ > 52 2	$\frac{6}{p}$	n	<pt3 c</pt3 	pT3 c	p^{*}	n	pN pl 0 1	N <i>p</i>	1 cM 0	[cM 1	p^*	n	Ovaria n	Othe r	p^*	n	Serous	Other p^* 1	G1/G	G G 3	p ; n	n o	ye s	p^*	n	absent	present	p^{z}	n	no	yes	p^*	n no	yes	p^*
CD45 stromal	1 5		1	1 5			0.37 1	7 1 1		/	l 5		1	1 5			0.51 6	15		/ 1	5		/ 1	;		0.24 2	15			1	14			0.176	14		1
PE ≤ PT		6 6	5		1	11			10)	7	5			10	2			12			12		3	9			9	3			2	9		9	2	
PE > PT		1 2	2		1	2			1		2	1			2	1			3			3		2	1			2	1			2	1		2	1	
CD45	1		0.4	6 1			1	1		, .	L		1	1			1	15		, 1	l		, 1			0.33	15			1	14			1	14		1
intratumoral	5		7	5			-	1		1	5		-	5			-	10		1 5	5		' 5	5		3	10			1				1	11		1
PE ≤ PT		6 8	3		2	12			10)	8	6			11	3			14			14		4	10			10	4			4	9		10	3	
PE > PT		1 ()		0	1			1		1	0			1	0			1			1		1	0			1	0			0	1		1	0	
CD3 stromal	1 5		0.2	8 I 5			0.52 4	2 1		/	5		0.32 9	2 1 5			1	15		/ 5	5		/ 5	;		1	15			1	14			0.520	14		1
$PE \le PT$		1 4	Ł		0	5			4		2	3			4	1			5			5		2	3			4	1			2	2		3	1	
PE > PT		6 4	Ł		2	8			7		7	3			8	2			10			10		3	7			7	3			2	8		8	2	
CD3 intratumoral	1^{1}_{5}		0.6 9	11 5			0.46 7	5 1 1		/	l 5		0.11 9	l 1 5			1	15		1 5	5		1 5	;		1	15			0.569	14			0.559	14		1
PE ≤ PT		3 5	5		2	6			4		3	5			6	2			8	/		8	/	3	5			5	3			3	4		5	2	
PE > PT		4 3	3		0	7			7		6	1			6	1			7			7		2	5			6	1			1	6		6	1	
CD8 stromal	1 5		0.6 9	1 1 5			0.2	1 1		/	L 5		0.60 8) 1 5			0.2	15		/ 5	5		/ 1	5		1	15			0.569	14			0.559	14		1
PE ≤ PT		3 5	5		0	8			6		4	4			5	3			8			8		3	5			5	3			3	4		5	2	
PE > PT		4 3	3		2	5			5		5	2			7	0			7			7		2	5			6	1			1	6		6	1	
CD8 intratumoral	$1 \frac{1}{5}$		0.1 9	11 5			0.48 6	3 1 1		/	l 5		0.62 2	2 1 5			0.52 5	15		/ 1	5		/ 1	;		0.32 9	15			1	14			0.245	14		1
PE≤PT		1 5	5		0	6			4		3	3			4	2			6			6		3	3			4	2			3	3		5	1	
PE > PT		6 3	3		2	7			7		6	3			8	1			9			9		2	7			7	2			1	7		6	2	
PD-1 stromal	1 5		0.1 9	1 1 5			0.48 6	3 1 1		/	l 5		0.62 2	2 1 5			0.22 9	15		/ 1	5		/ 1	;		0.58 0	15			0.604	14			0.085	14		0.538
PE ≤ PT		6 3	3		2	7			6		6	3			6	3			9			9		4	5			6	3			4	4		7	1	
PE > PT		1 5	5		0	6			5		3	3			6	0			6			6		1	5			5	1			0	6		4	2	
PD-1	1		0.2	0 1			1	1		, .	L		0.48	31			0.37	15		, 1	Į		, 1			1	15			1	14			0.505	14		1
intratumoral	5		0	5			1	1		/ <u></u>	5		6	5			1	15		1 5	5		′ 5	5		1	15			1	14			0.505	14		1
$PE \le PT$		5 8	3		2	11			9		7	6			11	2			13			13		4	9			9	4			3	9		9	3	
PE > PT		2 ()		0	2			2		2	0			1	1			2			2		1	1			2	0			1	1		2	0	
PD-L1 Positivity	1 5		1	1 5			1	1 1		/	1 5		0.60 4) 1 5			1	15		/ 1	5		/ 1 5	;		0.23 1	15			1	14			0.251	14		0.505
PE ≤ PT		5 6	5		2	9			8		6	5			9	2			11			11		5	6			8	3			4	6		7	3	
PE > PT		2 2	2		0	4			3		3	1			3	1			4			4		0	4			3	1			0	4		4	0	

Table S5. Ratios of immune cell phenotypes between primary tumor and peritoneal lesion in relation to clinicopathological factors.

n: number of patients, PT: Primary tumor, PE: Peritoneal lesion, p: pathological, c: clinical, T: extent of primary tumor, N: regional lymph node metastasis, M: distant metastasis, PD-1: programmed cell-death protein 1, PD-L1: programmed cell-death ligand 1. * p-value calculated by Fisher's exact two-tailed test.

Detion t Marsher	Definition Transaction According to Dath desired Demont	Location of Corresp	onding Lesions
Patient Number	Primary Tumor Site—According to Pathological Report	Omentum	Peritoneum
5064	Ovary	Greater Omentum	
5085	Ovary		Peritoneum*
5188	Ovary	Greater Omentum	
4747	Ovary	Greater Omentum	Sigma
4757	Ovary		Peritoneum*
4976	Ovary		Peritoneum*
5074	Ovary	Greater Omentum	
5146	Ovary	Greater Omentum	
5196	Ovary	Greater Omentum	
5199	Ovary	Greater Omentum	
5243	Ovary	Greater Omentum	
5306	Ovary	Greater Omentum	Peritoneum*
4751	Fallopian Tube	Greater Omentum	
4754	Ovary		Peritoneum*
5072	Ovary	Greater Omentum	Sigma
5082	Fallopian Tube		Sigma
5097	Ovary	Greater Omentum	
5170	Fallopian Tube	Greater Omentum	
5216	Fallopian Tube	Greater Omentum	
5264	Ovary	Greater Omentum	Diaphragm
5266	Fallopian Tube	Greater Omentum	
5277	Peritoneum	Greater Omentum	Uterus
5281	Peritoneum	Greater Omentum	Diaphragm
4843	Ovary	Greater Omentum	
4940	Ovary		Peritoneum*
5075	Ovary		Liver Capsule
5242	Ovary	Greater Omentum	
5070	Ovary	Greater Omentum	
5088	Ovary	Greater Omentum	
5103	Ovary		Sigma
5121	Ovary	Greater Omentum	
5137	Ovary		Peritoneum*

Table S6. Description of primary tumor site and corresponding lesions per patient.

* not further specified.

 Table 7. Description of rating method.

Rating	Individual Cells	Cluster
0	none	none
1	scattered	none
2	a few	none
3	a few	small, locally restricted
4	many	small and big, locally restricted
5	many	big, confluent





Article Integrin α2β1 Represents a Prognostic and Predictive Biomarker in Primary Ovarian Cancer

Katharina Dötzer¹, Friederike Schlüter¹, Franz Edler von Koch², Christine E. Brambs³, Sabine Anthuber⁴, Sergio Frangini ⁵, Bastian Czogalla^{6,7}, Alexander Burges^{6,7}, Jens Werner^{1,7}, Sven Mahner^{6,7} and Barbara Mayer^{1,7,*}

- ¹ Department of General, Visceral and Transplant Surgery, University Hospital, Ludwig-Maximilians-University Munich, Marchioninistraße 15, 81377 Munich, Germany; katharina.doetzer@med.uni-muenchen.de (K.D.); friederike1.schlueter@med.uni-muenchen.de (F.S.); jens.werner@med.uni-muenchen.de (J.W.)
- ² Gynecology and Obstetrics Clinic, Klinikum Dritter Orden, Menzinger Straße 44, 80638 Munich, Germany; franz.koch@dritter-orden.de
- ³ Department of Obstetrics and Gynecology, Klinikum Rechts der Isar, Technical University Munich, Ismaninger Straße 22, 81675 Munich, Germany; christine.brambs@tum.de or christine.brambs@luks.ch
- ⁴ Department of Obstetrics and Gynecology, Starnberg Hospital, Oßwaldstraße 1, 82319 Starnberg, Germany; sabine.anthuber@klinikum-starnberg.de
- ⁵ Department of Obstetrics and Gynecology, Munich Clinic Harlaching, Sanatoriumsplatz 2, 81545 Munich, Germany; sergio.frangini@muenchen-klinik.de
- ^b Department of Obstetrics and Gynecology, University Hospital, Ludwig-Maximilians-University Munich, Marchioninistraße 15, 81377 Munich, Germany; bastian.czogalla@med.uni-muenchen.de (B.C.);
- alexander.burges@med.uni-muenchen.de (A.B.); sven.mahner@med.uni-muenchen.de (S.M.)
 ⁷ German Cancer Consortium (DKTK), Partner Site Munich, Pettenkoferstraße 8a, 80336 Munich, Germany
- Correspondence: barbara.mayer@med.uni-muenchen.de; Tel.: +49-89-4400-76438

Abstract: Currently, the same first-line chemotherapy is administered to almost all patients suffering from primary ovarian cancer. The high recurrence rate emphasizes the need for precise drug treatment in primary ovarian cancer. Being crucial in ovarian cancer progression and chemotherapeutic resistance, integrins became promising therapeutic targets. To evaluate its prognostic and predictive value, in the present study, the expression of integrin $\alpha 2\beta 1$ was analyzed immunohistochemically and correlated with the survival data and other therapy-relevant biomarkers. The significant correlation of a high $\alpha 2\beta$ 1-expression with the estrogen receptor alpha (ER α ; p = 0.035) and epithelial growth factor receptor (EGFR; p = 0.027) was observed. In addition, high $\alpha 2\beta$ 1-expression was significantly associated with a low number of tumor-infiltrating immune cells (CD3 intratumoral, p = 0.017; CD3 stromal, p = 0.035; PD-1 intratumoral, p = 0.002; PD-1 stromal, p = 0.049) and the lack of PD-L1 expression (p = 0.005). In Kaplan–Meier survival analysis, patients with a high expression of integrin $\alpha 2\beta 1$ revealed a significant shorter progression-free survival (PFS, p = 0.035) and platinum-free interval (PFI, p = 0.034). In the multivariate Cox regression analysis, integrin $\alpha 2\beta 1$ was confirmed as an independent prognostic factor for both PFS (p = 0.021) and PFI (p = 0.020). Dual expression of integrin $\alpha 2\beta 1$ and the hepatocyte growth factor receptor (HGFR; PFS/PFI, p = 0.004) and CD44v6 (PFS, p = 0.000; PFI, p = 0.001; overall survival [OS], p = 0.025) impaired survival. Integrin $\alpha 2\beta 1$ was established as a prognostic and predictive marker in primary ovarian cancer with the potential to stratify patients for chemotherapy and immunotherapy, and to design new targeted treatment strategies.

Keywords: primary ovarian cancer; integrin $\alpha 2\beta 1$; prognostic factor; predictive factor; immune infiltrate; targeted therapy; personalized medicine

1. Introduction

Several clinicopathological factors, such as advanced tumor stage and residual tumor after surgery, have been established as strong prognostic factors in primary ovarian



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Copyright: © 2021 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). cancer [1]. In addition, a few tumor biological characteristics have been identified as prognostic markers. Examples are distinct gene signatures [2] or a high number of T-cells [3]. Although recently new promising candidates were detected [4], predictive markers are rare in primary ovarian cancer. Two targeted therapy approaches are recommended under current guidelines, namely vascular endothelial growth factor (VEGF) inhibition and poly (ADP-ribose) polymerase (PARP) inhibition, which are both administered in addition to the standard chemotherapy [5–7]. For VEGF inhibition, no predictive biomarker is available to select appropriate patients for anti-angiogenic therapy. Similarly, BRCA mutation or HRD status, which so far represent a prerequisite for some of the PARP inhibition treatments, need to be re-evaluated [8]. Thus, robust biomarkers for precise prognosis and treatment response are urgently required in primary ovarian cancer. This importance is emphasized by the fact that, despite standard therapy combining radical surgery and adjuvant platinum-based chemotherapy, 70–80% of the patients suffer from relapse [9].

Integrins are transmembrane cell adhesion molecules, which mediate cell–cell and cell–extracellular matrix (ECM) interaction. Currently, 18 α -subunits and 8 β -subunits are identified, forming a variety of integrin heterodimers [10]. Due to their ability of inside-out and outside-in signaling, they are known to be involved in migration, invasion, and metastasis promoting tumor progression in several cancer types [11–13]. Considering the mechanism of ovarian cancer metastasis by spreading in the peritoneal fluid and attaching to the omental and peritoneal tissue [14–16], integrins seem to be a promising therapeutic target in ovarian cancer.

While there is already some information about ovarian cancer and other β 1heterodimers, such as integrins $\alpha 4\beta1$ and $\alpha 5\beta1$ [17], less information is available about integrin $\alpha 2\beta1$. The main ligand of integrin $\alpha 2\beta1$ is collagen type I, but binding to other collagen types, laminins, and other ECM-proteins is also possible [18,19]. Expression of integrin $\alpha 2\beta1$ is not only observed on the epithelial cells, but also on the endothelial cells, platelets, white blood cells, and fibroblasts [20,21].

Previous studies indicate a role of integrin $\alpha 2\beta 1$ in chemotherapy resistance [22,23], which constitutes a special interest for ovarian cancer. In the present study, the expression of integrin $\alpha 2\beta 1$ in primary ovarian cancer and its prognostic and predictive role will be evaluated.

2. Materials and Methods

2.1. Study Population

Forty-eight patients diagnosed with a primary, chemonaive ovarian, fallopian tube, or peritoneal cancer from the SpheroID-Study were included. Patients suffering from another neoplasia within the last five years were excluded. Patients were recruited between September 2012 and January 2015 from five ovarian cancer centers, namely University Hospital, LMU Munich (n = 16), Klinikum Dritter Orden (n = 15), Klinikum rechts der Isar, Technical University Munich (n = 7), Munich clinic Harlaching (n = 5), and Starnberg Hospital (n = 5). Standardized surgical resection and pathological analysis was conducted by the recruiting hospital. Patient-, tumor- and treatment-related data for correlations were given in the routine reports and delivered in a pseudonymized form. Survival analysis was performed after the completion of chemotherapy. Seven patients with no chemotherapy or a reduced number of treatment cycles (≤ 2) had to be excluded. Progression-free survival (PFS) was defined as the time from surgical treatment to relapse or progression. Platinumfree interval (PFI) was defined as the time from end of the chemotherapy to relapse or progression. Overall survival (OS) was defined as the time from surgical treatment to death. Data from patients who did not die and had no relapse or progression were censored at the date of their last visit.

2.2. Immunohistochemistry

After surgical removal, tumor samples were snap frozen in liquid nitrogen. Serial cryosections (5 μ m) were performed. The samples were stained immunohistochemically

using the avidin–biotin–peroxidase method [24]. Tissue sections were fixed either in acetone for 8 min or, for the antigens ER α and PgR, in formalin for 3 min and afterwards in a citrate buffer for 7 min at 90 °C. Blocking of unspecific Fc receptors was performed with 10% AB Serum (Biotest, Dreieich, Germany) in either PBS (acetone fixation) or in a TRIS–HCl buffer (formalin fixation) for 20 min. Endogenous biotin was blocked with a two-step avidin–biotin blocking kit (Vector Laboratories, Burlingame, CA, USA) according to the manufacturer's instructions for 20 min. Primary antibodies were applied for one hour. Details about primary and secondary antibodies and working concentrations, including the appropriate positive and negative controls, are given in Table 1. Secondary biotinylated antibodies and peroxidase conjugated streptavidin (Dianova, Hamburg, Germany) were incubated for 30 min each.

Antigen	Clone	Species	Fixation	Use of Kit	wc (µg/mL)	Supplier	Cut-Off for Positivity
Primary antib	odies						
Integrin α2β1	BHA2.1	m	Acetone	-	2.50	Millipore, Burlington, MA, USA	≥20%
ERα	1D5	m	Formalin	+	2.50	Dako, Santa Clara, CA, USA	$\geq 1\%$
PR	PgR 636	m	Formalin	+	2.50	Dako, Santa Clara, CA, USA	$\geq 1\%$
HER-2/neu	4B5	r	Acetone	-	1.50	Ventana, Roche, Basel, CH	$\geq 10\%$ (Intensity
EGFR	H11	m	Acetone	-	2.94	Dako, Santa Clara, CA, USA	>50%
HGFR	SP44	r	Acetone	-	2.12	Spring Bioscience, Pleasanton, CA, USA	≥50%
IGF1R	23-41	m	Acetone	+	4.00	invitrogen, Carlsbad, CA, USA	$\geq 80\%$
MUC-1	Ma552	m	Acetone	-	0.50	Monosan, Uden, NL	\geq 70%
CD44v6	VFF-18	m	Acetone	-	1.00	affymetrix eBioscience, Santa Clara, CA, USA	≥10%
Integrin αVβ3	LM609	m	Acetone	-	5.00	Millipore, Burlington, MA, USA	\geq 20%
CD3	UCHT1	m	Acetone	-	1.25	BD Biosciences, Franklin Lakes, NJ, USA	
CD8	C8/144B	m	Acetone	+	3.00	Dako, Santa Clara, CA, USA	
PD-1	MIH4	m	Acetone	+	10.00	affymetrix eBioscience, Santa Clara, CA, USA	
PD-L1	MIH1	m	Acetone	+	10.00	affymetrix eBioscience, Santa Clara, CA, USA	$\geq 1\%$
Positive contro	ols						
Epithelial Antigen	Ber-EP4	m	Acetone	-	2.50	Dako, Santa Clara, CA, USA	
CD45	2B11 + PD7/26	m	Acetone	-	4.50	Dako, Santa Clara, CA, USA	
Isotype contro	ls						
MOPC 21	MOPC 21	m		-	5.00	Sigma-Aldrich, St. Louis, MO. USA	
	MOPC 21	m		+	4.00	Sigma-Aldrich, St. Louis, MO, USA	
	MOPC 21	m		+	10.00	Sigma-Aldrich, St. Louis, MO, USA	
DA1E	DA1E	r		-	2.12	Cell Signaling, Danvers, MA, USA	
Biotin conjuga	ted secondary	antibodies					
	111-065- 114	g anti r			7.00	Jackson Immunoresearch, West Grove, PA, USA	
	315-065- 048	r anti m			0.75	Jackson Immunoresearch, West Grove, PA, USA	

Table 1. Biomarkers and antibodies.

Legend: wc: working concentration, m: mouse, r: rabbit, g: goat, all used antibodies' isotype was IgG1. ERα: estrogen receptor α, PR: progesterone receptor, Her-2/neu: human epidermal growth factor receptor 2, EGFR: epidermal growth factor receptor, HGFR: hepatocyte growth factor receptor, IGF1R: insulin-like growth factor 1, MUC-1: mucin-1, PD-1: programmed cell death protein 1, PD-L1: programmed cell death protein 1, PD-L1: programmed 1.

2.3. Evaluation of Biomarker Expression

Sections were evaluated semiquantitatively using a light microscope (Figure S1). The percentage of positively stained carcinoma cells was evaluated for each antigen. Tumors were defined as hormone receptor-positive if $\geq 1\%$ of the cancer cells revealed a nuclear staining of ER or PR [25]. Her2/neu expression was scored according to breast cancer [26] and gastric cancer [27] guidelines. Due to the lack of further references, the other biomarkers' expression was estimated as a percentage of positive cancer cells in 10% steps. Validation was conducted by a second observer (FS). In the absence of standardized cut-offs for other biomarkers, cut-offs were evaluated according to the biphasic distribution or the group size (see Table 1). Quantitative evaluation of CD3, CD8, and PD-1, and semiquantitative evaluation of PD-L1 was performed according to Dotzer et al. [24].

2.4. Statistical Analysis

Clinicopathological factors were grouped by clinical relevance. Integrin expression was correlated with clinicopathological factors, other biomarkers' expression, and immune infiltrate using the Fisher's exact two-tailed test. Univariate analysis was performed by calculating cumulative survival probabilities with the Kaplan–Meier method and comparing them with a log-rank test. A Cox regression model was used for the multivariate analysis of survival. *p*-values < 0.05 were considered to be statistically significant. All statistical analyses were performed using IBM SPSS Statistics 23 (Armonk, NY, USA).

3. Results

3.1. Patient Characteristic

The clinicopathological data are shown in Table 2. Forty-eight patients were included in this study. The mean age at time of diagnosis was 62 years. Most patients suffered from high-grade, serous ovarian carcinoma in an advanced FIGO (Fédération Internationale de Gynécologie et d'Obstétrique) stage with the presence of ascites. Complete surgical resection without macroscopic residual tumor was achieved in 72.9% of all patients. In total, 83.4% of the patients received chemotherapy based on carboplatin and paclitaxel. The median OS was 42 months, the median PFS was 22 months, and the median PFI was 17 months.

		n or Value	%
Age	mean/median range	62/66 years 24–83 years	
FIGO Stage	I or II	0	0.0%
	III	34	70.8%
	IV	14	29.2%
рТ	pT2	5	10.4%
	pT3	43	89.6%
pN	pN0	6	12.5%
	pN1	31	64.6%
	Nx	11	22.9%
cM	cM0	34	70.8%
	cM1	14	29.2%
Primary Tumor Site	Ovarian	39	81.3%
	Fallopian Tube	6	12.5%
	Peritoneal	3	6.3%
Histological Subtype	Serous	44	91.7%
	Other	4	8.4%
Grading	G1/G2	2	4.2%
	G3	46	95.8%

Table 2. Patient characteristics.

		n or Value	%
	yes	40	83.3%
Ascites	no	8	16.7%
Magragania Pasidual Tumar	None	35	72.9%
wiacioscopic Residual fumor	<1 cm	6	12.5%
after Surgery	>1 cm	7	14.6%
	С	4	8.3%
	C + P	15	31.3%
First-Line-Treatment	C + P + B	25	52.1%
	None	4	8.3%
	<6 months	2	4.2%
Release offer Chemotherenzy	6–12 months	12	25.0%
Relapse after Chemotherapy	>12 months	28	58.3%
	none or non-sufficient chemotherapy	6	12.5%

Table 2. Cont.

Legend: n: number of patients, Nx: no evaluation of lymph node status, C: carboplatin, P: paclitaxel, B: bevacizumab.

Survival data are summarized in Table 2. The presence of distant metastases (FIGO IV) was related to a shorter OS (p = 0.015) and tended to predict a shorter PFS (p = 0.081) and PFI (p = 0.068). Furthermore, patients with a macroscopic residual tumor after surgery showed a significant shorter OS (p = 0.041), PFS (p = 0.008) and PFI (p = 0.01).

3.2. Prognostic and Predictive Impact of Integrin α2β1

High integrin $\alpha 2\beta 1$ expression in primary ovarian cancer was found to be associated with an unfavorable prognosis. Patients with a high expression of integrin $\alpha 2\beta 1$ showed a median PFS of 16 months, which was significantly shorter compared to patients with low $\alpha 2\beta 1$ expression (PFS 29 months, p = 0.035). In addition, high expression of integrin $\alpha 2\beta 1$ predicted a shorter PFI (11 months) in contrast to patients with a low $\alpha 2\beta 1$ -expressing primary tumor (25 months, p = 0.034). Most importantly, a high expression of integrin $\alpha 2\beta 1$ in primary ovarian cancer was found to be an independent prognostic factor for a shorter PFS (HR 2.46, CI 95% 1.14–5.29, p = 0.021) and a shorter PFI (HR 2.44, CI 95% 1.14–5.26, p = 0.022). No impact of the extent of $\alpha 2\beta 1$ expression of integrin $\alpha 2\beta 1$ and clinicopathological factors could be found.

Table 3. Univariate and multivariate survival analysis of clinicopathological factors and integrin $\alpha 2\beta 1$.

				PFS				PFI		(os
	n	Log	-Rank	MV Cox Regre	ession	Log	-Rank	MV Cox Regre	ession	Log	-Rank
		MS	р	HR (CI 95%)	р	MS	p	HR (CI 95%)	р	MS	р
Age \leq 62 years Age > 62 years	19 23	22 22	0.965			17 17	0.970			nr 42	0.193
<pt3c pT3c</pt3c 	7 35	27 22	0.665			22 17	0.679			45 42	0.928
pN0 pN1	5 28	29 22	0.163			17 22	0.145			45 42	0.929
cM0 cM1	29 13	27 16	0.081	2.06 (0.92-4.62)	0.081	22 11	0.068	2.10 (0.94-4.69)	0.072	nr 30	0.015
G1/G2 G3	2 40	14 22	0.579			8 17	0.610			30 42	0.843
Ascites absent Ascites present	6 36	35 19	0.147			30 15	0.139			42 38	0.408

				PFS				PFI		(OS
	n	Log	-Rank	MV Cox Regression		Log-Rank		MV Cox Regression		Log	Rank
		MS	p	HR (CI 95%)	р	MS	р	HR (CI 95%)	р	MS	р
MR Tumor absent	30	27				22				45	
MR Tumor present	12	13	0.008	2.19 (1.03–4.68)	0.043	9	0.010	2.10 (0.99-4.51)	0.057	26	0.041
Integrin $\alpha 2\beta 1$ low	27	29				25				45	
Integrin α2β1 high	15	16	0.035	2.46 (1.14–5.29)	0.021	11	0.034	2.45 (1.14-5.26)	0.022	30	0.155

Table 3. Cont.

Legend: n: number of patients, Cox regression: multivariate Cox regression, MS: median survival (in months) in Kaplan–Meier estimator, HR: hazard ratio, CI: confidence interval, MR Tumor: macroscopic residual tumor; nr: median survival not reached.

3.3. Correlation of Integrin $\alpha 2\beta 1$ with Other Biomarkers

In almost all patients (17 out of 18, 94.4%), a high expression of integrin $\alpha 2\beta 1$ significantly correlated with a high expression of ER α (p = 0.035). Furthermore, a high expression of integrin $\alpha 2\beta 1$ could be found more frequently in patients with a high expression of EGFR (7 out of 10, 70%) compared to patients with a low expression of EGFR (11 out of 38, 28.9%, p = 0.027, Table 4).

Table 4. Correlation between	n integrin $\alpha 2$	β1 and other	biomarkers.
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					Integrin α2β1	
			n	<20%	≥ 20%	p #
			48			0.035
	ERα	<1%		10	1	
		$\geq 1\%$		20	17	
			48			0.127
	PR	<1%		22	9	
		$\geq 1\%$		8	9	
			48			1
	Her-2/neu	negative		22	13	
Growth Factor-Receptor		positive		8	5	
			48			0.027
	EGFR	<50%		27	11	
		\geq 50%		3	7	
			48			0.133
	HGFR	<50%		16	5	
		\geq 50%		14	13	
			48			0.451
	IGF1R	<80%		4	4	
		\geq 80%		26	14	
			48			0.765
	MUC-1	<70%		14	7	
		\geq 70%		16	11	
Cell-Adhesion-			48			0.103
Molecule	CD44v6	<10%		24	10	
		≥10%		6	8	
			48			0.19
	Integrin αvβ3	<20%		24	11	
		\geq 20%		6	7	

Legend: n: number of patients, [#]: *p*-value calculated by Fisher's exact two-tailed test.

3.4. Prognostic and Predictive Impact of Integrin &2B1 Combined with Other Biomarkers

The dual expression of integrin $\alpha 2\beta 1$ and various growth factor receptors revealed an impact on PFS and PFI (Table 5). Patients with a high expression of integrin $\alpha 2\beta 1$ and a positive Her-2/neu status showed a shorter PFS (p = 0.043) and PFI (p = 0.037) than patients with a low expression of integrin $\alpha 2\beta 1$, Her-2/neu, or both. Combined high expression of integrin $\alpha 2\beta 1$ and IGF1R correlated significantly with a shorter PFS (p = 0.045) and PFI (p = 0.043). Most interestingly, a high expression of integrin $\alpha 2\beta 1$ and HGFR was related to a shorter PFS (p = 0.004) and PFI (p = 0.004) and impaired prognosis in comparison to integrin $\alpha 2\beta 1$ as single biomarker.

		P	FS	Р	FI	(OS
	n	MS	p *	MS	p *	MS	<i>p</i> *
Integrin $\alpha 2\beta 1$ high	15	16 20	0.035	11	0.034	30 45	0.155
Integrin a2p1 low	27	29		25		45	
Integrin $\alpha 2\beta 1$ high/ER α high	14	16	0.078	11	0.073	30	0.287
Remaining combinations [#]	28	27	0.078	22	0.075	42	0.207
Integrin $\alpha 2\beta 1$ high/PR high	8	16	0 574	1119	0 579	27	0 526
Remaining combinations #	34	24	0.374	19	0.578	42	0.526
Integrin $\alpha 2\beta 1$ high/Her-2/neu +	5	21	0.042	15	0.007	36	0.600
Remaining combinations #	37	27	0.043	22	0.037	42	0.698
Integrin $\alpha 2\beta 1$ high/EGFR high	6	14	0.000	8	0.000	30	0.400
Remaining combinations #	36	22	0.289	17	0.290	42	0.482
Integrin $\alpha 2\beta 1$ high/HGFR high	11	15	0.004	10	0.004	27	0.054
Remaining combinations #	31	29	0.004	25	0.004	45	0.054
Integrin $\alpha 2\beta 1$ high/IGFR high	11	16	0.045	11	0.042	36	0.291
Remaining combinations [#]	31	27	0.045	22	0.043	42	0.381
Integrin $\alpha 2\beta 1$ high/MUC-1 high	9	14	0.0(2	9		27	0.057
Remaining combinations #	33	27	0.063	22	0.055	42	0.257
Integrin $\alpha 2\beta 1$ high/CD44v6 high	6	13	0.000	9	0.001	19	0.005
Remaining combinations #	36	27	0.000	22	0.001	42	0.025
Integrin $\alpha 2\beta 1$ high/Integrin $\alpha v\beta 3$	5	35	0.222	30	0.220	nr	0.1(2
Remaining combinations [#]	37	22	0.322	17	0.320	42	0.162

Table 5. Univariate survival analysis of dual expression of integrin $\alpha 2\beta 1$ and other biomarkers.

Legend: n: number of patients, MS: median survival (in months) in Kaplan–Meier estimator, *: *p*-value calculated by log-rank test. [#] The remaining combinations represent tumor samples which were integrin $\alpha 2\beta 1$ high/biomarker X low, integrin $\alpha 2\beta 1$ low/biomarker X high, or integrin $\alpha 2\beta 1$ low/biomarker X low. nr: median survival not reached.

Likewise, a high expression of both integrin $\alpha 2\beta 1$ and CD44v6 was found to be a strong factor in a poor prognosis that correlated with a shorter PFS (p = 0.000), PFI (p = 0.001) and a reduced OS (p = 0.025, Table 5).

3.5. Correlation of Integrin $\alpha 2\beta 1$ and Immune Infiltrate

In patients with a high expression of integrin $\alpha 2\beta 1$, low numbers of stromal and intratumoral CD3+ cells were found (14 out of 18, 77.8%, p = 0.035 and p = 0.017, Table 6). Furthermore, most tumors with a high expression of integrin $\alpha 2\beta 1$ showed a low density of stromal (16 out of 18, 88.9%, p = 0.049) and intratumoral (17 out of 18, 94.4%, p = 0.002) PD-1+ cells. PD-L1 positivity was found more often in tumors with a low expression of integrin $\alpha 2\beta 1$ (23 out of 30, 76.7%) compared to samples with a high expression (6 out of 18, 33.3%; p = 0.005). No correlations for CD8+ cells have been found.

Immune Infiltrate				Integrin α2β	1
minune minuate	-	n	<20%	≥ 20%	p #
		48			0.034
CD3 stromal	Low		13	14	
	High		17	4	
		48			0.017
CD3 intratumoral	Low		12	14	
	High		18	4	
		48			0.133
CD8 stromal	Low		14	13	
	High		16	5	
		48			0.363
CD8 intratumoral	Low		17	13	
	High		13	5	
		48			0.049
PD-1 stromal	Low		18	16	
	High		12	2	
		48			0.002
PD-1 intratumoral	Low		15	17	
	High		15	1	
		48			0.005
PD-L1 positivity	No		7	12	
- •	Yes		23	6	

Table 6. Correlations between integrin $\alpha 2\beta 1$ and the immune infiltrate.

Legend: n: number of patients, [#]: *p*-value as calculated by Fisher's exact two-tailed test.

4. Discussion

In the present study, integrin $\alpha 2\beta 1$ was identified as a potential new prognostic and predictive marker in primary ovarian cancer.

A high expression of integrin $\alpha 2\beta 1$ was identified as a marker for a poor prognosis with equal strength, as reported for the established clinical factors: FIGO stage and macroscopic residual tumor after surgical resection. The positive correlation between a high expression of the integrin $\beta 1$ chain and short survival is documented for various tumor entities [28–30]. In particular, integrin $\alpha 5\beta 1$ is already known to be an unfavorable prognostic factor for ovarian cancer [31], but also for cervical, gastric, and non-small-cell lung cancer [32–34].

Integrin $\alpha 2\beta 1$ is involved in many steps of cancer progression. Binding to components of the extracellular matrix (ECM), integrin $\alpha 2\beta 1$ mediates tumor cell invasion and metastasis [35–37]. This step is promoted by crosstalk with growth factor receptors [38,39]. Interestingly, in the present study, a combined expression of integrin $\alpha 2\beta 1$ with ER α and EGFR was observed. Furthermore, the signaling of integrin $\alpha 2\beta 1$ can induce chemoresistance. This mechanism was observed for chemotherapies containing paclitaxel [23,40], gemcitabine [41], and etoposide [42].

Early relapse and resistance to platinum-based chemotherapy are key problems in the treatment of ovarian cancer [43]. Therefore, the predictive value for the treatment response of integrin $\alpha 2\beta 1$ was analyzed in the present study. Patients with a high expression of integrin $\alpha 2\beta 1$ were observed to have a shorter median PFI. In particular, $\beta 1$ integrins are already known to promote platinum resistance in ovarian cancer. The mechanisms of this effect are still unclear. Intracellular signaling initiated by binding to the ECM seems to be fundamental for cell adhesion-mediated drug resistance (CAM-DR) [44,45]. One of the main ECM molecules involved in this concept is collagen type I [46], which is the central binding partner of integrin $\alpha 2\beta 1$ [18]. These molecular interactions suggest that the

heterodimer $\alpha 2\beta 1$ contributes to CAM-DR. Therefore, targeting integrin $\alpha 2\beta 1$ represents a promising strategy for overcoming platinum resistance in primary ovarian cancer.

In addition, a high expression of integrin $\alpha 2\beta 1$ was observed in patients with a low density of stromal and intratumoral CD3+ as well as PD-1+ cells. Inversely, more than 75% of patients with a low expression of integrin $\alpha 2\beta 1$ showed PD-L1 positivity, which represents an established predictive biomarker for immunotherapy [47]. Several integrins are related to an immunosuppressive tumor microenvironment [48,49]. For example, αv -integrins are major activators of latent TGF- β , which is involved in immunotherapy resistance [50]. The present data suggest that integrin $\alpha 2\beta 1$ might play a similar role. Recently, immunotherapy became a promising approach in ovarian cancer [51,52], and phase III studies with checkpoint inhibitors in combination with platinum-based chemotherapy are already ongoing (NCT03038100, NCT03740165, NCT03737643). Low expression of integrin $\alpha 2\beta 1$, therefore, could be a potential predictive marker for immunotherapy in ovarian cancer. Taken together, integrin $\alpha 2\beta 1$ represents a stratification marker for patients receiving platinum-based chemotherapy and immunotherapy.

Inhibition of integrin $\alpha 2\beta 1$ should be considered as a targeted therapy in ovarian cancer. Several molecules and antibodies have been developed and evaluated for integrin $\alpha 2\beta 1$ inhibition in other entities.

Anti-tumoral activity was shown in prostate cancer in vivo using the monoclonal antibody GBR-500 [53]. E-7820 is a sulphonamide derivative that inhibits the expression of α 2-mRNA. In Phase I studies, treatment was associated with a stable disease in a variety of malignancies [54,55]. Phase II studies are ongoing to evaluate the combination with chemotherapy in colon carcinoma (NCT01347645, NCT01133990, NCT00309179). Another β 1-antibody could improve the efficiency of platinum-based chemotherapy in non-small-cell lung cancer [56]. However, despite these promising approaches, the complex biology of heterodimers with promiscuous ligands, allosteric activation, and multiple intracellular signaling pathways might hinder successful treatment strategies [13,57,58].

Furthermore, the results of this study also indicate the potential efficiency of dual inhibition. Patients with a combined high expression of integrin $\alpha 2\beta 1$ and HGFR or CD44v6 showed a very short median PFS and PFI, indicating a worse prognosis and platinum resistance.

Dual targeting has become a promising strategy in ovarian cancer. Its efficiency was proven in tumor spheroid and mouse models [59,60]; thus, various phase I studies are ongoing (NCT03895788, NCT03695380, NCT04315233). In future studies, dual inhibition including integrin $\alpha 2\beta$ 1-antagonists should be considered for patients with an appropriate biomarker profile.

The main limitation of this study is the small cohort, though it is representative and comparable to cohorts of other clinical trials. The promising role of integrin $\alpha 2\beta 1$ as a new prognostic and predictive biomarker in primary ovarian cancer needs to be confirmed by an enlarged study.

5. Conclusions

In the present study, integrin $\alpha 2\beta 1$ was identified as a prognostic and predictive marker in primary ovarian cancer. High expression of integrin $\alpha 2\beta 1$ correlated with a short PFS. Prognosis was even worse in integrin $\alpha 2\beta 1$ -positive tumors co-expressing HGFR or CD44v6. This finding might lead to new biomarker-directed treatment strategies in primary ovarian cancer. In addition, the high expression of integrin $\alpha 2\beta 1$ correlated with a short PFI, supporting the hypothesis that integrins mediate platinum resistance. Thus, a high expression of integrin $\alpha 2\beta 1$ might represent a stratification marker for personalized treatment.

Supplementary Materials: The following are available online at https://www.mdpi.com/2227-905 9/9/3/289/s1, Figure S1: Immunohistochemical stainings.

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