Drug-Resistant Urothelial Cancer Cell Lines Display Diverse Sensitivity Profiles to Potential Second-Line Therapeutics[1,2]

Stefan Vallo[*,†], Martin Michaelis[‡], Florian Rothweiler[*], Georg Bartsch[†], Kilian M. Gust[†], Dominik M. Limbart[*], Franz Rödel[§], Felix Wezel[¶], Axel Haferkamp[†] and Jindrich Cinatl Jr.[*]

[*]Institute of Medical Virology, University Hospital Frankfurt, Frankfurt am Main, Germany; [†]Department of Urology, University Hospital Frankfurt, Frankfurt am Main, Germany; [‡]Centre for Molecular Processing and School of Biosciences, University of Kent, Canterbury, United Kingdom; [§]Department of Radiation Therapy and Oncology, University Hospital Frankfurt, Frankfurt am Main, Germany; [¶]Department of Urology, University Hospital Mannheim, Mannheim, Germany

Abstract

Combination chemotherapy with gemcitabine and cisplatin in patients with metastatic urothelial cancer of the bladder frequently results in the development of acquired drug resistance. Availability of cell culture models with acquired resistance could help to identify candidate treatments for an efficient second-line therapy. Six cisplatin- and six gemcitabine-resistant cell lines were established. Cell viability assays were performed to evaluate the sensitivity to 16 different chemotherapeutic substances. The activity of the drug transporter ATP-binding cassette transporter, subfamily B, member 1 (ABCB1, a critical mediator of multidrug resistance in cancer) was evaluated using fluorescent ABCB1 substrates. For functional assessment, cells overexpressing ABCB1 were generated by transduction with a lentiviral vector encoding for ABCB1, while zosuquidar was used for selective inhibition. In this study, 8 of 12 gemcitabine- or cisplatin-resistant cell lines were cross-resistant to carboplatin, 5 to pemetrexed, 4 to methotrexate, 3 to oxaliplatin, 5-fluorouracil, and paclitaxel, and 2 to cabazitaxel, larotaxel, docetaxel, topotecan, doxorubicin, and mitomycin c, and 1 of 12 cell lines was cross-resistant to vinflunine and vinblastine. In one cell line with acquired resistance to gemcitabine (TCC-SUP[r]GEMCI[20]), cross-resistance seemed to be mediated by ABCB1 expression. Our model identified the vinca alkaloids vinblastine and vinflunine, in Europe an already approved second-line therapeutic for metastatic bladder cancer, as the most effective compounds in urothelial cancer cells with acquired resistance to gemcitabine or cisplatin. These results demonstrate that this *in vitro* model can reproduce clinically relevant results and may be suitable to identify novel substances for the treatment of metastatic bladder cancer.

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Introduction

Patients with metastatic urothelial cancer of the bladder are treated with cisplatin containing systemic chemotherapies (e.g., gemcitabine/cisplatin, GC) as a standard of care [1,2]. Unfortunately, the treatment success is limited resulting in a median survival of 12 to 14 months. Treatment failure is commonly caused by development of resistance to chemotherapy [1,2].

ATP-binding cassette transporter, subfamily B, member 1 (ABCB1) is a cell membrane efflux pump with broad substrate specificity. Overexpression of ABCB1 in tumor cells develops mostly as a specific response to ABCB1 substrates (e.g., vinca alkaloids, taxanes, or anthracyclines) and confers resistance to these substances. However, ABCB1 may also be upregulated as part of a generalized stress response to different toxic drugs (such as gemcitabine and cisplatin), which are not ABCB1 substrates [3,4]. Expression of ABCB1 was detected in both pre-chemotherapy and post-chemotherapy tumor tissue samples from patients with bladder cancer with higher expression in post-chemotherapy patients [5–11]. Therefore, efficient second-line chemotherapies or targeted therapies for the treatment of bladder cancer need to be tested especially in a context of specific resistance mechanisms such as ABCB1 overexpression.

Development of acquired cancer cell drug resistance is difficult to study in a clinical setting. Since acquisition of tumor biopsies represents an invasive procedure, possibilities to obtain serial tumor biopsies from patients under chemotherapy are limited by technical as well as ethical barriers [12]. Moreover, significance of biopsies may be affected by intratumor heterogeneity [13]. "Liquid biopsies" including circulating tumor cells and tumor DNA may be valuable sources for detection of molecular changes associated with resistance in the future [14] but may be unsuitable for functional studies. Therefore, experimental *in vitro* models are needed to identify potential markers of resistance and novel drug targets.

Drug-adapted cancer cell lines have been successfully used to study cancer cell mechanisms of resistance [15,16]; however, comprehensive cell line panels are missing. A panel of 18 urothelial cancer cell lines consisting of six parental chemosensitive cell lines and their gemcitabine- or cisplatin-resistant sublines was used to study the activity of 16 anticancer drugs. The cell lines are part of the Resistant Cancer Cell Line collection. This collection consists of cell lines of 15 different cancer entities including the six gemcitabine- and six cisplatin-resistant urothelial cancer cell lines that were used here.

Materials and Methods

Drugs

Cisplatin (solvent: 0.9% aqueous NaCl solution) was purchased from Gry-Pharma (Kirchzarten, Germany), gemcitabine (solvent: 0.9% aqueous NaCl solution) from Lilly (Bad Homburg, Germany), vinflunine [solvent: phosphate-buffered saline (PBS)] from Pierre Fabre (Freiburg, Germany), pemetrexed (solvent: DMSO) from Lilly, methotrexate (solvent: PBS) from Hexal (Holzkirchen, Germany), carboplatin (solvent: 5% aqueous glucose solution) from Hexal, oxaliplatin (solvent: PBS) from Teva (Basel, Switzerland), paclitaxel (solvent: DMSO) from Bristol-Myers Squibb (New York, NY), topotecan (solvent: dH$_2$O) from GlaxoSmithKline (London, United Kingdom), docetaxel (solvent: DMSO) from Sanofi (Paris, France), cabazitaxel (solvent: DMSO) from Sanofi, larotaxel (solvent: DMSO) from Shanghai Fuhe Chemistry Technology (Shanghai, China), vinblastine (solvent: PBS) from Teva, doxorubicin

(solvent: 0.9% aqueous NaCl solution) from Sigma-Aldrich (St Louis, MO), mitomycin c (solvent: dH$_2$O) from Medac (Wedel, Germany), and 5-fluorouracil (solvent: 0.9% aqueous NaCl solution) from Medac.

Cell Lines and Lentiviral Transduction

The cell lines RT112, RT4, 5637, T24, HT1376, and TCC-SUP were obtained from the American Type Culture Collection (Manassas, VA). Drug-resistant sublines were established by continuous exposure to increasing drug concentrations and are part of the Resistant Cancer Cell Line (RCCL) collection (http://www.kent.ac.uk/stms/cmp/RCCL/RCCLabout): RT112^{r}CDDP^{1000} (cisplatin-resistant, 1000 ng/ml cisplatin), RT112^{r}GEMCI^{20} (gemcitabine-resistant, 20 ng/ml gemcitabine), RT4^{r}CDDP^{1000}, RT4^{r}GEMCI^{10}, 5637^{r}CDDP^{1000}, 5637^{r}GEMCI^{20}, T24^{r}CDDP^{1000}, T24^{r}GEMCI^{20}, T24^{r}VBL^{20} (vinblastine-resistant, 20 ng/ml vinblastine), HT1376^{r}CDDP^{1000}, HT1376^{r}GEMCI^{20}, TCC-SUP^{r}CDDP^{1000}, TCC-SUP^{r}GEMCI^{20}, and TCC-SUP^{r}VBL^{20}.

Cell line adaptation was started with drug concentrations that were two-fold higher than the respective IC$_{50}$. The doses were stepwise increased during subculturing until resistance to clinically achievable plasma concentration was reached. The establishment of readily growing resistant cell lines required 1 to 2 years in dependence on the used cell line and the drug.

The ABCB1-expressing cell lines TCC-SUP^{ABCB1} and T24^{ABCB1} and the corresponding control cell lines TCC-SUP^{CER2} and T24^{CER2} were established by lentiviral transduction using the Lentiviral Gene Ontology vector technology as described previously [17,18].

All cell lines were grown in Iscove's modified Dulbecco's medium (IMDM) supplemented with 10% fetal calf serum (FCS; Gibco, Karlsruhe, Germany). Cell line authentication was performed by short tandem repeats (STR) profiling.

Growth Curves

To determine cell growth kinetics, 4000 cells per cm^2 were seeded in cell culture flasks containing IMDM supplemented with 10% FCS. Cell counts were determined using a Neubauer chamber in the presence of trypan blue. Doubling time (DT) was calculated using the formula DT = culture time/cell doubling. Cell doubling = ln(Nf/Ni)/ln2, where Ni represents seeded cell number and Nf represents the harvested cell number [19].

Cell Viability Assay

Cell viability was determined by the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) dye reduction assay after 120-hour incubation as described previously [20]. Drug resistance was defined by resistance factors defined as IC$_{50}$ drug-resistant cells/IC$_{50}$ parental cells. Cell lines were regarded to be resistant to a drug if the resistance factor was >2 [21].

Flow Cytometry

Antibodies directed against ABCB1 (20 µl per sample with an antibody concentration of 25 ng/µl; Alexis Biochemicals through AXXORA Deutschland, Lörrach, Germany) followed by secondary antibodies labeled with phycoerythrin were used to detect protein expression by flow cytometry (FACSCalibur; BD Biosciences, Heidelberg, Germany). Mouse IgG2a antibodies were used as isotype control.

For washout experiments, cells were incubated for 1 hour with 1 µM rhodamine 123 (R123, ABCB1 substrate). Zosuquidar (Sigma-Aldrich), an inhibitor of ABCB1, was added immediately. Cells were resuspended in supplemented growth medium, and cellular fluorescence was measured at FL1 channel by flow cytometry.

Statistical Analysis

Results are expressed as mean ± SD of at least three independent experiments. For statistical analysis, Student's t test, analysis of variance, and Student-Newman-Keuls test were performed whenever applicable. Significance was defined at values of $P \le .05$.

Results

Cell Growth Kinetics

Four of six gemcitabine-resistant sublines showed decreased growth rates compared to their parental cell lines [5637 (DT: 0.96 day) vs 5637^{r}GEMCI^{20} (DT: 1.74 day); HT1376 (DT: 1.29 day) vs HT1376^{r}GEMCI^{20} (DT: 1.74 day); TCC-SUP (DT: 2.05 day) vs TCC-SUP^{r}GEMCI^{20} (DT: 4.91 day); T24 (DT: 0.97 day) vs T24^{r}GEMCI^{20} (DT: 1.08 day)], while no significant differences were found for RT112 cells [RT112 (DT: 1.07 day) vs RT112-^{r}GEMCI^{20} (DT: 1.11 day)] and RT4 cells [RT4 (DT: 2.70 day) vs RT4^{r}GEMCI^{10} (DT: 2.34 day)]. Three of six cisplatin-resistant sublines [RT112^{r}CDDP^{1000} (DT: 0.96 day), RT4^{r}CDDP^{1000} (DT: 1.65 day), and TCC-SUP^{r}CDDP^{1000} (DT: 1.30 day)] displayed enhanced growth rates compared to their parental cell lines, while growth rate of 5637^{r}CDDP^{1000} (DT: 1.23 day) cells was decreased

relative to 5637 cells. For HT1376 vs HT1376^{r}CDDP^{1000} (DT: 1.30 day) and T24 vs T24^{r}CDDP^{1000} (DT: 1.03 day), no significant difference in growth rate was found (Figure 1).

Cross-Resistance Profiles

The effects of a panel of 16 anticancer drugs were determined on the viability of all 18 urothelial cancer cell lines by MTT assay. Cisplatin-resistant cell lines showed resistance factors to cisplatin (IC$_{50}$ resistant cell line/IC$_{50}$ parental cell line) ranging from 2.78 (HT1376^{r}CDDP^{1000}) to 28.86 (TCC-SUP^{r}CDDP^{1000}). Gemcitabine-resistant cell lines displayed resistance factors to gemcitabine ranging from 7.11 (RT4^{r}GEMCI^{10}) to 73.28 (RT112^{r}GEMCI^{20}) relative to parental cell lines (Suppl. Table 1).

The parental cell line panel included cell lines derived from low-risk carcinomas (RT4 and 5637) rarely requiring systemic chemotherapy in $vivo$ [22,23] and those from high-risk carcinomas (HT1376, RT112, T24, and TCC-SUP) that are commonly treated by chemotherapy when metastasized. Gemcitabine-resistant sublines of the low-risk carcinoma cell lines RT4 and 5637 showed resistance to three of the investigated 16 anticancer drugs (RT4^{r}GEMCI^{10}: gemcitabine, 5-fluorouracil, and carboplatin; 5637^{r}GEMCI^{20}: gemcitabine, methotrexate, and pemetrexed). Cisplatin-resistant sublines were resistant to three (RT4^{r}CDDP^{1000}: cisplatin, carboplatin, and 5-fluorouracil) and four (5637^{r}CDDP^{1000}: cisplatin, carboplatin, methotrexate, and topotecan) of the 16 drugs (Suppl. Table 1).

Gemcitabine-resistant sublines derived from high-risk urothelial carcinoma RT112, T24, HT1376, and TCC-SUP showed resistance
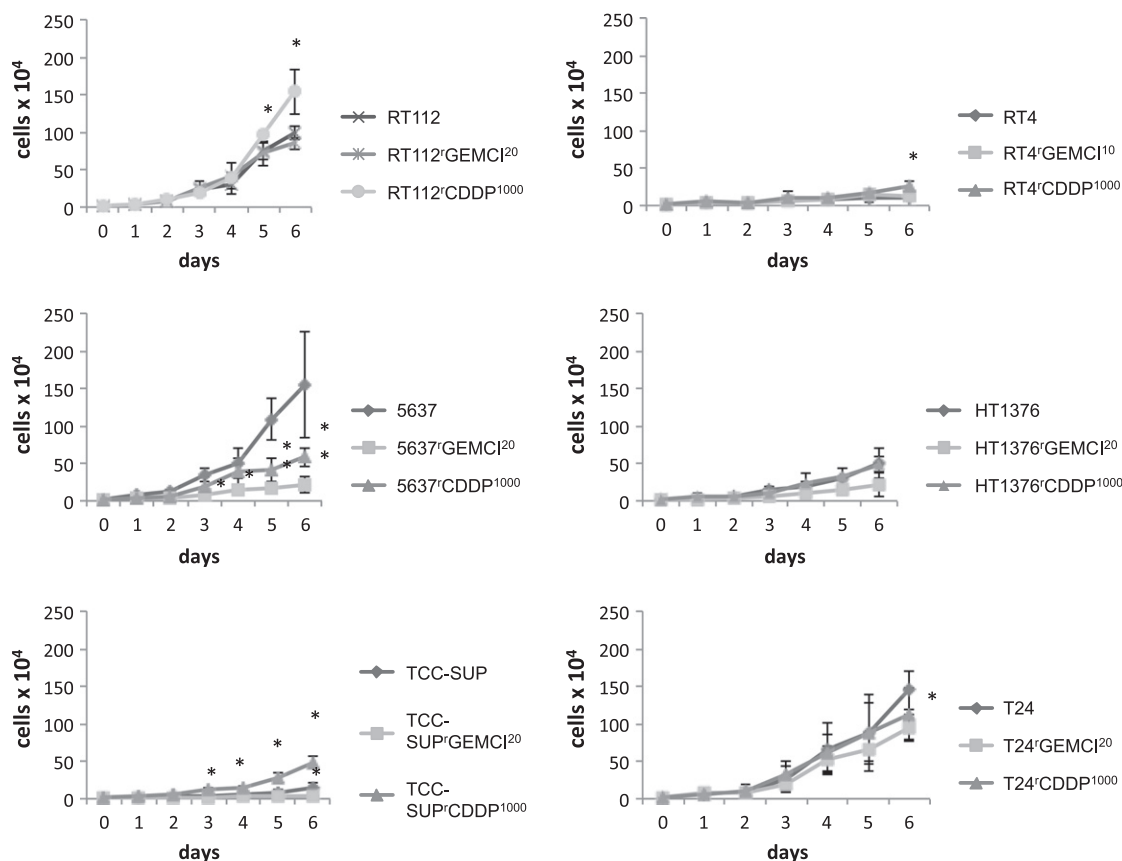

Figure 1. Growth curves of urothelial carcinoma cell lines; 4000 cells per cm^2 were seeded in cell culture flasks at day 0 containing IMDM supplemented with 10% FCS. Cell counts were determined using a Neubauer chamber in the presence of trypan blue. Values are displayed as mean ± SD. *$P \le .05$ relative to parental cell line.

to up to 11 of the tested substances of the 16 anticancer agents. Cisplatin-resistant sublines of RT112, T24, HT1376, and TCC-SUP were cross-resistant to up to five agents. In summary, most pronounced cross-resistance to other chemotherapeutic agents was observed in gemcitabine-resistant sublines of high-risk urothelial carcinoma cells (Suppl. Table 1).

Moreover, resistance profiles differed among the investigated drugs. Eight of 12 investigated resistant cell lines were cross-resistant to carboplatin, 5 of 12 to pemetrexed, 4 of 12 to methotrexate, 3 of 12 to oxaliplatin, 5-fluorouracil, and paclitaxel, and 2 of 12 to cabazitaxel, larotaxel, docetaxel, topotecan, doxorubicin, and mitomycin c. Only one resistant cell line (TCC-SUP^{r}GEMCI^{20}) was cross-resistant to vinflunine and vinblastine (Figure 2 and Suppl. Table 1).

ABCB1 Expression in Drug-Resistant Urothelial Carcinoma Cell Lines

Since ABCB1 overexpression is a major mechanism of resistance to chemotherapy, we evaluated its role in this model of urothelial bladder cancer cell lines with acquired drug resistance. TCC-SUP-^{r}GEMCI^{20} and T24^{r}GEMCI^{20} cells showed cross-resistance to ABCB1 substrates docetaxel, paclitaxel, and doxorubicin (Suppl. Table 1). The ABCB1 inhibitor zosuquidar sensitized TCC-SUP-^{r}GEMCI^{20} cells to vinflunine and vinblastine but not to gemcitabine

(Suppl. Table 2). Vinflunine was described to be a weaker ABCB1 substrate than other vinca alkaloids [24]. In accordance, the relative resistance IC$_{50}$ TCC-SUP^{r}GEMCI^{20}/IC$_{50}$ TCC-SUP and IC$_{50}$ T24^{r}GEMCI^{20}/IC$_{50}$ T24 was lower for vinflunine than for vinblastine (Suppl. Tables 1–3).

R123 is a fluorescent dye that is transported by ABCB1. A flow cytometric assay with R123 was used to determine functional activity of ABCB1 [25]. Flow cytometry indicated a strong increase of R123 fluorescence after treatment with zosuquidar in TCC-SUP-^{r}GEMCI^{20}, TCC-SUP^{r}VBL^{20}, TCC-SUP^{ABCB1}, T24^{r}GEMCI^{20}, T24^{r}VBL^{20}, and T24^{ABCB1} cells compared to cell lines that served as a control (Figures 3 and 4).

Next, we compared ABCB1 expression and drug sensitivity profiles in TCC-SUP and T24, TCC-SUP^{r}GEMCI^{20} and T24^{r}GEMCI^{20}, TCC-SUP^{ABCB1} and T24^{ABCB1} (TCC-SUP and T24 cells transduced with a lentiviral vector encoding for ABCB1), TCC-SUP^{CER2} and T24^{CER2} (TCC-SUP and T24 cells transduced with a control vector), and TCC-SUP^{r}VBL^{20} and T24^{r}VBL^{20} (TCC-SUP and T24 cells with acquired resistance to vinblastine, ABCB1 substrate). Successful transduction of the ABCB1 encoding plasmid in TCC-SUP^{ABCB1} and T24^{ABCB1} cells was verified by a significant increase of ABCB1 expression compared to cells transduced with a control vector (Figures 3 and 4). Compared to TCC-SUP^{ABCB1}, T24^{ABCB1}, TCC-SUP^{r}VBL^{20}, and T24^{r}VBL^{20}, gemcitabine-
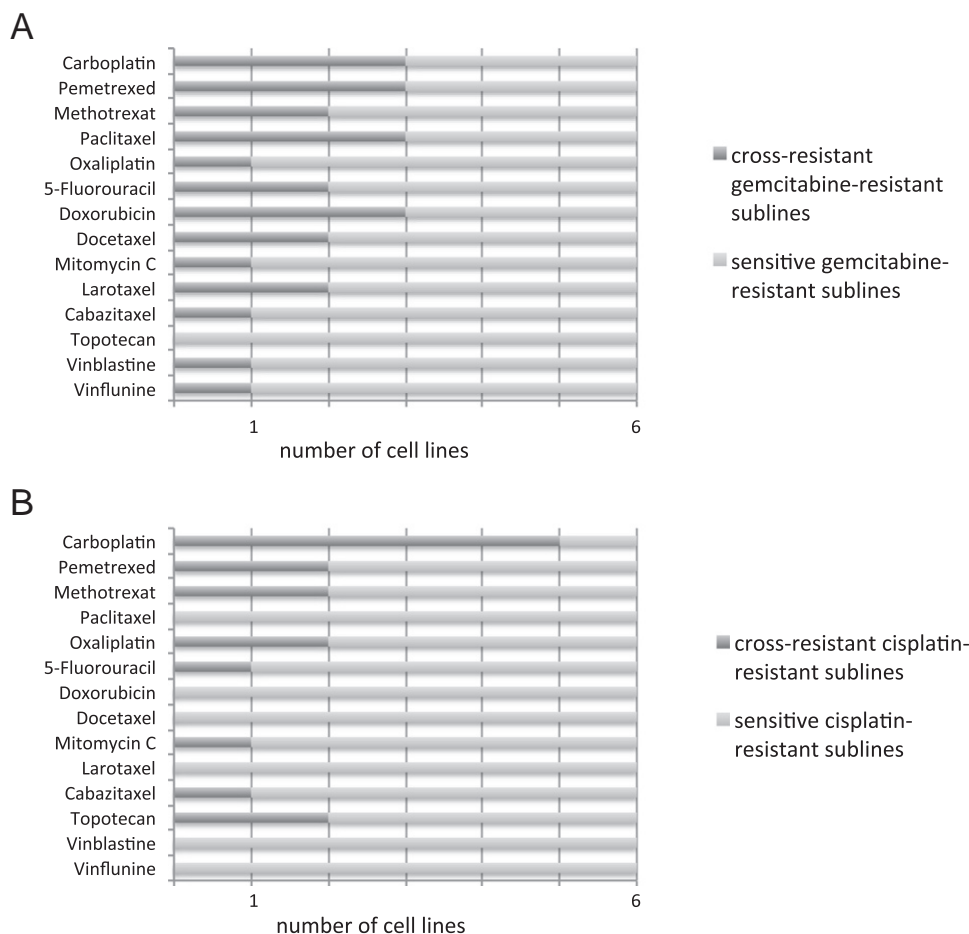

Figure 2. (A) The number of gemcitabine-resistant sublines that displayed cross-resistance to additional anticancer drugs is presented. Cross-resistance was defined as IC$_{50}$ (as determined by MTT assay) resistant subline/IC$_{50}$ respective parental cell line >2. (B) Cisplatin-resistant sublines that displayed cross-resistance to additional anticancer drugs.
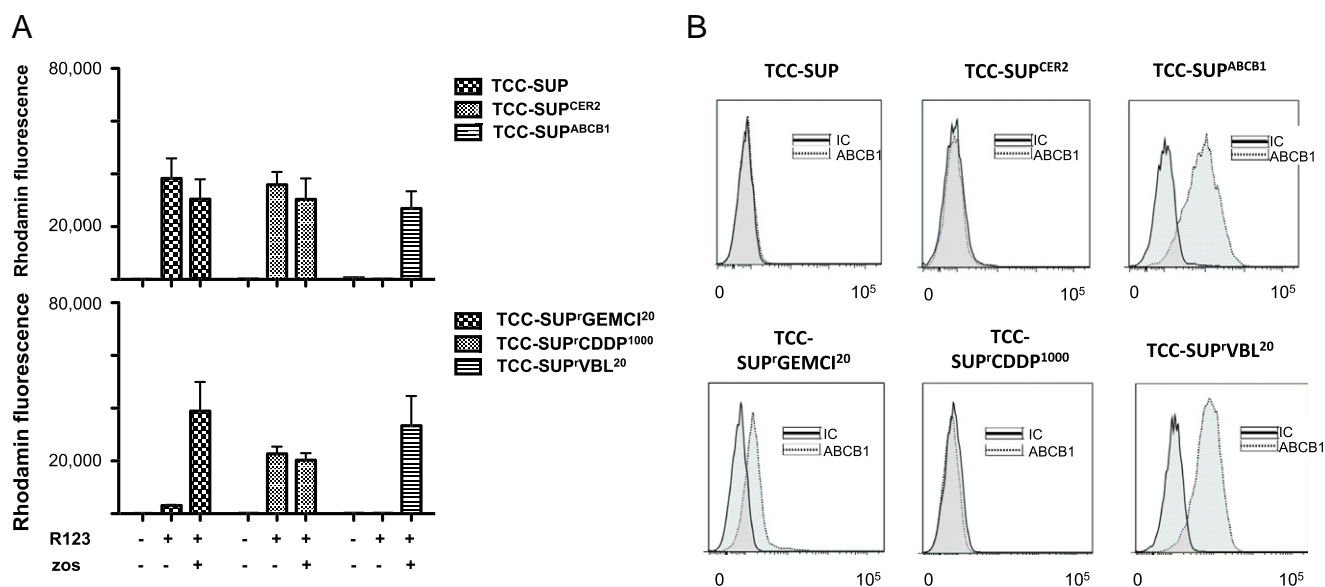
Figure 3. (A) R123 fluorescence of TCC-SUP, TCC-SUP^{CER2}, TCC-SUP^{ABCB1}, TCC-SUP^{r}GEMCI^{20}, TCC-SUP^{r}CDDP^{1000}, and TCC-SUP-^{r}VBL^{20} cells after staining with R123 alone and in combination with 1.25 μM zosuquidar. Values are means ± SD. (B) ABCB1 expression in TCC-SUP, TCC-SUP^{CER2}, TCC-SUP^{ABCB1}, TCC-SUP^{r}GEMCI^{20}, TCC-SUP^{r}CDDP^{1000}, and TCC-SUP^{r}VBL^{20} cells. IC, isotype control. *$P \leq$ .05 relative to parental cell line.

resistant sublines of TCC-SUP and T24 cells showed lower ABCB1 expression (Figures 3 and 4). Zosuquidar sensitized ABCB1-expressing cell lines to ABCB1 substrates with exemption of T24^{r}GEMCI^{20} cells (Suppl. Tables 2 and 3).

Discussion

In this study, we established a panel of urothelial cancer cell lines with acquired resistance to gemcitabine or cisplatin, the standard therapeutics for patients with metastasized urothelial cancer of the bladder [1].

First, we compared tumor cell growth differences, since rapidly dividing tumor cells might be more vulnerable to chemotherapy. There was no consistent correlation between cell growth kinetics and drug sensitivity in the investigated cell lines. This suggests that the cell growth kinetics are not critical for the drug response in our models.

The most effective compounds among our anticancer drug panel were vinflunine and vinblastine, since most gemcitabine- or cisplatin-resistant cell lines were still sensitive to these drugs. Notably, vinflunine was approved by the European Medicines Agency for second-line treatment of urothelial bladder cancer on the basis of a
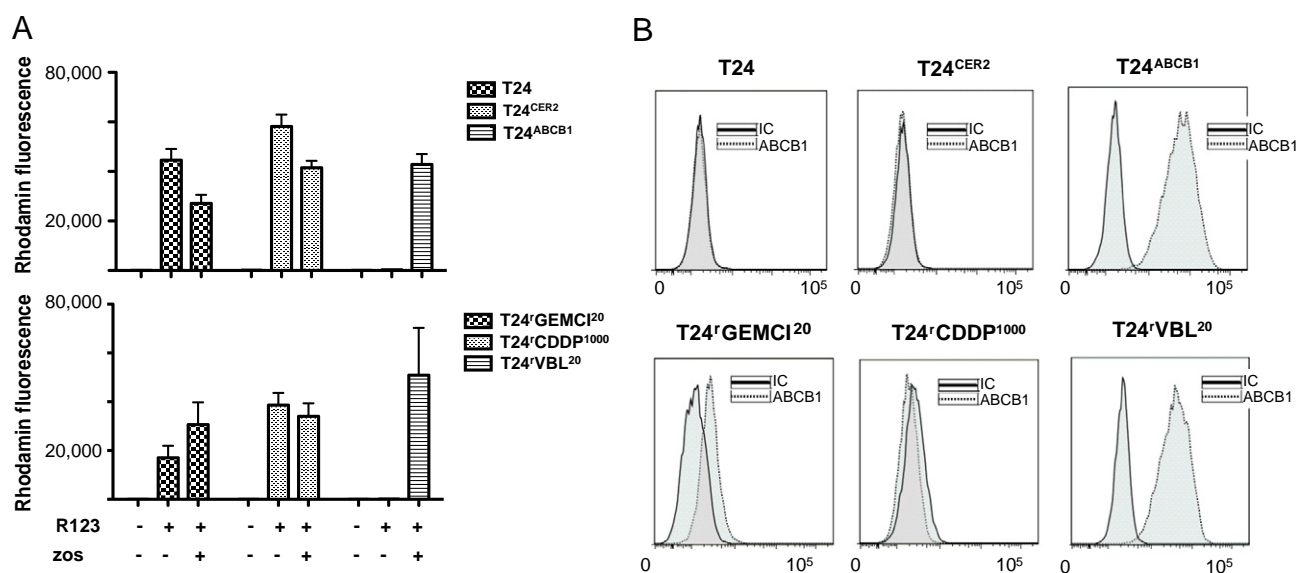
Figure 4. (A) R123 fluorescence of T24, T24^{CER2}, T24^{ABCB1}, T24^{r}GEMCI^{20}, T24^{r}CDDP^{1000}, and T24^{r}VBL^{20} cells after staining of R123 alone and in combination with 1.25 μM zosuquidar. Values are means ± SD. (B) ABCB1 expression in T24, T24^{CER2}, T24^{ABCB1}, T24^{r}GEMCI^{20}, T24^{r}CDDP^{1000}, and T24^{r}VBL^{20} cells. *$P \leq$ .05 relative to parental cell line.

survival advantage of 2.4 months over best supportive care [26]. Vinblastine is a constituent of methotrexate, vinblastine, doxorubicin, and cisplatin (MVAC), an alternative therapy protocol for metasized urothelial cancer [27]. This suggests that testing of drug candidates in drug-resistant cell lines holds potential for identification of next-line therapies.

Several studies demonstrated ABCB1 overexpression in tumor cells established from different human carcinomas (e.g., ovarian, stomach, colon) after adaptation to cisplatin [28–30]. Since cisplatin is not an ABCB1 substrate, this is probably the consequence of a long-term non-specific stress response [31,32]. Interestingly, only gemcitabine- but not cisplatin-resistant cells displayed ABCB1 up-regulation in our study, being to our knowledge the first report describing gemcitabine-induced ABCB1 up-regulation in urothelial tumor cells. Thus, ABCB1 expression may affect the efficacy of candidate drugs for second-line therapies of urothelial cancer after GC failure. In addition, ABCB1 may also modulate the malignant properties of cancer cells (e.g., cell survival, cell proliferation, and cell invasion) independently of the transporter-mediated drug efflux [4,33]. Indeed, ABCB1 expression correlated with an advanced tumor grade or with an increasing risk of recurrence in urothelial carcinoma patients [7,9]. Therefore, it will be important to show to which extent GC treatment is associated to an increased ABCB1 expression.

Cabazitaxel is a taxane that was recently approved for treatment of castration-resistant prostate cancer [34]. Currently, two clinical trials that investigate cabazitaxel in advanced bladder cancer are ongoing (NCT01616875 and NCT01668459). In contrast to taxanes including paclitaxel and docetaxel that have been used for decades, cabazitaxel is a weaker ABCB1 substrate [34]. In accordance, ABCB1-expressing TCC-SUP^{r}GEMCI^{20} cells remained sensitive to cabazitaxel. Interestingly, cabazitaxel's resistance profile differed from those of the other three taxanes in our study. TCC-SUP-^{r}GEMCI^{20} and T24^{r}GEMCI^{20} cells displayed cross-resistance to docetaxel, paclitaxel, and larotaxel, another taxane under clinical evaluation [35]. Moreover, HT1376^{r}GEMCI^{20} cells were cross-resistant to larotaxel and paclitaxel. In contrast, RT112^{r}GEMCI^{20} and RT112^{r}CDDP^{1000} cells showed cross-resistance to cabazitaxel. Our results show how complex the effects of apparently closely related compounds can be and that cancer cell line panels are suitable to identify such differences.

Among platinum derivates, cisplatin and carboplatin are thought to share a very similar mode of anticancer action [36,37]. In urothelial carcinoma, cisplatin was found to be superior to carboplatin in a randomized phase 2 study [38]. The European Association of Urology recommends carboplatin as a less toxic alternative in patients that are unfit for cisplatin because of bad performance status or elevated creatinine levels [39]. In concert with the anticipated closely related mechanisms of action of cisplatin and carboplatin [36,37], five of six cisplatin-resistant cell lines were also resistant to carboplatin. Interestingly, also three of six gemcitabine-resistant sublines displayed decreased sensitivity to cisplatin (RT112^{r}GEMCI^{20}, TCC-SUP-^{r}GEMCI^{20}, and T24^{r}GEMCI^{20}), and additionally, three of six gemcitabine-resistant sublines showed reduced sensitivity to carbo-platin (RT112^{r}GEMCI^{20}, TCC-SUP^{r}GEMCI^{20}, and RT4-^{r}GEMCI^{10}). Therefore, cross-resistance against cisplatin and carboplatin seems to be common after gemcitabine resistance.

Oxaliplatin supposedly differs in its anticancer mechanism of action from those exerted by cisplatin and carboplatin [40]. In concordance with this, oxaliplatin differed clearly in its activity profile from the cisplatin and carboplatin efficacy patterns. Only one gemcitabine- (TCC-SUP^{r}GEMCI^{20}) and two cisplatin-resistant cell lines (RT112^{r}CDDP^{1000} and TCC-SUP^{r}CDDP^{1000}) showed cross-resistance to oxaliplatin. In this context, gemcitabine/oxaliplatin combination therapy was suggested as an alternative for urothelial cancer patients unfit for cisplatin [41,42].

Conclusions

Here, we established a novel panel of gemcitabine- and cisplatin-resistant urothelial cancer cell lines. Cross-resistance profiles identified vinflunine, the European Medicines Agency–approved second-line therapeutic for urothelial cancer, together with vinblastine as the most effective drugs. This emphasizes the potential of panels of cancer cell lines with acquired drug resistance as preclinical models for the identification of potential next-line therapies after treatment failure. Notably, ABCB1 expression was detected in two gemcitabine-resistant cell lines although gemcitabine is not an ABCB1 substrate. The clinical relevance of these findings needs to be further investigated. Larger cell line panels may better reflect the complex processes of resistance formation in urothelial cancer cells [15]. Thus, the panel of drug-resistant urothelial carcinoma cell lines will be further expanded and characterized during ongoing research.

Supplementary data to this article can be found online at http://dx.doi.org/10.1016/j.tranon.2015.04.002.

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