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High testosterone levels in prostate tissue obtained by needle biopsy correlate with poor-prognosis factors in prostate cancer patients

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Abstract

Background: There is currently no consensus on the correlations between androgen concentrations in prostate tissue and blood and stage and pathological grade of prostate cancer. In this study, we used a newly-developed ultra-sensitive liquid-chromatography tandem mass spectrometry method to measure testosterone (T) and dihydrotestosterone (DHT) concentrations in blood and needle biopsy prostate specimens from patients with prostate cancer.

Methods: We analyzed androgen levels in 196 men diagnosed with prostate cancer. All patients had undergone systematic needle biopsy, and an additional needle biopsy from the peripheral zone was conducted for the simultaneous determination of T and DHT. We analyzed the relationships between T and DHT levels in tissue and blood and Gleason score, clinical stage, and percentage of positive biopsy cores, using multivariate analysis.

Results: The median T and DHT levels in blood were 3551.0 pg/mL and 330.5 pg/mL, respectively. There was a strong correlation between serum T and DHT. The median T and DHT levels in prostate tissue were 0.5667 pg/mg and 7.0625 pg/mg, respectively. In multivariate analysis, serum prostate-specific antigen and tissue T levels were significantly associated with poor prognosis; high T levels in prostate tissue were significantly related to high Gleason score (p = 0.041), advanced clinical stage (p = 0.002), and a high percentage of positive biopsy cores (p = 0.001).

Conclusions: The results of this study indicate that high T levels in prostate tissue are related to high Gleason score, advanced clinical stage, and a high percentage of positive biopsy cores in patients with prostate cancer. T level in needle biopsy specimens may therefore be a useful prognostic factor in prostate cancer patients.

Keywords: Prostate cancer, Androgen, Testosterone, Dihydrotestosterone

Background

Prostate cancer is the most common internal cancer and the second most frequent cause of cancer-related deaths among men in the United States. Although the incidence of prostate cancer in Japan is lower than in the United States, it has been gradually increasing in recent years. The etiology of prostate cancer is unclear, but it is thought to be multifactorial, with genetic, dietary, and environmental causes. Although prostate cancer initially responds to androgen ablation therapy, most patients ultimately become hormone-refractory and show treatment failure.

The ability to predict prostate tumor behavior is important, because more intensive treatment is necessary to prevent the development of castration-resistant prostate cancer (CRPC). Pathological grade and clinical stage can strongly predict tumor aggressiveness, but no useful molecular markers have yet been identified. Several previous studies have reported blood and prostate tissue levels of testosterone (T) and dihydrotestosterone (DHT) in patients with prostate cancer, but these studies have involved small sample sizes, and several have measured the levels using radioimmunoassays (RIAs), which require a large amount of tissue (≥20 mg) for...
androgen measurement [1,2]. There have been few reports regarding the correlation between prostate cancer aggressiveness and androgen concentrations measured in smaller prostate tissue samples, such as those obtained by needle biopsy.

Advancements in liquid-chromatography tandem mass spectrometry (LC-MS/MS) methods mean that T and DHT levels can be measured in small tissue samples with high sensitivity and reliability [3-6]. LC-MS/MS can be used to measure androgen concentrations in tissue samples as small as those obtained by a single needle biopsy (approximately 3 mg), and the latest LC-MS/MS technique is more than 10 times as sensitive as the RIAs used in the past, especially in the lower concentration range [7].

Previous reports revealed that T levels were higher and DHT levels lower in prostate cancer tissues compared with tissues from patients with benign prostatic hyperplasia, although there is currently no consensus on androgen concentrations in prostate cancer tissues from men with different stages and with different pathological grades of disease [6,8-10]. Moreover, the relationship between tissue androgen concentrations and tumor behavior in prostate cancer is not clear. In the present study, we measured androgen (T and DHT) levels in blood and prostate tissues using LC-MS/MS and analyzed the correlations between these levels and prognostic factors in patients with prostate cancer.

Methods

Patients

A total of 359 patients with suspected prostate cancer underwent prostate needle biopsy for primary pathological diagnosis at major cancer treatment facilities in Japan between April 2000 and July 2003. Blood samples were also collected. All blood samples were taken between 09.00 h and 15.00 h to minimize the effect of daily T variations. Patients underwent a systematic needle biopsy. An additional needle biopsy sample was taken from the peripheral zone of the prostate as a chemical biopsy, for the simultaneous determination of T and DHT. The tissues were immediately frozen at −70°C until hormone analysis. Patients underwent biopsy for DHT. The concentrations of T and DHT were simultaneously determined in smaller prostate tissue samples, such as those obtained by needle biopsy.

The derivatization reagent was prepared as follows: 10 mg DAP, 20 mg MNBAn, and 25 mg of PA were dissolved in 1 mL of tetrahydrofuran and the mixture was agitated until it became cloudy or crystals appeared. The reagent solution was used after 3 min [5]. Serum prostate-specific antigen (PSA) levels were measured using a DPC Imrise third generation PSA assay kit.

LC-MS/MS

Serum hormone levels were determined by LC-MS/MS, as described by Yamashita et al. [5]. T and DHT levels in prostate tissue samples were measured using an API-5000 triple-stage quadrupole mass spectrometer (Applied

### Table 1 Patient characteristics

<table>
<thead>
<tr>
<th>No. patients</th>
<th>196</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age, years (mean, SD)</td>
<td>70.6 (7.334)</td>
</tr>
<tr>
<td>PSA, ng/mL (median, 95% CI)</td>
<td>11.5 (32.43–82.73)</td>
</tr>
<tr>
<td>Prostate volume, cm³ (median, 95% CI)</td>
<td>27.7 (29.83–34.91)</td>
</tr>
<tr>
<td>Gleason score ≤7, ≥8 (%)</td>
<td>130 (67.4), 63 (32.6)</td>
</tr>
<tr>
<td>Clinical stage ≤III, ≥IV (%)</td>
<td>166 (86.0), 27 (14.0)</td>
</tr>
<tr>
<td>% Positive core &lt;30%, ≥30% (%)</td>
<td>143 (74.1), 50 (25.9)</td>
</tr>
</tbody>
</table>

SD: Standard deviation; PSA: Prostate-specific antigen; CI: Confidence interval.
Tissue hormone analysis

Each frozen prostate tissue sample (1–3 mg, 5–13 mm) was weighed and then a piece of the sample was cut off with scissors. Purified water (0.5 mL) was added to the cut sample, which was then homogenized for 20 s using an Ultra-Turrax homogenizer with an ice-cooling bath, and then washed with 0.5 mL water. Ethanol (3 mL) was added and the mixture was allowed to stand at room temperature for 30 min. After dilution of the reaction mixture with 1% acetic acid solution (v/v, 1 mL) to stop the reaction, the resulting derivative was loaded onto a Bond Elut C18 cartridge pre-conditioned with methanol (6 mL) and purified water (6 mL). The cartridge was washed with purified water (1 mL) followed by 30% acetonitrile solution (v/v, 3 mL). T and DHT were subsequently eluted with 70% acetonitrile solution (v/v, 3 mL) and the solvent was removed using a centrifugation evaporator.

The residue was dissolved in 100 μL reagent mixture, prepared as described above. TEA (20 μL) was added and the resulting mixture was allowed to stand at room temperature for 30 min. After dilution of the reaction mixture with 1% acetic acid solution (v/v, 1 mL) to stop the reaction, the resulting derivative was loaded onto a Bond Elut C18 cartridge pre-conditioned with methanol and purified water. The cartridge was washed with purified water (1 mL) followed by 40% acetonitrile solution (v/v, 3 mL). The derivatives were then eluted with 80% acetonitrile solution (v/v, 3 mL). The solvent was evaporated to dryness using a centrifugation evaporator at 53–55°C, the residue was dissolved in 40% acetonitrile solution (v/v, 100 μL), and a 40-μL aliquot of the solution was subjected to LC-ESI-MS/MS.

Validation of analytical methods

The inter- and intra-assay accuracies and precisions of the LC-MS/MS were tested as described by Shibata et al. [11].

Cancerous and noncancerous regions in the prostate

Hormone-naive prostate cancer specimens were obtained from 10 patients by radical prostatectomy. Each prostate specimen was cut into two mirror-image 2.5-mm-thick fragments, each of which was cut into a further 90 sections. All 90 sections were used for the androgen assays and histopathology diagnosis, to analyze the relationships between tissue androgen concentrations with histopathological findings. There was no significant difference in tissue T or DHT levels between cancerous and noncancerous sections of individual prostate cancer sections (mean T levels, p = 0.796; mean DHT levels, p = 0.912). The tissues used for androgen concentration measurements in this study were not examined by a pathologist, because the whole of the samples were used for androgen measurement. Cancer diagnosis was based on other, simultaneously-obtained biopsy specimens. These results indicated that the results of androgen measurements were unaffected by the use of cancerous or noncancerous lesions.

Histopathology

All prostate biopsy samples were reviewed by a central pathologist. Pathological grading was performed according to the Gleason classification system. Histological diagnoses were made by the central pathologist blinded to the patients’ serum and prostate androgen measurements.

Statistical analysis

We compared tissue and blood androgen levels using Pearson’s correlation coefficient. If some factors were found to be correlated with other factors, the correlated factors were not analyzed simultaneously because of multicollinearity. We also analyzed the relationships between T and DHT levels in tissue and blood and prognostic factors including Gleason score, clinical stage, and % positive cores by multivariate analyses using a logistic regression model. All data were analyzed using IBM SPSS ver. 21 software. Each test was two-sided, and p values <0.05 were considered significant.

Results

Androgen levels in blood and prostate tissue obtained by needle biopsy

The androgen levels in prostate tissue and peripheral blood are shown in Table 2. Serum T levels were almost 10-fold higher than DHT levels in peripheral blood,
whereas DHT levels in prostate tissue were almost 10-fold higher than tissue T levels. These serum and tissue androgen levels were similar to those in other reports concerning prostate cancer patients.

Correlations between T and DHT levels in serum and prostate tissue are shown in Table 3. There were strong correlations between T and DHT blood levels (Pearson correlation coefficient: 0.76996) and we therefore did not analyze these two factors simultaneously, because of multicollinearity.

### Gleason score

The relationships between Gleason score and androgen levels and other clinical factors are shown in Table 4. Logistic regression identified high serum PSA (hazard ratio [HR]: 1.022, p = 0.001), low blood T levels (HR: 0.706, p = 0.012) and high tissue T levels (HR: 1.388, p = 0.041) as factors significantly associated with high Gleason score.

### Clinical stage

High serum PSA levels (HR: 1.009, p = 0.010) and high tissue T levels (HR: 1.713, p = 0.002) were significantly related to advanced clinical stage (Table 4).

### % Cancer-positive cores

High serum PSA (HR: 1.017, p = 0.005), low tissue DHT (HR: 0.878, p = 0.003) and high tissue T levels (HR: 2.432, p = 0.001) were significantly related to a high % positive cores (Table 4).

In summary, multivariate analysis demonstrated that: (1) these measurements could be conducted using the small amounts of tissue obtained by needle biopsy, rather than excised prostate tissues; and (2) androgen measurements could be carried out much more precisely than by RIA, by using LC-MS/MS. Nishiyama and colleagues used LC-MS/MS and reported that 47 patients with prostate cancer with Gleason scores of 7–10 had low DHT levels in the prostate, as shown by univariate analysis [9]. In the current study, we found that low tissue DHT levels were associated with a high % positive cores.

Previous studies of tissue androgen measurements have had several problems, including small sample sizes, tissue-handling issues [15], problems with methodological accuracy and concerns about confounding factors. The tissue content of DHT decreases rapidly within 2 h at 37°C, and the tissue samples used in the present study were therefore frozen immediately (at –70°C) until analysis. To ensure the quality and precision of our measurement methods, we used prostate tissues spiked with various amounts of T or DHT to determine the precision of the LC/MS-MS method, and performed multivariate analysisafter controlling for confounding factors.

Our results confirmed that high T levels in prostate tissue are related to high Gleason score, advanced clinical stage.
Table 4 Correlation between prognostic factors and androgen concentrations in blood and prostate tissue*

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Gleason score</th>
<th>Clinical stage</th>
<th>% positive cores</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>≥8 ≤7</td>
<td>≥IV ≤III</td>
<td>≥30% &lt;30%</td>
</tr>
<tr>
<td>No. of patients</td>
<td>63 130</td>
<td>27 166</td>
<td>50 143</td>
</tr>
<tr>
<td>Median age (SD), years</td>
<td>73.0 (7.1) 69.5 (7.2)</td>
<td>0.076 1.050 0.995–1.108</td>
<td>75.0 (7.3) 70.0 (7.2)</td>
</tr>
<tr>
<td>Median serum PSA (95% CI, pg/mL)</td>
<td>32.1 (69.2-216.2)</td>
<td>8.76 (12.3-20.2)</td>
<td>0.001 1.022 1.009–1.036</td>
</tr>
<tr>
<td>Median prostate volume, mL</td>
<td>27.0 (28.5-40.0)</td>
<td>27.9 (28.8-34.0)</td>
<td>0.387 0.991 0.970–1.012</td>
</tr>
<tr>
<td>Median concentration of T (95% CI) (blood) (/1000), pg/mL</td>
<td>3.11 (2.90-3.62)</td>
<td>3.64 (3.67-4.14)</td>
<td>0.012 0.706 0.538–0.926</td>
</tr>
<tr>
<td>Median concentration of DHT (95% CI) (tissue), pg/mg</td>
<td>5.31 (6.15-10.27)</td>
<td>7.31 (9.26-13.02)</td>
<td>0.222 0.974 0.935–1.016</td>
</tr>
<tr>
<td>Median concentration of T (95% CI) (tissue), pg/mg</td>
<td>0.90 (1.20-2.23)</td>
<td>0.46 (0.68-1.09)</td>
<td>0.041 1.388 1.013–1.900</td>
</tr>
</tbody>
</table>

*Multivariate analysis by logistic regression. SD: Standard deviation; HR: Hazard ratio; CI: Confidence interval.
and a high % positive cores. Low DHT levels in prostate tissue were also associated with a high % positive cores. Thomas et al. reported that prostate-tissue expression levels of 5alpha-reductase type 1 and type 2 were increased in high-grade, compared with low-grade prostate cancer, which might explain the relationships between high T and low DHT prostate tissue levels and poor prognostic factors [16], de Winter et al. reported that the intensity of androgen receptor immunostaining was decreased in more aggressive tumors [17], suggesting that androgen receptor heterogeneity might also offer an explanation of our findings.

Based on our results, we propose that prostate tissue T level measured by chemical biopsy may be a useful prognostic factor, in addition to Gleason score, clinical stage and % positive biopsy cores, for aiding the design of suitable therapeutic programs for prostate cancer. It is important that the tissue T level as a prognostic marker is determined simultaneously with the biopsy, to allow identification of the individual patient’s prognostic factors before making decisions about treatment options. Moreover, tissue T could help to subclassify Gleason 7 needle biopsies into a high-T and normal-T group, to distinguish between those patients with more aggressive disease and those with more indolent disease. This would be useful for selecting suitable patients for active surveillance.

Our results revealed that low serum T was associated with a Gleason score ≥8. In this regard, Garcia-Cruz et al. reported that low T was associated with high D’Amico classification, by multivariate analysis [18]. Hoffman et al. also reported that low free T indicated an increased risk of a biopsy Gleason score ≥8 (11% vs 0%, p = 0.025). Our results are compatible with the hypothesis that low serum androgen levels may influence poor-prognosis factors in prostate cancer [19], but no mechanism for this relationship has yet been established.

In our previous study, we first reported that CRPC may be predicted by prostate tissue T and DHT levels in a single biopsy specimen, obtained as a chemical biopsy, in patients undergoing cancer checkups, whether for prostate cancer or not [11]. Our present finding that high tissue T levels might be associated with poor prognostic factors may thus help predict prostate cancer aggressiveness. The correlation of high tissue T levels with poor prognostic parameters (Gleason score, advanced clinical stage, high % positive cores) detected in this study suggests that tissue T may have a prognostic role, though no prognostic information was available in the current study. Further studies with larger numbers of patients and longer follow-up are warranted to confirm the validity of high tissue T levels as a prognostic marker.

Conclusions
We conclude that high serum PSA levels and high tissue T levels in men with prostate cancer are significantly associated with indicators of poor prognosis, including advanced clinical stage, high Gleason score and % positive cores. Tissues T levels determined from biopsy specimens, in combination with pathological features, may be one of several useful diagnostic tools for predicting the prognosis and determining a suitable therapeutic program for patients with prostate cancer.

Additional file

Additional file 1: Ethics committee.

Abbreviations
LC-MS/MS: Liquid-chromatography tandem mass spectrometry; T: Testosterone; DHT: Dihydrotestosterone; PSA: Prostate-specific antigen; CRPC: Castration-resistant prostate cancer; RIA: Radioimmunoassay.

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

Authors’ contributions
YM and KZ designed the study. YM contributed to statistical analysis and wrote the manuscript. SM and KS supported the statistical analysis. HU, SU, YS, NM, TI, AM, YS, NN and HS contributed to collecting the tissue samples and clinical data. SH contributed to measurement of androgen concentrations. MH contributed to pathological examination. All authors have read and approved the final manuscript.

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