Magnetic resonance imaging-directed biopsy improves the prediction of prostate cancer aggressiveness compared with a 12-core transrectal ultrasound-guided prostate biopsy

JIE ZHANG¹, JIANJUN XIU¹, YIN DONG¹, MUWEN WANG², XUE HAN¹, YEJUN QIN³, ZHAOQIN HUANG¹, SHIFENG CAI¹, XIANSHUN YUAN¹ and QINGWEI LIU¹

¹Department of Radiology, ²Minimally Invasive Urology Center and ³Department of Pathology, Provincial Hospital, Shandong University, Jinan 250021, P.R. China

Received October 22, 2013; Accepted February 17, 2014

DOI: 10.3892/mmr.2014.1994

Abstract. The Gleason grading system is a fundamental indicator of the aggressive nature of prostate cancer (PCa). Diffusion-weighted imaging (DWI) and magnetic resonance spectroscopy (MRS) are methods for the assessment of PCa aggressiveness. The present study was designed to prospectively investigate whether transrectal ultrasound (TRUS)-guided MR imaging (MRI)-directed biopsies (TRUS-MR-Dbs) improve the prediction of PCa aggressiveness in comparison with 12-core TRUS-guided biopsies (TRUS-Gbs). A total of 518 patients underwent pre-biopsy multi-parametric MRI to identify the clinically suspicious PCa regions. TRUS-MR-Dbs were performed on patients with suspected PCa by MRI in addition to TRUS-Gbs. Only patients who underwent radical prostatectomy (RP) were included in the comparative analysis. TRUS-biopsy was directed to those areas within suspicious regions with a minimum apparent diffusion coefficient obtained by DWI or with a maximum (choline + creatine)/citrate ratio obtained by MRS. The highest Gleason grades (HGGs) and the Gleason scores (GSs) of specimens were identified. The biopsies and RP results were evaluated using a McNemar test or χ² analyses using Fisher’ exact tests. MRI results were positive in 254 (49.0%) of the 518 patients. TRUS-MR-Dbs detected 165/254 (65.0%) cancer cases and TRUS-Gbs detected 190/518 (36.7%) cancer cases. Forty patients underwent RP. The TRUS-MR-Dbs method demonstrated a higher concordance rate (CR) with RP (89.6%) than TRUS-Gbs (72.9%, P=0.008) for the overall HGG. The CRs with RP for TRUS-MR-Dbs vs. those for TRUS-Gbs were 100 vs. 85.7% (P=0.5), 87.5 vs. 68.8% (P=0.031) and 50 vs. 50% (P=1) for HGG3, HGG4 and HGG5, respectively. The HGG CRs with RP for DWI-directed biopsies (DWI-Dbs) vs. MRS-directed biopsies (MRS-Dbs) were 77.1 vs. 50.0% (P=0.015) for the overall tumors, 80.0 vs. 40.0% (P=0.003) for peripheral zone tumors and 69.2 vs. 76.9% (P=1) for transition zone tumors. A total of 37 (77.1%) and 25 (52.1%; P=0.007) tumors were assigned accurate GS for TRUS-MR-Dbs and TRUS-Gbs, respectively. The results revealed that TRUS-MR-Dbs improved the prediction of PCa aggressiveness and that DWI-Dbs had a superior performance when compared with MRS-Dbs in the peripheral zone.

Introduction

The precise determination of prostate cancer (PCa) aggressiveness facilitates the application of more personalized treatment regimens and improves the prediction of the prognosis for patients. The Gleason grade may be used to indicate the pathological characteristics of PCa. The Gleason score (GS) is a fundamental biological manifestation of the aggressiveness and prognosis of PCa (1,2). The 12-core extended systematic transrectal ultrasound (TRUS)-guided biopsy (TRUS-Gb) is currently an accepted standard method to detect tumors and identify the tumors’ Gleason grades. This contributed to the detection of low-volume, low-risk tumors, but the TRUS-Gb-determined GS is often increased in numerous patients receiving radical prostatectomy (RP) (3,4).

Multi-parametric magnetic resonance imaging (MRI) methods, including T2-weighted imaging (T2WI), diffusion-weighted imaging (DWI), MR spectroscopy (MRS) and dynamic contrast-enhanced (DCE) imaging, are important in the accurate detection of PCa (5,6). MRI-targeted biopsy is a potential alternative to improve the detection of PCa (7-10). DWI evaluates the random Brownian motion properties of water molecules in tissues, which are sensitive to water diffusion restriction. MRS is a non-invasive imaging modality to obtain metabolic information about tumors. DWI and MRS have demonstrated promising results in assessing PCa aggressiveness (11-13).
The purpose of the present study was to investigate whether TRUS-guided MRI-directed biopsies (TRUS-MR-Dbs), including DWI-directed biopsies (DWI-Dbs) and MRS-directed biopsies (MRS-Dbs), may improve the prediction of PCA aggressiveness in patients when compared with the 12-core TRUS-Gbs.

Materials and methods

Ethics statement. The Ethical Committee of the Provincial Hospital Affiliated to Shandong University (Shandong, China) approved this prospective study and written informed consent was obtained from all of the participants.

Patients. The general practice of urologists in China and preference of the majority of patients is to undergo MRI examination prior to biopsy rather than following biopsy. As a routine examination, patients with clinically suspicious PCa had been excluded. The Ethical Committee of the Provincial Hospital approved this prospective study and written informed consent was obtained from all of the participants.

MRI. All MRI examinations were performed on a 3.0-T system (Magnetom verio; SIEMENS, Munich, Bavaria, Germany) with integrated eight-channel pelvic phased-array surface coils and spine coils for signal reception. All patients were imaged in a supine, head first position. The sequences with large FOVs were used to observe nodal metastases and distant spread. Coronal and/or sagittal enhanced T1-weighted imaging as well as volume interpolated body examination with large FOVs were optional and are not listed in Table I. T2WI was performed with a turbo spin-echo technique. DWI was performed using a single-shot echo-planar imaging technique. Apparent diffusion coefficient (ADC) maps were automatically generated from the DWI by the scanner. The MRS was performed using 3D chemical shifting imaging techniques based on a point-resolved spectroscopic sequence with sufficient lipid and water suppression. Eight saturation bands were used to minimize the contamination from adjacent structures of the prostate. Prior to approval for evaluation, a spectroscopist validated the spectra by examining them with regard to the correct positions, signal-to-noise ratio (SNR) >5:1, full width at half maximum (FWHM) ≤15 Hz, a relatively steady baseline and the absence of lipid signals.

Table I. MRI procedures and the corresponding parameters.

<table>
<thead>
<tr>
<th>Sequences</th>
<th>TR (ms)</th>
<th>TE (ms)</th>
<th>ST (mm)</th>
<th>Average</th>
<th>FOV (cm²)</th>
<th>Matrix</th>
</tr>
</thead>
<tbody>
<tr>
<td>Axial T2WI (TSE)</td>
<td>3110.00</td>
<td>101.00</td>
<td>3</td>
<td>2</td>
<td>20.0x20.0</td>
<td>320x256</td>
</tr>
<tr>
<td>Coronal T2WI (TSE)</td>
<td>2950.00</td>
<td>96.00</td>
<td>3</td>
<td>2</td>
<td>20.0x20.0</td>
<td>320x256</td>
</tr>
<tr>
<td>Sagittal T2WI (TSE)</td>
<td>3410.00</td>
<td>102.00</td>
<td>3</td>
<td>2</td>
<td>20.0x20.0</td>
<td>320x256</td>
</tr>
<tr>
<td>MRS³ (CSI-PRESS)</td>
<td>750.00</td>
<td>145.00</td>
<td>-</td>
<td>6</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>DWI² (EPI)</td>
<td>6200.00</td>
<td>93.00</td>
<td>3</td>
<td>6</td>
<td>20.0x20.0</td>
<td>160x120</td>
</tr>
<tr>
<td>Axial T1WI (TSE)</td>
<td>467.00-645.00</td>
<td>9.80</td>
<td>4</td>
<td>1</td>
<td>20.0x20.0</td>
<td>512x384</td>
</tr>
<tr>
<td>Axial T2WI (TSE)</td>
<td>4070.00</td>
<td>93.00</td>
<td>5</td>
<td>1</td>
<td>38.0x28.5</td>
<td>320x168</td>
</tr>
<tr>
<td>Axial T1WI (VIBE)⁵</td>
<td>3.90</td>
<td>1.40</td>
<td>3</td>
<td>1</td>
<td>38.0x30.8</td>
<td>320x182</td>
</tr>
<tr>
<td>Axial T1WI-DCE (VIBE)⁵</td>
<td>5.21</td>
<td>1.80</td>
<td>3</td>
<td>1</td>
<td>20.0x20.0</td>
<td>224x160</td>
</tr>
</tbody>
</table>

³resolution was 7x7x7 mm³; b-values = 0 and 800 sec/mm²; the sequence was performed prior to and following the injection of the contrast agent. TR, repetition time; TE, echo time; ST, slice thickness; FOV, field of view; TSE, turbo spin-echo sequence; EPI, single-shot echo-planar imaging; CSI-PRESS, 3D chemical shifting imaging techniques based on point-resolved spectroscopic sequence; DCE, dynamic contrast enhancement imaging; VIBE, volume interpolated body examination; MRI, magnetic resonance imaging; T1WI, T1-weighted imaging; T2WI, T2-weighted imaging; DWI, diffusion-weighted imaging; MRS, magnetic resonance spectroscopy.

Two senior radiologists determined the PCa suspicious regions in consensus using the combined information of multi-parametric MRI. A lesion fulfilling two or more of the following criteria was regarded as a PCa suspicious region: i) An area with homogeneous low-signal-intensity and mass effect in the T2W images; ii) focal hyperintensity in diffusion-weighted images and corresponding hypointensity in ADC maps; iii) a normal choline (Cho) + creatine (Cr)/citrate (Cit) ratio (CC/C) = 0.22±0.12 in the peripheral zone (PZ) and (CC/C) = 0.34±0.14 in the transition zone (TZ) (14) and CC/C >0.34 [the mean ± standard deviation] in PZ and CC/C >0.48 in TZ and iv) a lesion that enhances rapidly and is washout of contrast agent in delay phase.
Biopsy and histopathological analysis. The median interval time from MRI examination to biopsy was three days (range, 1-12 days). Considering the areas with the minimum ADC or with the maximum CC/C within suspicious regions may represent the most abnormal part with the highest cellularity or metabolism (15,16). These areas were used to target the biopsies to represent the highest grade of the tumor (Figs. 1 and 2). The patients with suspected PCa on MR images under-
went TRUS-MR-Dbs besides TRUS-Gbs (12-core) using an ultrasound device (DC-7; Mindray Medical International Limited, Shenzhen, China). The biopsy was performed by an urologist and a radiologist. The radiologist was also involved in the interpretation of the MR images. The steps of biopsy were as follows: In step 1, the 12-core systematic TRUS-Gbs were performed on all patients. In step 2, the area with minimum ADC in the suspicious regions was visually matched with the corresponding location identified by TRUS. One- to two-core biopsy was performed in this area while monitoring by TRUS. This step is known as DWI-Dbs. In step 3, the area with the maximum CC/C in the suspicious regions was visually matched with the corresponding location identified using TRUS. One- to two-core biopsy was performed in this area while monitoring by TRUS. This step is known as MRS-Dbs. In step 4, one- to four-core biopsy was performed under the TRUS in the suspicious regions in MR images in addition to the biopsies performed in steps 2 and 3.

If there were no suspicious regions of PCa on MRI, only step 1 was performed. If the minimum ADC and the maximum CC/C appeared in the same site, steps 2 or 3 were omitted. The combination of steps 2-4 is referred to as TRUS-MR-Dbs in the present study. The highest Gleason grade (HGG) for TRUS-MR-Dbs was obtained from step 2 and/or step 3. The Gleason score (GS) for TRUS-MR-Dbs was obtained from step 2, step 3 and step 4. The HGG and GS for TRUS-Gbs were obtained from step 1.

TRUS was able to provide dynamic anatomy images of arbitrary sections. The majority of PCa cases were present as hypoechoic masses. However, the sonographic findings of PCa were non-specific. Numerous tumors were iso-echoic on TRUS. To achieve a good spatial consistency between ultrasound scans and MRI sections in performing TRUS-MR-Dbs (step 2, 3 and 4), certain criteria were required to be met for the biopsy: The gland morphology determined using ultrasound axial and sagittal scans being similar to that obtained with the axial and sagittal T2W images, in regard to specific anatomic structures (i.e. seminal vesicles, verumontanum, urethra and a number of hyperplasia-nodules) as landmarks. The biopsy cores were labeled to specify the location of the biopsy and marked on the T2W images at the corresponding sites.

Collectively, 12-18 cores were obtained from each patient. A median of four cores (range, two-six), which contained a median of two cores (range, one-three) in the most abnormal parts, were obtained from the suspicious regions for PCa in each patient.

The sites of suspicious PCa and their number were recorded on MRI for each patient. TRUS-MR-Dbs cores corresponding to each suspicious tumor on MRI were also recorded. The sites of 12 cores of TRUS-Gbs were relatively fixed. In the present study, only the patients who underwent RP were included to predict the aggressiveness of the PCa. The number of tumors was confirmed by RP pathology. For the patients who underwent RP, the urologist, the pathologist and the radiologist confirmed in consensus which cores of TRUS-Gbs and which cores of TRUS-MR-Dbs corresponded with each tumor using the biopsy records, the biopsy/RP pathology and the MR images as references. The biopsy cores corresponding to each tumor in a patient were then confirmed by the two biopsy methods. Only the tumors visible on MR images were included.

The median interval time from biopsy to RP was nine days (2-33 days). Pathological sections of prostate from RP were matched with MR sections based on the level sextant locations by a pathologist and a radiologist. For cancerous biopsy cores and RP specimens, the GS of each tumor were determined by a pathologist with 15 years of experience, by the sum of the primary and secondary Gleason grades. The corresponding HGG of the specimens from the most aberrant regions was identified. The tumor volume was calculated by using the ellipsoid formula (0.52 x length x width x height).

### Statistical analysis

Cross-tabulation analysis was used to describe the biopsy and RP findings (including HGG and GS). For both TRUS-Gbs and TRUS-MR-Dbs, the concordance rates (CRs) with RP were determined for HGG and GS. McNemar tests or $\chi^2$ analyses with Fisher's exact tests were performed to determine the differences for CRs with RP between TRUS-MR-Dbs and TRUS-Gbs as well as between DWI-Dbs and MRS-Dbs. The independent samples t-test was used to determine the difference between the volumes of tumors with accurate GS and of tumors with lower GS for TRUS-MR-Dbs. The correlation between tumor volume and HGG was evaluated using Pearson's correlation. For all statistical analyses, $P<0.05$ was considered to indicate a statistically significant difference. Statistical analysis was performed with SPSS for Windows, version 17.0 (SPSS, Inc., Chicago, IL, USA).
Results

Detection of cancer by TRUS-Gb. MRI results were positive in 254 (49.0%) of the 518 patients. TRUS-MR-Db alone detected 165/254 (65.0%) of cancer cases. TRUS-Gb alone detected 190/518 (36.7%) of cancer cases. The overall number of patients with cancer was 196. The median age of the 196 patients was 73 years (range, 51-87 years). The 165 cancer cases detected by TRUS-MR-Db included 159 of the 190 cancer cases, which were detected by TRUS-Gb, and six that were not able to be detected by TRUS-Gb. These six cancer cases had a HGG ≥ 3. There were 31 cancer cases that were not detectable by TRUS-MR-Db, but by TRUS-Gb. Among these 31 cancer cases, 24 had a HGG of ≤ 3 and 26 were <0.5 cm in length.

RP findings. Out of the 165 patients, only 40 patients underwent RP and the quality of MR images and MRS of these patients matched the criteria. The clinical results of 40 patients with 48 tumors (8 patients, 2 tumors/patient; 32 patients, 1 tumor/patient) are presented in Table II. Tables III and IV summarize the biopsy and RP findings of these patients. TRUS-MR-Db demonstrated a higher CR with RP for overall HGG, 89.6% (43/48). For TRUS-Gb, it was 72.9% (35/48) (P=0.008). The CRs with RP for TRUS-MR-Db vs. those for TRUS-Gbs were 100% (14/14) vs. 85.7% (12/14; P=0.5) for tumors with HGG of 3 (HGG3); 87.5% (28/32) vs. 68.8% (22/32; P=0.031) for tumors with HGG4 and 50% (1/2) vs. 50% (1/2; P=1) for tumors with HGG5 (Fig. 3A). For biopsies with low grade (HGG3), the positive predictive value (PPV) for TRUS-MR-Db representing a true low grade was 77.8% (12/14), whereas for TRUS-Gb it was 54.5% (12/22; P=0.125). Undergrading of tumors compared with RP was present at 10.4% (5/48) for TRUS-MR-Db and 27.1% (13/48) for TRUS-Gb (P=0.008). Undergrading of tumors to HGG<4, which had been graded as HGG4 or HGG5 by RP, occurred at 29.4% (10/34) for TRUS-Gbs and 11.8% (4/34) for TRUS-MR-Db (P=0.031). No overgrading was observed for the biopsies.

The HGG CRs with RP determined by DWI-Db vs. MRS-Db were 77.1% (37/48) vs. 50 (24) (0.015) for overall HGG, 89.6% (43/48) vs. 72.9 (35) (P=0.008) for tumors with HGG of 3 (HGG3); 87.5% (28/32) vs. 68.8% (22/32; P=0.031) for tumors with HGG4 and 50% (1/2) vs. 50% (1/2; P=1) for tumors with HGG5 (Fig. 3A). For biopsies with low grade (HGG3), the positive predictive value (PPV) for TRUS-MR-Db representing a true low grade was 77.8% (12/14), whereas for TRUS-Gb it was 54.5% (12/22; P=0.125). Undergrading of tumors compared with RP was present at 10.4% (5/48) for TRUS-MR-Db and 27.1% (13/48) for TRUS-Gb (P=0.008). Undergrading of tumors to HGG<4, which had been graded as HGG4 or HGG5 by RP, occurred at 29.4% (10/34) for TRUS-Gbs and 11.8% (4/34) for TRUS-MR-Db (P=0.031). No overgrading was observed for the biopsies.

GS findings. A total of 37 (77.1%) and 25 (52.1%; P=0.004) tumors were assigned the same GS as with RP by
By TRUS-MR-Dbs and TRUS-Gbs, respectively (Fig. 3B). By
TRUS-MR-Dbs and TRUS-Gbs, 9 (18.8%) and 23 (47.9%;
P<0.001) tumors were assigned a lower GS than that assigned
by RP, respectively. Two (4.5%) tumors were assigned a
higher GS than that assigned by RP for TRUS-MR-Dbs. The
GSs determined by biopsies and RP are presented in Table V.
The volume of tumors assigned the same GS as RP for
TRUS-MR-Dbs (5.3±4.0 cm³) was significantly smaller than
that of the tumors assigned lower GS (9.1±5.3 cm³; P=0.02).
The tumor volume demonstrated a positive correlation with
the HGG (r=0.396; P=0.005).

Discussion
In the present prospective study, both TRUS-MR-Dbs and
TRUS-Gbs were performed on the same patient, and assessed

Table V. Cross tabulation for biopsies and radical prostatectomy results based on GS grouping.

<table>
<thead>
<tr>
<th>Biopsy type</th>
<th>Biopsy result</th>
<th>2+3</th>
<th>3+2</th>
<th>3+3</th>
<th>3+4</th>
<th>4+3</th>
<th>4+4</th>
<th>4+5</th>
<th>PPV, % (n)</th>
</tr>
</thead>
<tbody>
<tr>
<td>TRUS-Gb</td>
<td>GS0</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0.0(0/2)</td>
</tr>
<tr>
<td></td>
<td>2+3</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>50.0(1/2)</td>
</tr>
<tr>
<td></td>
<td>3+2</td>
<td>0</td>
<td>3</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>60.0(3/5)</td>
</tr>
<tr>
<td></td>
<td>3+3</td>
<td>0</td>
<td>0</td>
<td>5</td>
<td>6</td>
<td>4</td>
<td>0</td>
<td>0</td>
<td>33.3(5/15)</td>
</tr>
<tr>
<td></td>
<td>3+4</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>7</td>
<td>6</td>
<td>0</td>
<td>0</td>
<td>53.8(7/13)</td>
</tr>
<tr>
<td></td>
<td>4+3</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>6</td>
<td>1</td>
<td>0</td>
<td>1</td>
<td>85.7(6/7)</td>
</tr>
<tr>
<td></td>
<td>4+4</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>2</td>
<td>1</td>
<td>66.7(2/3)</td>
</tr>
<tr>
<td></td>
<td>4+5</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>100.0(1/1)</td>
</tr>
<tr>
<td>CR, % (n)</td>
<td>100(1/1)</td>
<td>60(3/5)</td>
<td>62.5(5/8)</td>
<td>53.8(7/13)</td>
<td>37.5(6/16)</td>
<td>66.7(2/3)</td>
<td>50(1/2)</td>
<td>52.1(25/48)</td>
<td></td>
</tr>
<tr>
<td>TRUS-MR-Db</td>
<td>2+3</td>
<td>1</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>50.0(1/2)</td>
</tr>
<tr>
<td></td>
<td>3+2</td>
<td>0</td>
<td>4</td>
<td>1</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>80.0(4/5)</td>
</tr>
<tr>
<td></td>
<td>3+3</td>
<td>0</td>
<td>0</td>
<td>7</td>
<td>2</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>63.6(7/11)</td>
</tr>
<tr>
<td></td>
<td>3+4</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>10</td>
<td>2</td>
<td>0</td>
<td>0</td>
<td>83.3(10/12)</td>
</tr>
<tr>
<td></td>
<td>4+3</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>11</td>
<td>0</td>
<td>0</td>
<td>91.7(11/12)</td>
</tr>
<tr>
<td></td>
<td>4+4</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>3</td>
<td>1</td>
<td>60.0(3/5)</td>
</tr>
<tr>
<td></td>
<td>4+5</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>100.0(1/1)</td>
</tr>
<tr>
<td>CR, % (n)</td>
<td>100(1/1)</td>
<td>80(4/5)</td>
<td>87.5(7/8)</td>
<td>76.9(10/13)</td>
<td>68.8(11/16)</td>
<td>100(3/3)</td>
<td>50(1/2)</td>
<td>77.1(37/48)</td>
<td></td>
</tr>
</tbody>
</table>

GS, Gleason score; GS0, not detected; TRUS-Gb, 12-core transrectal ultrasound guided biopsy; TRUS-MR-Db, transrectal ultrasound-guided MR imaging-directed biopsy; CR, concordance rate; PPV, positive predictive values.

Figure 3. Concordance rates. (A) Concordance rates according to HGG categorization; 12-core TRUS-Gb vs. TRUS-MR-Db. (B) Concordance rates according to Gleason score categorization; 12-core TRUS-Gb vs. TRUS-MR-Db. HGG, highest Gleason grade; TRUS-Gb, 12-core transrectal ultrasound guided biopsy; TRUS-MR-Db, transrectal ultrasound-guided magnetic resonance imaging-directed biopsy.
in terms of how clinical factors for different patients (i.e., age, PSA, prostate weight) affect the biopsy Gleason grade CRs with RP (17). The most abnormal ADC or CC/C areas for PCa following multi-parametric imaging were used to target the biopsies. These areas were expected to have corresponding tumors with the highest grading (12,18). Therefore, HGG CRs for biopsies with RP were mainly investigated in the present study. To the best of our knowledge, this may be the first prospective report on the use of the combination of DWI and MRS to direct TRUS-guided biopsy to obtain PCa specimens that may be more representative of the true RP Gleason grade.

Although TRUS-Gb increased the number of detected cancer cases, the majority or those that were not detected by TRUS-MR-Db exhibited an HGG ≤3 or a length <0.5 cm. TRUS-MR-Db used fewer biopsy cores (a median value of four cores) in a fewer number of patients (49% of patients) and identified a lower number of low grade or microfocal tumors. These results were consistent with the findings of previous studies and reviews (8,19,20), where it was suggested that TRUS-MR-Db may avoid the unnecessary diagnosis of insignificant PCa.

The Gleason grades as determined by TRUS-MR-Dbs demonstrated a higher CR with RP compared with TRUS-Gbs. Biopsy tends to underestimate tumor grades, which may result in the undertreatment of patients. For example, patients with high grades (HGG4 and HGG5) may be incorrectly receiving a treatment that is also used for patients with low grades (HGG≤4). The rate of underestimation compared with RP for TRUS-MR-Db was significantly lower than that for TRUS-Gb (P=0.031). These results were consistent with results of previous retrospective studies on the ability of DWI and MRS to assess tumor aggressiveness and serve as biomarkers to improve pretreatment prediction of HGG (12,13,21).

However, TRUS-MR-Gbs only improved the prediction rates for tumors with HGG4 but not for tumors with HGG3 and 5. For tumors with HGG3, the GS of tumors visible on MR images was 3+3, followed by 3+2 in the present study, which demonstrated that the structure/grade of low-grade tumors was relatively homogeneous or mainly of grade 3. Furthermore, the results of the present study revealed that a small tumor volume was associated with a low grade. Therefore, as long as the tumor was detected by 12-core TRUS-Gbs, the probability of detection of a component with grade 3 may be greater. San Francisco et al (22) have reported that tumors with low Gleason grade detected by extended biopsies (≥10 cores) had significantly higher concordance (88%) with the prostatectomy Gleason grade. In addition, the 12-core biopsy was performed under ultrasound guidance in the present study. The intensity of the relatively small low-grade tumors visible on MR images was relatively homogeneous, which may correspond to a homogeneous structure/grade in the low-grade tumors in the present study. For HGG5, the number of tumors was markedly low (2), which may be not sufficient for a reliable analysis. In addition, the intensity of high grade tumors was often heterogeneous, which may be associated with tumor necrosis. In addition, high grade tumors were often large, which increased the difficulty of the spatial consistency between MR images and ultrasound scans. Therefore, TRUS-MR-Dbs did not improve the prediction rate for HGG5.

DWI-Dbs performance was significantly superior compared with MRS-Dbs in assessing the HGG of PCa (P=0.015). TRUS-guided DWI-Dbs and MRS-Dbs improved the accuracy in the detection of PCa (19,23,24). However, DWI-Db and TRUS-Gb CRs with RP for HGGs were not significantly different (P=0.727). CRs with RP for MRS-Dbs were lower than for TRUS-Gb (P=0.013; Table IV). The negative correlation between ADC and Gleason grade and the positive correlation between CC/C and Gleason grade have been reported in previous studies (13,18,25). Despite this, it is possible that the minimum ADC or maximum CC/C of a tumor may not fully represent the highest grade of a tumor, particularly for CC/C. For only 50% of tumors, the HGGs were accurately predicted with the maximum CC/C. However, TRUS-MR-Dbs (the combination of DWI-Dbs and MRS-Dbs) may improve the CRs with RP compared with TRUS-Gbs. A previous prospective study reported that DWI targeted MR-guided biopsy significantly improved the pretreatment assessment of PCa aggressiveness compared with 10-core TRUS-guided biopsy (26). The possible reasons for the different results are as follows: (i) The location accuracy for TRUS-MR-Db is lower than that for MR-guided biopsy and (ii) the 12-core biopsy further improves the CRs with RP compared with 10-core biopsy (22). Despite this, the combination of DWI- and MRS-Dbs had an improved performance as compared with TRUS-Gb. DWI-Db has a superior performance to MRS-Dbs in detecting HGG in predicting HGG for PZ tumors (P=0.003), but was equal to MRS-Dbs for TZ tumors (P=1). It may be associated with relatively strict requirements for MRS, including uniformity of a static magnetic field and stability of radiofrequency. Although eight saturation bands were used to minimize the contamination from adjacent structures of the prostate, a number of artifacts may have remained. The PZ adjacent to the rectum and pelvic fat is more susceptible to rectum gas, rectum peristalsis and lipid contamination. Voxels with poor quality MRS or artifacts were excluded. The majority of the excluded voxels were in PZ tumors. A number of the excluded voxels may have had the highest grade and certain included voxels may have contained hidden artifacts. Although DWI may also be affected by these artifacts, the impact on DWI was far less than that on MRS. A number of the artifacts of DWI may have been corrected or compensated in a better way. A retrospective study by Kobus et al (11) also reported that DWI assesses the PCa aggressiveness better in PZ.

The GS of PCa has a dominant role in the evaluation of tumor aggressiveness. The GS CRs with RP determined by TRUS-MR-Dbs and TRUS-Gbs were also compared. The GS determined by TRUS-MR-Dbs demonstrated a higher CR (P=0.004) and a lower underestimated rate (P<0.001) with RP than that determined by TRUS-Gbs; however, TRUS-MR-Dbs overestimated two tumors. The mean volume of tumors assigned accurate GS was smaller than that of underestimated tumors.

There were several limitations in the present study, including the relatively low number of tumors examined, particularly the number of tumors with HGG5 (only two), which may reduce the validity of the results. In China, the incidence of PCa is lower than in Western countries. Although the incidence of PCa has significantly increased in recent years, the PSA...
screening is not commonly applied and the majority of patients are identified as having high grade PCAs when diagnosed (27) and have lost the opportunity for RP. In China, patients with PCAs are relatively old (median age >70 years) and therefore, patients tend to select a therapy associated with reduced injury, such as endocrine therapy instead of RP. Almost all patients with HGG5 had lost the opportunity of RP in the Provincial Hospital (Shandong, China), which may be the reason for the low number of HGG5 tumors. The CRs with RP of each GS group of biopsies were calculated, but statistical comparisons for the corresponding GS groups between TRUS-MR-Dbs and TRUS-Gbs were not performed due to the low number of each GS group tumors.

A second limitation was the spatial consistency between ultrasound scans and MRI sections. Although measures were taken to solve this problem, it was not easy to achieve the exact spatial consistency. An MRI-ultrasound fusion platform may improve the detection of tumors due to its relatively accurate spatial consistency (19,28,29). To the best of our knowledge, this platform has not been used to predict tumor aggressiveness thus far. MRI-guided biopsy may locate the tumor more accurately and has been used to improve the prediction of PCA aggressiveness (26). However performing target biopsy under MRI-guidance is expensive, time-consuming and not widely available. TRUS-guided biopsies with the visual facilitations of MRI (DWI or MRS in the present study) have been prospectively demonstrated to contribute to the detection of PCAs (7,24), and is currently the most widely used diagnostic strategy. The present study used the same biopsy method (TRUS-MR-Db) to detect the most abnormal part of PCa. Although the spatial consistency with TRUS-MR-Db between the ultrasound scans and MRI sections may be poorer than with the MRI-ultrasound fusion platform, TRUS-MR-Db was satisfactory to a certain extent. Therefore, TRUS-MR-Dbs (combination of DWI- and MRS-Dbs) demonstrated higher HGG and GS CRs for overall tumors with RP compared with TRUS-Gbs.

In conclusion, DWI- and MRS-directed biopsies at the most abnormal (lowest ADC or highest CC/C) sites may be necessary in the pretreatment and prediction of tumor aggressiveness in PCAs. DWI-Dbs are more effective than MRS-Dbs, particularly for tumors in the PZ.

Acknowledgements

The present study was supported by the Shandong Province Science and Technology Development Plan (nos. 2012GSF11820 and 2012YD18053). The authors are grateful to the language services agent, Beijing chunfenglv, for the references (nos. 2012GSF11820 and 2012YD18053). The authors are thankful to the language services agent, Beijing chunfenglv, for their help.

References


