Original Article

Relation of stem cell markers ALDH1 and CD44 with clinicopathological factors in urothelial carcinomas of urinary bladder

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Abstract: Molecular studies are ongoing in regards to superficial urothelial carcinoma of the bladder (UCB) either to define targeted therapy or to better select aggressive therapy candidates and also to delineate the outcome of the disease. In this study, we aimed to present the impact of ALDH1 and CD44 as stem cell markers in tumorigenesis and their prognostic value in urothelial carcinoma. We investigated ALDH1 and CD44 immunohistochemically in paraffin-embedded material of 125 non-muscle-invasive (NMI) cases in 163 UCB patients. In the NMI-UCB subgroup, we found ALDH1 to be significantly correlated with all poor prognostic factors, including high stage (≥pT2), high grade, recurrence and progression development and poor survey (P=0.001) in contrast to CD44 expression (P>0.05). Although ALDH1 expression had a good correlation with a poor clinical course of UCB, it could be used as a molecular marker to determine the best treatment strategy and could contribute to the development of targeted therapies.

Keywords: ALDH1, CD44, bladder carcinoma, prognostic value

Introduction

As one the most common human malignancies, urothelial carcinomas of the bladder (UCB) can present with non-muscle-invasive (NMI; pTa or pT1) tumors that can be treated with conservative approaches. However, recurrences and grade and/or stage progression are occasionally seen. Nonetheless, muscle-invasive (MI; ≥pT2) bladder cancers have high risks of disease progression and metastasis. These unfavorable prognoses of MI tumors have been used to propose additional prognostic indicators to identify the subset of NMI tumors that are likely to progress [1-3]. Within this context, the roles of stem cell markers have intensified in recent years in tumorigenesis and prognosis due to emerging cancer stem cell theory [4]. The cancer stem cells have the capacity for self-renewal and are also thought to be able to

differentiate into other cancer stem cell types [5, 6].

Aldehyde dehydrogenase 1 (ALDH1) and CD44 expressions are related with stem cells and are supposed to be predictors of cancer stem cells in a variety of tumors [4, 6-9]. ALDH1 is an enzyme that takes part in the synthesis and regulation of retinoic acid (RA). It plays a crucial role in the differentiation and regulation of the self-renewing ability of either normal or cancer stem cells [10].

It has been reported that ALDH1 expression has a relationship with the prognostic factors and/or resistance to treatment in breast [11], colorectal [12], lung [13], ovarian [14], esophageal [15], stomach [16], and bladder [17] cancers. Therefore, in our study, we aimed to investigate the expression of CD44 and ALDH1 as

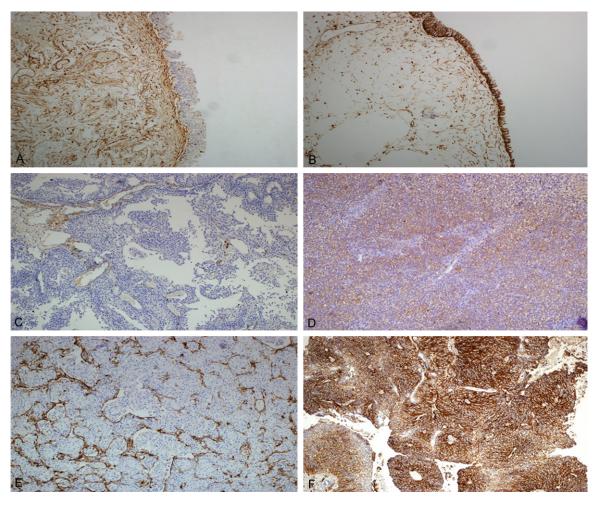


Figure 1. ALDH1 (A) and CD44 (B) expressions in normal urinary bladder (×100); positive immunohistochemical staining of ALDH1 in high grade UBC (×100) (C); negative immunohistochemical staining of ALDH1 in NMI-UCB (Low grade) (×100) (D); positive membranous immunohistochemical staining of CD44 in UBC (×100) (E); Loss of immunoreactivity of CD44 (×100) (F).

stem cell markers in UCB to determine the possible relationship between clinicopathological factors that have been proven to play a role in the prognosis in advance.

Materials and methods

This study included 163 cases of UCB that were diagnosed between 2006-2014 at the Istanbul Medeniyet University Goztepe Training and Research Hospital Pathology Department. All the sections were reviewed to confirm the original diagnosis, were staged according to the 2010 American Joint Committee on Cancer guidelines [18], and were graded according to the 2004 World Health Organization classification system [19] by two expert pathologists (SS and BC). All cases were graded according to the

degree of nuclear pleomorphism, anaplasia, and architectural distortion [19].

Age, gender, grade, pT stage, presence of carcinoma *in situ*, recurrence rates, progression, and survey data of cases were compared with the immunohistochemical findings. Cases were divided into the two main groups of non-muscular invasion transitional cell carcinoma (NMI-UCB) or muscular invasion transitional cell carcinoma (MI-UCB) according to the depth of invasion and the pT stage (group 1: pT2 and group pT3).

Tissue microarray construction (TMA)

Paraffin-embedded tumor tissues and hemotoxylin-eosin (H & E) slides were compared to

Table 1. The features of patients and immunohistochemical findings

| | Total | NMI (Ta-T1) | | |
|-------------|------------|-------------|--|--|
| | n (%) | n (%) | | |
| Gender | | | | |
| Male | 109 (66.9) | 90 (72.0) | | |
| Female | 54 (33.1) | 35 (28.0) | | |
| Age | | | | |
| <65 years | 73 (44.8) | 56 (44.8) | | |
| ≥65 years | 90 (55.2) | 69 (55.2) | | |
| First Grade | | | | |
| Low | 75 (46.0) | 69 (55.2) | | |
| High | 88 (54.0) | 56 (44.8) | | |
| CIS | | | | |
| Absent | 140 (85.9) | 110 (88.0) | | |
| Present | 23 (14.1) | 15 (12.0) | | |
| Status | | | | |
| Live | 123 (75.5) | 107 (85.6) | | |
| Exitus | 40 (24.5) | 18 (14.4) | | |
| Recurrence | | | | |
| Absent | 84 (51.5) | 70 (56.0) | | |
| Present | 79 (48.5) | 55 (44.0) | | |
| Progression | | | | |
| Absent | 113 (69.3) | 98 (78.4) | | |
| Present | 50 (30.7) | 27 (21.6) | | |
| ALDH1 | | | | |
| Negative | 102 (62.6) | 89 (71.2) | | |
| Positive | 61 (37.4) | 36 (28.8) | | |
| CD 44 | | | | |
| Negative | 38 (23.3) | 30 (24.0) | | |
| Positive | 125 (76.7) | 95 (76.0) | | |

cylinder samples with a4-mm diameter, using a manual tissue microarrayer (Quick Ray; Unitma Co. Ltd., Seoul, Korea), and these blocks were prepared for immunochemical staining.

Immunohistochemistry and scoring

Sections 4 μm in thickness were taken from 163 UCB tissues and compared with the H & E sections. Immunohistochemical staining for ALDH1 and CD44 was performed on step sections of TMA blocks according to the Bond Max Autostainer (Leica Biosystems) manufacturer's procedure and protocol.

CD44 staining at the membrane of the tumor cells was evaluated with scores of zero to 3 +

for 0-10, 10-25, 25-50, and more than 50 percent of tumor cell positivity, respectively. The staining intensity was classified from 1-3 (1, weak; 2, moderate; and 3, strong) [20, 21].

Cytoplasmic ALDH1 immunoreactivity was scored based upon the percent of tumor cell positivity from 0-100 and the staining intensity from 1-3 (1, weak; 2, moderate; and 3, strong). The product of the percentage of positive cells and the staining intensity was then divided by 4, making the reactivity score range from 0 to 100. The results of immunostaining were classified as negative or a score of 0, 1, 2 and 3 when the reactivity score was <3, 3-25, 26-50 and ≥50 percent, respectively [22].

Scores were calculated by multiplying the rate and intensity as follows: for CD44, 0-2 was scored as0 and 3-9 was scored as 1, while for ALDH1, 0-1 was scored as 0 and 2-9 received a score of 1, which were re-classified as semi-quantitative.

Statistical analysis

Data were analyzed by descriptive statistical methods (mean, standard deviation, median, frequency, and rate) and for the comparison of qualitative data using Pearson's chi-square test and Fisher Yates Continuity Correction, Freeman Halton (Monte Carlo) tests. For the immunohistochemical evaluation of variable cross-compliance, Spearman's correlation analysis was employed. Significance was set at P<0.05.

Clinicopathological data were evaluated using a Log Rank test and univariate and multivariate Cox regression analyses. Statistical analyses were performed using the Number Cruncher Statistical System (NCSS, 2007) and Power Analysis and Sample Size (PASS, 2008) statistical software (NCSS LLC, Kaysville, Utah, USA).

Results

According to the tumor stage, 125 patients (76.7%) were classified as NMI-UCB (Ta-T1), and 38 patients (23.3%) were placed in the MI-UCB (T2-T4) subgroup. The general follow-up period ranged from to to 72 months with an average of 33.50±14.39 months. ALDH1 immunoreactivity was observed mainly in the cytoplasm, whereas CD44 immunoreactivity was

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Table 2. Clinicopathological parameters and cancer stem cell markers in total UCB group

| | | | ALDH1 | | | CD 44 | |
|--------------------|------------------|-----------|------------|-----------|-----------|------------|--------------------|
| UCB | | Absent | Present | D.\/=\ | Absent | Present | P Value |
| | | n (%) | n (%) | - P Value | n (%) | n (%) | |
| Age | <65 years (n=73) | 49 (67.1) | 24 (32.9) | 0.280° | 15 (20.5) | 58 (79.5) | 0.572b |
| | ≥65 years (n=90) | 53 (58.9) | 37 (41.1) | | 23 (25.6) | 67 (74.4) | |
| Gender | Male (n=109) | 69 (63.3) | 40 (36.7) | 0.786ª | 26 (23.9) | 83 (76.1) | 0.972b |
| | Female (n=54) | 33 (61.1) | 21 (38.9) | | 12 (22.2) | 42 (77.8) | |
| First pT | Ta-T1 (n=125) | 89 (71.2) | 36 (28.8) | 0.001**,b | 30 (24.0) | 95 (76.0) | 0.875 ^b |
| | T2-T4 (n=38) | 13 (34.2) | 25 (65.8) | | 8 (21.1) | 30 (78.9) | |
| First Grade | Low (n=75) | 57 (76.0) | 18 (24.0) | 0.001**,a | 15 (20.0) | 60 (80.0) | 0.461 ^b |
| | High (n=88) | 45 (51.1) | 43 (48.9) | | 23 (26.1) | 65 (73.9) | |
| CIS | Negative (n=140) | 95 (67.9) | 45 (32.1) | 0.001**,b | 34 (24.3) | 106 (75.7) | 0.646b |
| | Positive (n=23) | 7 (30.4) | 16 (69.6) | | 4 (17.4) | 19 (82.6) | |
| Recurrence | Absent (n=84) | 67 (79.8) | 17 (20.2) | 0.001**,b | 18 (21.4) | 66 (78.6) | 0.688b |
| | Present (n=79) | 35 (44.3) | 44 (55.7) | | 20 (25.3) | 59 (74.7) | |
| Progression | Absent (n=113) | 87 (77.0) | 26 (23.0) | 0.001**,b | 25 (22.1) | 88 (77.9) | 0.735 ^b |
| | Present (n=50) | 15 (30.0) | 35 (70.0) | | 13 (26.0) | 37 (74.0) | |
| Status (Mortality) | Live (n=123) | 92 (74.8) | 31 (25, 2) | 0.001**,b | 24 (19.5) | 99 (80.5) | 0.072 ^b |
| | Exitus (n=40) | 10 (25.0) | 30 (75.0) | | 14 (35.0) | 26 (65.0) | |

^aPearson Ki-square Test; ^bFisher-Freeman-Halton Test; **P<0.01.

Table 3. Clinicopathological parameters and cancer stem cell markers in NMI group

| | | ALDH 1 | | CD 44 | | | | |
|--------------------|------------------|-----------|-----------|-----------|-----------|-----------|--------------------|--|
| NMI (Ta-T1) | | Absent | Present | – P Value | Absent | Present | - D\/olus | |
| | | n (%) | n (%) | P value | n (%) | n (%) | P Value | |
| Age | <65 years (n=73) | 42 (75.0) | 14 (25.0) | 0.518° | 11 (19.6) | 45 (80.4) | 0.414° | |
| | ≥65 years (n=90) | 47 (68.1) | 22 (31.9) | | 19 (27.5) | 50 (72.5) | | |
| Gender | Male (n=109) | 64 (71.1) | 26 (28.9) | 1.000° | 23 (25.6) | 67 (74.4) | 0.675° | |
| | Female (n=54) | 25 (71.4) | 10 (28.6) | | 7 (20.0) | 28 (80.0) | | |
| First Grade | Low (n=75) | 57 (82.6) | 12 (17.4) | 0.003**,c | 13 (18.8) | 56 (81.2) | 0.198° | |
| | High (n=88) | 32 (57.1) | 24 (42.9) | | 17 (30.4) | 39 (69.6) | | |
| CIS | Negative (n=140) | 83 (75.5) | 27 (24.5) | 0.012*,b | 29 (26.4) | 81 (73.6) | 0.116 ^b | |
| | Positive (n=23) | 6 (40.0) | 9 (60.0) | | 1 (6.7) | 14 (93.3) | | |
| Recurrence | Absent (n=84) | 59 (84.3) | 11 (15.7) | 0.001**,c | 16 (22.9) | 54 (77.1) | 0.899⁰ | |
| | Present (n=79) | 30 (54.5) | 25 (45.5) | | 14 (25.5) | 41 (74.5) | | |
| Progression | Absent (n=113) | 79 (80.6) | 19 (19.4) | 0.001**,c | 23 (23.5) | 75 (76.5) | 0.992° | |
| | Present (n=50) | 10 (37.0) | 17 (63.0) | | 7 (25.9) | 20 (74.1) | | |
| Status (Mortality) | Live (n=123) | 83 (77.6) | 24 (22.4) | 0.001**,c | 24 (22.4) | 83 (77.6) | 0.372b | |
| | Exitus (n=40) | 6 (33.3) | 12 (66.7) | | 6 (33.3) | 12 (66.7) | | |

 $^{{}^{\}text{a}}\text{Pearson Ki-sqareTest;} \ {}^{\text{b}}\text{Fisher-Freeman-Halton Test;} \ {}^{\text{c}}\text{Yates' Continuity Correction Test;} \ {}^{\text{+}}\text{P}<0.05\ {}^{\text{+}}\text{P}<0.01.$

seen in the cytoplasmic membrane. If focal staining on the basal layer were excluded, ALDH1 immunoreactivity was not observed in the 20 control cases with normal bladder transitional epithelium (**Figure 1A**). CD44 expres-

sion was represented in the normal bladder epithelium (Figure 1B).

In all UCB cases, ALDH1 expression was not observed in 102 (62.6%) patients (**Figure 1C**),

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Table 4. ALDH1 and CD44 assessment using Cox regression analysis in UCB

| | Recurrence | | | | Progression | | | Overall Survival | | |
|----------|------------|--------------|---------|-------|--------------|---------|-------|------------------|---------|--|
| | HR | 95% CI | P Value | HR | 95% CI | P Value | HR | 95% CI | P Value | |
| ALDH1 | | | | | | | | | | |
| Age>65 | 2.986 | 1.303-6.841 | 0.010* | 2.990 | 1.330-6.718 | 0.008** | 2.651 | 1.170-6.006 | 0.019* | |
| Gender | 1.022 | 0.455-2.293 | 0.958 | 0.916 | 0.408-2.056 | 0.832 | 1.100 | 0.477-2.536 | 0.823 | |
| First pT | 3.693 | 1.778-7.670 | 0.001** | 4.026 | 2,004-8.090 | 0.001** | 3.139 | 1.563-6.301 | 0.001 | |
| Grade | 1.138 | 0.463-2.800 | 0.778 | 1.858 | 0.826-4.179 | 0.134 | 1.962 | 0.874-4.404 | 0.102 | |
| CIS | 0.974 | 0.395-2.407 | 0.955 | 0.871 | 0.346-2.193 | 0.769 | 1.006 | 0.409-2.477 | 0.990 | |
| ALDH1 | 4.590 | 2.042-10.319 | 0.001** | 3.222 | 1.454-7.140 | 0,004** | 3.109 | 1.383-6,985 | 0.006** | |
| CD44 | | | | | | | | | | |
| Age>65 | 2.834 | 1.319-6.088 | 0.008** | 3.110 | 1.427-6.780 | 0.004** | 2.602 | 1.195-5.667 | 0.016* | |
| Gender | 1.129 | 0.554-2.343 | 0.745 | 1.134 | 0.540-2.382 | 0.740 | 1.463 | 0.687-3.116 | 0.324 | |
| First pT | 5.306 | 2.526-11.144 | 0.001** | 6.279 | 3.114-12.663 | 0.001** | 4.832 | 2.393-9.760 | 0.001** | |
| Grade | 1.711 | 0.700-4.183 | 0.239 | 2.198 | 0.957-5.045 | 0.063 | 2.177 | 0.956-4.953 | 0.064 | |
| CIS | 1.227 | 0.519-2.899 | 0.641 | 1.164 | 0.488-2.778 | 0.732 | 1,479 | 0.626-3.496 | 0.372 | |
| CD 44 | 0.548 | 0.283-1.059 | 0.074 | 0.516 | 0.266-1.00 | 0.050 | 0.508 | 0.260-0.995 | 0.058 | |

^{*}P<0.05; **P<0.01.

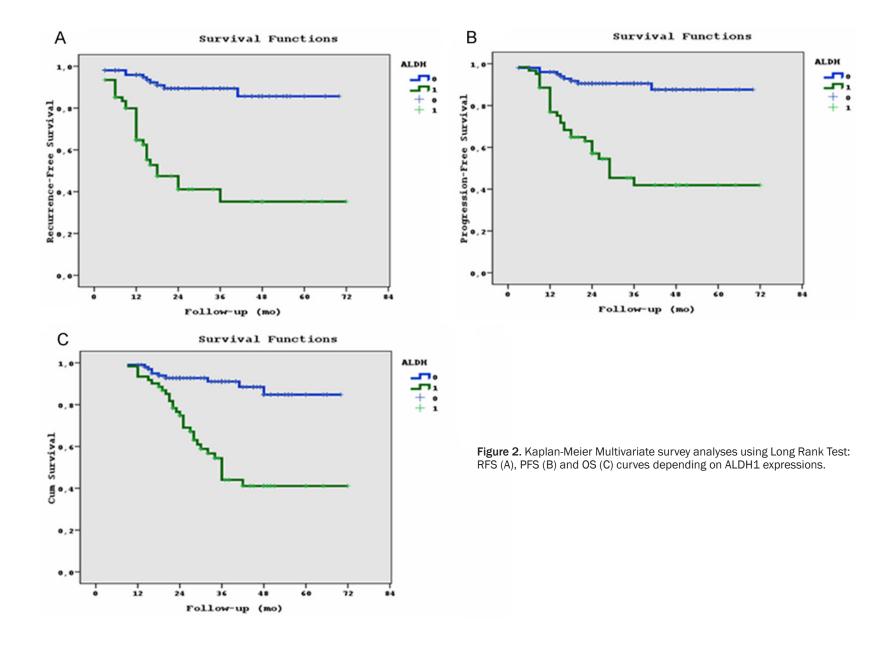
but 61 (37.4%) patients were positive (**Figure 1D**). In the NMU-UCB subgroups, these numerical values were 89 (71.2%) and 36 (28.8%), respectively. CD44 expression was negative in 38 (23.3%) patients (**Figure 1E**), while it was positive in 125 (76.7%) patients (**Figure 1F**). Assessment data for our clinicopathological and immunohistochemical findings are given in **Table 1**.

ALDH1 expression was significantly associated with carcinoma in situ (P<0.01) with a high stage (≥pT2) (P=0.001) and a higher grade (P<0.01) in UCB. In addition, ALDH1 expression in UCB as well as in the NMI-UCB subgroup was significantly associated with recurrence [P<0.05 in the NMI-UCB subgroup], progression to a higher stage, and low survey (P=0.001). The immune expression of CD44 had no statistical correlation with any descriptive data or clinicopathological parameters. The association between ALDH1 and CD44 expression and clinicopathological characteristics are summarized in **Table 2** and **Table 3**.

The recurrence-free follow-up period ranged between 3 and 72 months, with an average of 23.60±16.88 months. Additionally, the progression-free follow-up period was between 3 and 72 months with an average of 29.58±16.26 months.

In the multivariate Cox regression analysis, ALDH1 expression both in the total UCB group and the NMU-UCB subgroup was identified as a significant predictor that could determine tumor recurrence, progression, and survival. However, CD44 expression was unrelated to the prognostic parameters in Cox regression analysis (P>0.05) (Table 4).

Kaplan-Meier analysis was carried out using the Long Rank test and revealed that the recurrence-free survival rate and average survival time were 50.8% and 34.73±4:50 months in ALDH1-positive patients, respectively, whereas they were 90.2% and 62.82±2.17 months in ALDH1-negative cases, respectively. The progression-free survival rate was 50.8%, and the average survival time was 40.76±3.94 months in ALDH1-positive cases. In negative cases, the survival rate was 90.2%, and the average survey length was 63.67±1.91 months, so a statistically significant difference was present. A significant difference was also seen for the ALDH1positive patients whose overall survival rate was 50.8%; the average survival time was 44.85±3.35 months. In negative cases, the survival rate was 90.2% and the average survival time was 63.85±1.84 months (P<0.01). When compared with ALDH1 positivity, rates of recurrence-free time, progression-free time, and overall survival were statistically significantly increased in ALDH1 negativity (Kaplan-



Meier, log-rank test =0.001; P<0.01) (**Figure 2**). No statistically significant difference with CD44 expression was observed (P>0.05).

Discussion

The parameters that should be used to determine the appropriate medical or surgical management of UCB are still controversial. The European Organization for Research and Treatment of Cancer (EORTC) scoring system currently assists professionals in determining the indications for cystectomy based upon parameters such as tumor diameter, grade, stage (pT), recurrence rate, and presence of carcinoma in situ. In addition to these parameters, ongoing immunohistochemical and molecular studies have attempted to identify the molecules that play a role in tumorigenesis [23]. In order to determine the risk of recurrence and progression of superficial/non-invasive bladder carcinoma, these new molecules might give rise to developments for new treatment modalities.

Our study had several limitations, including the probability of tumor heterogeneity, the effect of a residual tumor on recurrence, and the moderately short follow-up periods. Other investigations on stem cell markers have also been associated with poor prognostic factors and the development of resistance to chemo-radiotherapy [8, 9, 24-27]. In addition, while several studies have reported a statistically significant association between CD44 and poor prognosis-surveillance [28-30], we did not identify any significant prognostic association of CD44 with UCB in our study (P>0.05). However, Keymoosi et al. recently identified a correlation of CD44 in combination with ALDH1 with a poor prognosis in bladder cancer [31].

In contrast, ALDH1 expression has been correlated with a poor prognosis and resistance to therapy in tumors of the breast [20, 25], colon [12, 22], stomach [16], lung [13], and pancreas [26], and it has been suggested to be a negative prognostic factor [27]. For bladder carcinoma, Kitamura et al. [17] reported that ALDH1 has an effective role in bladder carcinoma development as its expression was negative in control bladder tissue. In our study, the results indicated that ALDH1 expression was significantly associated with poor surveillance and poor prognostic factors, such as high-stage,

high grade, recurrence, progression, and presence of concomitant carcinoma in situ either in UCB (P<0.01) or in the NMU-UCB subgroup (P<0.01).

Moreover, Cox regression analyses and multivariate analyses detected a significant relationship with ALDH1 and survivals, such as the recurrence-free survival (RFS), progression-free survival (PFS), and the overall survival (OS), respectively. In light of these data, we suggest that ALDH1 might contribute to tumorigenesis in UCB. Additionally, the presence of ALDH1 could be a high risk factor for a poor prognosis, as ALDH1-positive tumors tend to be biologically aggressive. This hypothesis was also implied by our results that the expression of ALDH1 was correlated with indicators of poor prognosis, such as invasion into the muscularis propria and poorly differentiated tumors.

The present study showed that ALDH1 expression levels may be used as an indicator to determine the potential of tumor recurrence and progression within cases where T1 staged-TUR materials were present or the deep muscle layer was not seen.

Finally, we propose that inhibiting ALDH1 may provide new insights and can be an effective treatment for urothelial carcinoma eradication. This marker should be considered in the development of new diagnostic and therapeutic approaches for bladder cancer.

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Disclosure of conflict of interest

None.

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