Androgen receptor mRNA expression is a predictor for recurrence-free survival in non-muscle invasive bladder cancer

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Abstract

Background: Non-muscular invasive bladder cancer (NMIBC) has a high risk of recurrence. As androgen receptor (AR) reportedly affects bladder cancer, we assessed the correlation between NMIBC recurrence and tumor AR expression in Japanese patients.

Methods: We retrospectively reviewed 53 specimens of non-metastatic NMIBC, with recurrence-free survival (RFS) as the primary endpoint. We used real-time quantitative polymerase chain reaction to quantify AR mRNA expression. Kaplan–Meier product-limit estimators were used to assess RFS distribution, log-rank tests to analyze differences in RFS between high- and low-risk groups; and multivariate analyses of AR mRNA expression and other clinicopathological factors to predict independent factors for RFS.

Results: The high AR mRNA-expressing group (n = 43) tended to have a longer median RFS (not reached) than did the low-AR group (n = 10; 9.04 months; P = 0.112). Multivariate analysis showed female sex (hazard ratio [HR]: 7.360, 95% CI: 1.649–32.856, P = 0.009), tumor size ≥3 cm (HR: 23.697, 95% CI: 4.383–128.117, P < 0.001) and low AR mRNA expression (HR: 0.202, 95% CI: 0.048–0.841, P = 0.028) to be independent predictors of shorter RFS.

Conclusion: Our study showed that low AR mRNA expression level is an independent risk factor for RFS in Japanese patients with NMIBC. Further studies are necessary but AR expression might be a new indicator of recurrence of NMIBC.

Keywords: Bladder cancer, Non-muscular invasive, Androgen receptor, RT-qPCR, Recurrence

Background

Bladder cancer (BC) is a common malignancy, with an estimated 429,800 new cases and 165,100 deaths in 2012, worldwide [1]. Urothelial carcinoma is the most common type of BC, and it is stratified to non-muscular invasive bladder cancer (NMIBC) and muscular invasive bladder cancer (MIBC). Patients with MIBC have a high risk of disease progression and metastasis and usually need aggressive treatments, such as radical cystectomy or chemotherapy; whereas NMIBC has a better prognosis and can be treated curatively by transurethral resection of bladder tumor (TUR-BT). However, NMIBC has a high risk of recurrence and can progress to MIBC. To prevent recurrence and progression, intravesical instillation (IVI) of anthracycline or bacillus Calmette-Guérin (BCG) has been administered, but about 50% still recur. As the treatment has not changed for decades, new studies leading to new treatments are needed to prevent recurrence and progression.

Men tend to be diagnosed about four times more often than women [2], which implies that BC is an endocrine-related cancer. Androgen receptor (AR) signaling has been suggested to have an important role in BC occurrence and progression by previous studies [3–18].

Thus, in this study, we assessed the correlation between NMIBC recurrence and tumor AR expression in Japanese patients, by quantifying AR mRNA expression.
with real-time quantitative polymerase chain reaction (RT-qPCR).

Methods

Patients

We retrospectively reviewed 53 specimens of TUR-BT among 51 patients with non-metastatic NMIBC treated at Yokohama City University Medical Center and Yokohama City University Graduate School of Medicine from 2008 to 2015. In the two patients, specimens were obtained two times at the time of initial TUR-BT and at the time of recurrence. Samples for RT-qPCR came from biopsies performed during TUR-BT procedures. Both primary and recurrent NMIBC was diagnosed by TURBT. The experimental procedures were conducted in accordance with the ethical standards of the Helsinki Declaration. This study was approved by the institutional review board of Yokohama City University Medical Center. Informed consent to participate in the study were obtained from all subjects.

Clinical assessments

The primary endpoint of this study was recurrence-free survival (RFS), which was defined as the time between the date of the initial TUR-BT and the date when the tumor relapsed. Tumors detected in second-look TUR were not defined as recurrence, and RFS was counted from the first TUR-BT. Histopathological factors were assessed by pathologists specialized in uropathology.

Real-time PCR

AR mRNA level was assessed by quantitative RT-qPCR method. We obtained 53 BC tissue specimens by TUR-BT to analyze gene expression. Total RNA (0.5 μg), which was isolated from the bladder tissue specimens, using Isogen (NipponGene, Tokyo, Japan), was reverse transcribed using 1 μM oligo (dT) primers (Qiagen, Germantown, MD, USA) and 4 units of Omniscript reverse transcriptase (Qiagen) in a total volume of 20 μL. Gene expression for AR and human glyceraldehyde-3-phosphate dehydrogenase (GAPDH) were examined by real-time qPCR, using AR primers (Applied Biosystems, Foster City, CA, USA). Gene expressions were determined using TaqMan® Gene Expression Assays (Applied Biosystems, Grand Island, NY, USA). All data were normalized to GAPDH; expression of one tumor was assumed to be 1 and used as reference. The 2^−ΔΔCT method was used to calculate relative amounts of target genes.

Statistical analysis

The cut-off point for the AR mRNA level was determined by a web-based calculator cut-off finder (http://molpath.charite.de/cutoff) and median values were used as the cut-off points for other factors to categorize continuous measurements [19]. Spearman’s rank-order correlation was used to measure the association between AR mRNA expression level and clinical factors. A Kaplan–Meier product-limit estimator was used to assess RFS distribution. Log-rank tests were used to analyze differences in RFS between the high- and low-risk groups. Univariate and multivariate COX models were used to analyze factors for predicting RFS. We derived relative risks and 95% confidence intervals (95% CIs). All tests were two-sided; alpha = 0.05 was considered significant. All analyses were conducted using IBM SPSS Statistics software for Windows, version 22 (IBM Corp., Armonk, NY, USA).

Results

Patients’ characteristics are shown in Table 1. Patients’ median age at TUR-BT was 74.1 years (range: 40.8–88.8 years) and 48 (90.6%) patients were male. Of the 53 patients, 11 (20.8%) were recurrent BC and 9 patients (17.0%) were recurrent within 1 year. Tumor grade was low in 39 patients (73.6%) and high in 14 patients (26.4%). Pathological T-stage was pTa in 42 patients (79.2%) and pT1 in 11 patients (20.8%). Tumors were single in 24 patients (45.3%) and multiple in 29 patients (54.7%). Tumor size was ≥3 cm in 7 patients (13.2%); concurrent carcinoma in situ (CIS) was found in 2 patients (3.8%). Immediate IVI using anthracycline was administered in 51 patients (96.2%) and additional adjuvant IVI using either BCG or anthracycline was administered in 12 patients (22.6%). Six patients (11.5%) underwent second TURs.

We found no significant association between AR mRNA expression level and other clinicopathological factors (Spearman’s rank-order correlation; Table 2).

The Kaplan–Meier curve for RFS is shown in Fig. 1. Within a median observation period of 10.2 (range: 0.4–29.5) months, recurrence occurred in 17 patients (32.1%).

We stratified the 53 patients into two groups: 43 in the high AR mRNA-expressing group and 10 in the low AR group. The high AR group showed a tendency for longer median RFS (not reached) than did the low-AR group (9.04 months, P = 0.112; Fig. 2).

Results of univariate and multivariate analyses to identify risk factors for RFS are shown in Table 3. In univariate analysis, female sex (hazard ratio [HR]: 5.009, 95% CI: 1.589–15.787, P = 0.006) and tumor size ≥3 cm (HR: 3.487, 95% CI: 1.282–9.480, P = 0.0141) were significantly related to shorter RFS. Multivariate analysis showed that female sex (HR: 7.360, 95% CI: 1.649–32.856, P = 0.009), tumor size ≥3 cm (HR: 23.697, 95% CI: 4.383–128.117, P < 0.001) and low AR mRNA expression (HR: 0.202, 95% CI: 0.048–0.841, P = 0.028) were independent predictors of shorter RFS.
**Discussion**

Although the function of AR signaling in BC is unclear, it is known to play important roles in prostate cancer occurrence and progression, and reportedly affects kidney, lung, breast and liver cancers [20].

AR is a nuclear steroid hormone receptor, composed of several domains, N-terminal transactivation domain, DNA binding domain, a hinge region and a ligand-binding domain (LBD). In the cytoplasm, heat shock protein-90 (HSP90) binds with the LBD. As 5α-dihydrotestosterone binds to AR, AR releases HSP90 and translocates into the nucleus, binds to the androgen-response element, and promotes gene transcription [3–6, 21].

AR signaling and the urinary tract may be critically associated. Studies in animal models have shown AR to be present in several tissues in lower urinary tract and may affect growth, differentiation and maintenance of urinary bladder tissue [6, 7, 22, 23].

Reportedly, AR signaling has a pivotal role in BC occurrence and progression. Testosterone increased the risk of N-butyl-N-(4-hydroxybutyl) nitrosamine (BBN)-induced bladder tumors in female rats, whereas diethylstilbestrol, a synthetic form of female hormone estrogen, inhibited incidence in male rats [8]. Also, intact female rats administered with testosterone had higher incidences of both bladder calculi and tumors, but oophorectomized rats did not, which implies that testosterone in combination with estrogen may contribute to BC formation [9]. Moreover, flutamide, a non-steroidal antiandrogen therapy, inhibited BBN-induced bladder tumor incidence in rats, whereas finasteride, a 5α-reductase inhibitor, did not; this suggests that testosterone itself has an effect in bladder carcinogenesis but its converting form, 5α-dihydrotestosterone, does not [10].

Studies with animal models have indicated that AR has an important role in BC. Miyamoto et al. found that both male and female AR-knockout mice (ARKO) did not develop BBN-induced tumors, which suggests that AR has a crucial activity in bladder carcinogenesis [11]. That study also found that 25% of ARKO mice administered with dihydrotestosterone (DHT) developed tumors, compared with 50% of castrated male wild-type rats and 92% of intact wild-type male rats. These findings suggest that absence or low levels of androgen could restore AR signaling, or that androgens could induce bladder carcinogenesis independently of AR [7, 24]. Another study showed that mice lacking AR only in the urothelial tissues had lower incidence of BBN-induced BC and a higher survival rate than wild-type mice [12].

Therefore, the association between BC and AR has been the focus of several recent studies. Expression of AR have indicated to have high risk in bladder cancer occasion, but for recurrence, expression of AR have shown to contribute to longer RFS. An immunohistochemical study of 169 patients indicated that recurrence was less likely for patients with AR + BC specimens [13]; another study showed that loss of AR was strongly associated with higher grade and more invasive tumors [14]. A study of the predictive value of AR mRNA expression in pT1 NMIBC showed that high AR mRNA expression was an independent predictor for longer RFS and cancer-specific survival [15], which was similar to our present findings that high AR mRNA level was an independent predictor for RFS in NMIBC (including pTa and pT1).

Reportedly, AR pathway inhibitors have been shown to be effective in preventing BC recurrence. In patients

**Table 1 Patients’ characteristics**

| Age, median (range) | 74.1 (40.5–88.8) |
|---------------------|------------------|
| Sex, n (%)          |                  |
| Male                | 48 (90.6%)       |
| Female              | 5 (9.4%)         |
| Primary or recurrence, n (%) |              |
| Primary             | 42 (79.2%)       |
| Recurrence          | 11 (20.8%)       |
| Recurrence within one year, n (%) |          |
| Yes                | 9 (17.0%)        |
| No                 | 44 (83.0%)       |
| Tumor grade, n (%)  |                  |
| Low                | 39 (73.6%)       |
| High               | 14 (26.4%)       |
| T stage, n (%)      |                  |
| pTa                 | 42 (79.2%)       |
| pT1                | 11 (20.8%)       |
| Number of tumor, n (%) |               |
| Single              | 24 (45.3%)       |
| Multiple            | 29 (54.7%)       |
| Tumor size, n (%)   |                  |
| < 3 cm              | 46 (86.8%)       |
| ≥ 3 cm              | 7 (13.2%)        |
| Concurrent CIS, n (%) |             |
| Yes                 | 2 (3.8%)         |
| No                  | 51 (96.2%)       |
| Immediate intravesical chemotherapy, n(%) |            |
| Yes                 | 51 (96.2%)       |
| No                  | 2 (3.8%)         |
| Additional adjuvant intravesical chemotherapy, n (%) |         |
| Yes                 | 12 (22.6%)       |
| No                  | 41 (77.4%)       |
| Second TUR, n (%)   |                  |
| Yes                 | 6 (11.5%)        |
| No                  | 47 (88.5%)       |

CIS carcinoma in situ, TUR transurethral resection
with double primary cancers of the prostate and the blad-
ser, androgen-deprivation therapy (ADT) to treat their pro-
state cancers reduced the risk of recurrence of AR+ BC [16]. Enzalutamide, an AR-signaling inhibitor, is also re-
ported to inhibit AR+ BC cell growth in vivo [17]. BCG IVI has been the most effective therapy for NMIBC; its me-
chanism may be due to interaction between BCG and the AR pathway. DHT down-regulates NF-κB-mediated IL-6 expression, whereas AR blockers inhibit the effect of DHT. This result suggests that AR pathway inhibitors could improve the efficacy of BCG treatment [18].

As described above, previous clinical studies demon-
strated that higher AR expression in both mRNA levels and protein levels were correlated with favorable outcome in NMIBC [13–15]. Moreover, our current study also demonstrated that higher AR mRNA expression levels were associated with longer recurrence-free sur-
vival in NMIBC. On the other hand, enhanced AR sig-
aling pathway could play an important role in cancer initiation and progression in vitro and in vivo. There were discrepancies between our findings and previous reports from studies in vitro or in vivo [8, 10–12]. Even

| Table 2 | Spearman’s rank-order correlation between each variables |
|---------|--------------------------------------------------------|
|         | Sex     | Age     | T stage | Tumor grade | Tumor size | Number of tumor | Recurrence/Primary | Instillation | AR mRNA expression |
| Sex     | 1.000   | 0.071   | −0.006  | 0.099      | −0.126     | 0.034           | 0.153            | −0.020       | −0.174           |
| Age     | 0.071   | 1.000   | 0.056   | −0.160     | 0.063      | −0.093          | 0.056            | 0.010        | −0.009           |
| T stage | −0.006  | 0.056   | 1.000   | 0.432      | 0.075      | 0.372           | −0.262           | 0.057        | −0.110           |
| Tumor grade | 0.099 | −0.160  | 0.432   | 1.000      | 0.019      | 0.373           | −0.201           | 0.392        | −0.149           |
| Tumor size | −0.126 | 0.063   | 0.075   | 0.019      | 1.000      | −0.093          | −0.200           | −0.078       | 0.188            |
| Number of tumor | 0.034 | −0.093  | 0.372   | 0.373      | −0.093     | 1.000           | −0.095           | 0.492        | −0.051           |
| Recurrence/primary | 0.153 | 0.056   | −0.262  | −0.201     | 0.200      | −0.095          | 1.000            | 0.057        | −0.110           |
| Instillation | −0.020 | 0.010   | 0.057   | 0.392      | −0.078     | 0.492           | 0.057            | 1.000        | −0.085           |
| AR mRNA expression | −0.174 | −0.009  | −0.110  | −0.149     | 0.188      | −0.051          | −0.110           | −0.085       | 1.000            |

mRNA messenger ribonucleic acid

Fig. 1 Kaplan–Meier curve for RFS after TUR-BT. Within a median observation period of 10.2 months (range: 0.4–29.5 months), recurrence occurred in 17 patients (32.1%)
though the reasons for discrepancies are unknown, several possible hypothesis have been suggested. It has to be considered that AR expression, and thus the role of AR, might change during the progression of bladder cancer, as indicated by the previously cited studies [14]. In addition, Sikic et al. reported that different AR subtype (AR 1 and AR2) has different role in bladder cancer carcinogenesis [14].

The mechanism of AR signaling in BC is still unclear, therefore, further studies of the AR pathway in NMIBC are needed.

Our study had several limitations. First, it was a retrospective study with relatively few patients and a short follow-up period. Second, CIS was not analyzed in this study. CIS is an essential risk factor of recurrence and progression in NMIBC, but our study cohort had only two BCs with coexisting CIS. Finally, for verify the role of AR pathway in NMIBC, not only AR mRNA expression levels, but also protein expression levels should be investigated in current study. Future clinical studies would need to estimate an appropriate sample size to include enough of these patients and investigate mRNA and protein expression levels simultaneously.

**Conclusions**

Our study show that AR mRNA expression is an independent predictor for RFS in Japanese patients with NMIBC. The AR pathway is now attracting attention in BC research, and may provide new methods to treat bladder cancer and prevent its recurrence.

| Table 3 | Univariate and multivariate analyses for predicting recurrence-free survival |
|---------|--------------------------------------------------------------------------|
| **Sex** | Women vs men | 0.006 | 5.009 | 1.589 | 15.787 | 0.009 | 7.360 | 1.649 | 32.856 |
| **Age** | High vs low | 0.324 | 1.633 | 0.617 | 4.325 | 0.453 | 0.640 | 0.199 | 2.054 |
| **T stage** | pT1 vs pTa | 0.925 | 1.056 | 0.342 | 3.261 | 0.096 | 5.274 | 0.743 | 37.435 |
| **Tumor grade** | High vs low | 0.526 | 0.694 | 0.225 | 2.142 | 0.657 | 0.650 | 0.097 | 4.357 |
| **Tumor size** | ≥3 cm vs < 3 cm | 0.014 | 3.487 | 1.282 | 9.480 | <0.001 | 23.697 | 4.383 | 128.117 |
| **Number of tumor** | Multiple vs single | 0.189 | 0.527 | 0.203 | 1.370 | 0.477 | 1.688 | 0.398 | 7.148 |
| **Tumor recurrence** | Recurrence vs primary | 0.531 | 1.397 | 0.491 | 3.973 | 0.052 | 4.122 | 0.988 | 17.190 |
| **Instillation** | Yes vs no | 0.068 | 0.253 | 0.058 | 1.108 | 0.088 | 0.182 | 0.026 | 1.290 |
| **AR mRNA expression** | High vs low | 0.123 | 0.438 | 0.153 | 1.250 | 0.028 | 0.202 | 0.048 | 0.841 |

*mRNA* messenger ribonucleic acid, *HR* hazard ratio, *CI* confidence interval

**Fig. 2** Kaplan–Meier curve for recurrence-free survival (RFS), stratified by AR mRNA expression level. Green line: RFS curve for high AR mRNA expression level group; blue line: RFS for low AR mRNA expression level group. The high AR mRNA-expressing group tended to have a longer median RFS (not reached) than did the low-AR group (9.04 months; *P* = 0.112)
Abbreviations
AR: Androgen receptor; ARKO: AR-knockout (mice); BBN: N-butyl-N-(4-hydroxybutyl) nitrosamine; BC: Bladder cancer; BCG: Bacillus Calmette-Guérin; CIS: Carcinoma in situ; DHT: Dihydrotestosterone; HR: Hazard ratio; HSP90: Heat shock protein-90; IV: Intravesical instillation; LBD: Ligand-binding domain; MIBC: Muscular invasive bladder cancer; NMIBC: Non-muscular invasive bladder cancer; RFS: Recurrence-free survival; RT-qPCR: Real-time quantitative polymerase chain reaction; TUR-BT: Transurethral resection of bladder tumor; XIST: X-inactive specific transcript

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Authors’ contributions
Conceived and designed the experiments: MY, TK, KI, YM. Analyzed data: MY, TK, KI, YM. Performed the experiments: MY, TK, YI, HI. Wrote the paper: MY, TK, KI, MY, HU, YM. All authors read and approved the final manuscript.

Ethics approval and consent to participate
IRB approval for Yokohama City University. Informed consent was obtained from all individual participants included in the study.

Consent for publication
Not applicable.

Competing interests
The authors declare that they have no competing interests.

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