Tissue Microstructure Is Linked to MRI Parameters and Metabolite Levels in Prostate Cancer

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Introduction: Magnetic resonance imaging (MRI) can portray spatial variations in tumor heterogeneity, architecture, and its microenvironment in a non-destructive way. The objective of this study was to assess the relationship between MRI parameters measured on patients in vivo, individual metabolites measured in prostatectomy tissue ex vivo, and quantitative histopathology.

Materials and methods: Fresh frozen tissue samples (n = 53 from 15 patients) were extracted from transversal prostate slices and linked to in vivo MR images, allowing spatially matching of ex vivo measured metabolites with in vivo MR parameters. Color-based segmentation of cryosections of each tissue sample was used to identify luminal space, stroma, and nuclei.

Results: Cancer samples have significantly lower area percentage of lumen and higher area percentage of nuclei than non-cancer samples (p ≤ 0.001). Apparent diffusion coefficient is significantly correlated with percentage area of lumen (ρ = 0.6, p < 0.001) and percentage area of nuclei (p = −0.35, p = 0.01). There is a positive correlation (ρ = 0.31, p = 0.053) between citrate and percentage area of lumen. Choline is negatively correlated with lumen (ρ = −0.38, p = 0.02) and positively correlated with percentage area of nuclei (ρ = 0.38, p = 0.02).

Conclusion: Microstructures that are observed by histopathology are linked to MR characteristics and metabolite levels observed in prostate cancer.

Keywords: ADC, magnetic resonance imaging, citrate, choline, HR-MAS MRS, color-based segmentation

INTRODUCTION

Magnetic resonance imaging (MRI) plays an important role in the diagnostic work-up of prostate cancer patients (1). MRI can portray spatial variations in tumor heterogeneity, architecture, and its microenvironment in a non-destructive way. Metabolic and morphologic changes in prostate cancer tissue lead to changes in MRI and MR spectroscopy (MRS) parameters. In cancer areas of the prostate’s peripheral zone, T2 intensity and apparent diffusion coefficient (ADC) are reduced,
while the choline and creatine-to-citrate ratio is increased compared to non-cancer areas (2, 3). These cancer-related changes in MR imaging parameters may be caused by a combination of increased cellularity, reduced luminal space, and altered metabolism.

Attempts have been made to elucidate the relationship between MR visible metabolites and MR imaging parameters. A negative correlation between ADC and the metabolite ratio choline plus creatine-to-citrate has been demonstrated (4). Due to low spectral resolution in vivo, correlation between MRI parameters and individual metabolites (such as choline-containing compounds and citrate) has not yet been investigated. Ex vivo high-resolution spectroscopy data from magic angle spinning (HR-MAS) MRS from prostate cancer can be linked to in vivo MR parameters as previously described (5), and individual metabolites can therefore be correlated with MRI parameters.

In the prostate, healthy peripheral zone has heterogeneous tissue architecture and consists primarily of glandular lumen lined with secretory epithelium embedded within a stromal matrix (6). Healthy prostate tissue is characterized by high levels of citrate since the glandular secretory epithelial cells have the ability to produce, accumulate, and secrete citrate (7). The water content is low in stroma, but high in luminal space resulting in a relatively long $T_2$ and unrestricted water diffusion. As a result, healthy peripheral zone shows high signal intensity in $T_2$-weighted images and on ADC maps, while $T_2$ intensity and ADC are reduced in cancer (8–14). However, it has been demonstrated by Bourne et al. that the secretory epithelia within healthy peripheral zone represent a compartment of highly restricted water diffusion (6) and that healthy peripheral zone in general displays two $T_2$ components, one liquid-like component with long $T_2$ time originating from the luminal space and one component with a short $T_2$ time originating from stromal and epithelial tissues (15). Further, significant correlations between imaging parameters and histological features, such as luminal space, cell density, percentage of nuclei, and cytoplasm, have been demonstrated (16–18).

The objective of this study was to assess the relationship between MRI parameters ($T_2$ intensity and ADC) measured on patients in vivo, individual metabolites measured in prostatectomy tissue ex vivo and quantitative histopathological features (percentage nuclei, stroma, and luminal space). An overview of these relationships could give a better insight into the origin of the observed MRI and MRS signals and contribute to better understanding of the similarities and differences of these parameters.

**MATERIALS AND METHODS**

**Patients and Tissue Samples**

The Regional Committee for Medical Research Ethics approved the study, and patients gave informed written consent to participate. Fresh frozen tissue samples ($n = 53$ from 15 patients, median 3 (range 2–6) per patient) were extracted from transversal prostate slices and linked to in vivo MR parameters as previously described (5, 19). In short, a full transversal, fresh tissue slice (2 mm thick) was resected from the middle of the prostate and snap-frozen. Tissue samples (3 mm in diameter) were thereafter drilled out of the frozen tissue slice, and locations of removed samples were documented on a photo. Preoperative MR images best corresponding to the level of the resected tissue slice were identified, and circular regions of interest were outlined according to the location of the removed tissue samples. Patient and tissue sample characteristics are listed in Table 1.

**MR Imaging**

Magnetic resonance imaging was performed as previously described (2). In short, patients with biopsy proven prostate cancer underwent a preoperative multiparametric MR examination including $T_2$-weighted imaging ($T_2$WI), diffusion-weighted imaging (DWI), MR spectroscopic imaging [MRSI; results previously reported in Ref. (2, 5) and not shown here], and dynamic contrast-enhanced magnetic resonance imaging [DCE-MRI; results previously reported in Ref. (2) and not shown here] on a 3-T system (Magnetom Trio, Siemens Medical Solutions, Erlangen, Germany). Phased array body coil and spine coil elements were used for signal detection. $T_2$-weighted turbo spin echo images were obtained in three orthogonal planes. The transversal $T_2$-weighted images [TR/TE 4210 ms/104 ms, FOV 160 mm × 160 mm, matrix 320 × 256, slice thickness 3 mm, and acquisition time 5 min 47 s] were angulated perpendicular to the urethra to replicate the angle of slicing for histopathological analysis. Diffusion-weighted images [TR/TE 3500 ms/77 ms, FOV 340 mm × 168 mm, matrix 170 × 170, slice thickness 4 mm, four $b$-values (50, 300, 600, and 800 $s/mm^2$), and acquisition time 2 min 59 s] and dynamic contrast-enhanced images [TR/TE 4 ms/1.34 ms, FOV 280 mm × 227.6 mm, matrix 256 × 230.4, slice thickness 2 mm, temporal resolution 12.9 s, and total acquisition time 5 min 32 s] were equally angulated. $T_2$-weighted images and ADC maps were used to calculate $T_2$ intensities and ADC in regions of interest corresponding to tissue resection areas.

**HR-MAS MRS Experiment**

$^1$H HR-MAS MR spectra of the tissue samples were obtained using a 14.1-T spectrometer (Bruker Avance DRX 600, Bruker BioSpin).

**Table 1 | Characteristics of patients ($N = 15$) and tissue samples ($n = 53$).**

<table>
<thead>
<tr>
<th>Patient characteristics</th>
<th>Value, median (range)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (years)</td>
<td>63.7 (48.0–69.5)</td>
</tr>
<tr>
<td>sPSA (ng/ml)</td>
<td>12.0 (5.9–21.4)</td>
</tr>
<tr>
<td>Tissue sample characteristics</td>
<td>n - values (50, 300, 600, and 800 $s/mm^2$)</td>
</tr>
<tr>
<td>Non-cancer</td>
<td>14</td>
</tr>
<tr>
<td>6 (3 + 3)</td>
<td>11</td>
</tr>
<tr>
<td>7 (3 + 4, 4 + 3)</td>
<td>18 (4, 14)</td>
</tr>
<tr>
<td>8 (3 + 5, 4 + 4)</td>
<td>8 (2, 6)</td>
</tr>
<tr>
<td>9 (4 + 5, 5 + 4)</td>
<td>8 (6, 3)</td>
</tr>
<tr>
<td>Tumor load (%)a, median (range)</td>
<td>60 (10–90)</td>
</tr>
</tbody>
</table>

$s$PSA, serum prostate-specific antigen at time of surgery; $n =$ number of patients tissue samples.

*aPercentage tumor in tumor-containing tissue samples.*
of total area was used in the calculations. One tissue sample (GS 4 + 4) was excluded from color-based segmentation due to poor quality of the H&E slide. The metabolite concentrations from this tissue sample were included in the analyses when metabolites were correlated with MR parameters.

Statistics
The Kolmogorov–Smirnov test was used to test data normality. Linear mixed model was used for pairwise comparison of histological components and different Gleason scores. Linear mixed model was also used to evaluate the association between MR parameters, metabolites, and histological features. Parameters that were not normally distributed were log-transformed before being entered in the linear mixed model. To account for multiple samples per patient, patient identification was entered as a random effect on the intercept in the model. Spearman’s rank correlation (r) was calculated between in vivo MR parameters, ex vivo metabolite concentrations, and histological features since they were not all normally distributed. This measure does not take into account multiple samples per patient. Multiple comparisons were corrected for with the Benjamini and Hochberg false discovery rate. Adjusted p values <0.05 were considered significant. All statistical analyses were performed using SPSS (IBM SPSS Statistics 22.0), except Benjamini and Hochberg corrections, which were performed in Matlab (MATLAB R2009a, The MathWorks Inc., Natick, MA, USA).

RESULTS
An overview of histological components in cancer and non-cancer samples is given in Table 2. Cancer samples have significantly lower area percentage of lumen and higher area percentage of nuclei than non-cancer samples (p < 0.001). Percentage area of stroma is not significantly different between cancer and non-cancer samples (p = 0.3). There are no significant differences between histological parameters in tissue samples with different Gleason scores except for percentage area of lumen in Gleason score 9 samples, which are significantly lower than for Gleason scores 6 and 7 (Figure 2).

The in vivo-measured MR parameters, T2 intensity and ADC, are reduced in areas of cancer compared to non-cancer (Table 3) (p < 0.001). There is a trend toward reduced ADC and T2 intensity with higher Gleason score and a significantly lower ADC in Gleason score 9 samples compared to Gleason score 6 samples (p = 0.02). The ex vivo-measured metabolite concentrations of citrate and choline is reduced and increased, respectively, in cancer samples compared to non-cancer samples (Table 3). There is a trend toward increased choline and decreased citrate with increased Gleason score and significantly lower citrate in Gleason score 9 samples compared to Gleason score 6 samples (p = 0.01). There are no significant differences in choline between samples with different Gleason scores.

FIGURE 1  | (A) ADC map with ROIs from cancer (red outline) and non-cancer peripheral zone (green outline) areas corresponding to extracted tissue samples. (B) H&E stained slides (4x objective) of cryosections with Gleason 5 + 4 (red) and non-cancer (green) tissue. Red and green outline refer to tissue location in (A). (C) Close-up view of corresponding color-based segmentation with setting 2. Negative pixels (blue color) correspond to stroma; positive pixels (yellow and orange color) correspond to cytoplasm and nuclei; lumen is calculated as total number of pixels minus positive and negative (white color).

GmbH, Karlsruhe, Germany) and post-processed as previously described (20). Quantification of metabolites was performed by LC Model (21), as described by Giskeødegård et al. (20).

Histopathology and Color-Based Segmentation
A cryosection was taken from each tissue sample and stained with hematoxylin and eosin (H&E). These H&E-stained slides were digitized with 4x magnification, and color-based segmentation (Positive Pixel Count algorithm in ImageScope v.11, Aperio Technologies) was used to identify luminal space, stroma, and nuclei, as described by Langer et al. (16). In short, two hue and windows settings were used (setting 1: 0.1 for hue, 0.5 for window; setting 2: 0.7 for hue, 0.35 for window) and optimized for each histologic slide by adjusting the window on a test region such that negative pixels in setting 1 represented nuclei, negative pixels in setting 2 represented stroma, and positive pixels in setting 2 represented cytoplasm and nuclei. Lumen was calculated as total area minus positive and negative pixels in setting 2 (Figure 1). For all the components (lumen, stroma, and nuclei), percentage of total area was used in the calculations. One tissue sample (GS 4 + 4) was excluded from color-based segmentation due to poor quality of the H&E slide. The metabolite concentrations from this tissue sample were included in the analyses when metabolites were correlated with MR parameters.
There is an intermediate to strong positive correlation between ADC and percentage area of lumen ($p = 0.6$, $p < 0.001$). There is a weaker, but significant, negative correlation between ADC and percentage area of nuclei ($p = -0.35$, $p = 0.01$) (Table 4; Figure 3). ADC, percentage area of lumen, and percentage area of nuclei are all correlated with Gleason score in the tissue sample ($p = -0.58$, $p < 0.001$; $p = -0.62$, $p < 0.001$; and $p = 0.4$, $p = 0.01$, respectively) (Table 4). When only non-cancer samples ($n = 14$) are considered, there is still a strong correlation between ADC and lumen ($p = 0.7$, $p = 0.005$), while the correlation between ADC and nuclei is no longer significant ($p = -0.007$, $p = 0.982$).

In linear mixed model analysis, percentage area of lumen, nuclei, and Gleason score are significant covariates of ADC ($p < 0.001$, $p = 0.014$, and $p < 0.001$, respectively). Percentage area of stroma is not significantly correlated with ADC or Gleason score (Table 4).

There is a positive correlation ($p = 0.31$) between Citrate and percentage area of lumen, however after correcting for multiple testing, it is only borderline significant at the 0.05 level ($p = 0.053$). With linear mixed model, lumen is a significant covariate of citrate ($p = 0.005$). Choline is negatively correlated with lumen ($p = -0.38$, $p = 0.02$) and positively correlated with percentage area of nuclei ($p = 0.38$, $p = 0.02$). Both citrate and choline are significantly correlated with Gleason score ($p = -0.46$, $p = 0.002$; $p = 0.55$, $p < 0.001$, respectively). When only non-cancer samples are considered ($n = 14$) citrate is not significantly correlated with percentage area of lumen ($p = 0.15$, $p = 0.65$) and choline is not significantly correlated with lumen or nuclei ($p = 0.01$, $p = 0.95$; $p = 0.24$, $p = 0.41$, respectively). The relationship between percentage area of lumen, citrate, choline, and Gleason score is visualized in Figure 4. Spermine and creatine are not significantly correlated with any of the glandular components, but the ratios (choline + spermine + creatine)/citrate and (citrate + spermine + creatine)/choline are significantly correlated with percentage area of lumen ($p = 0.49$, $p = 0.002$; $p = -0.45$, $p = 0.004$, respectively) (Table 4).

**DISCUSSION**

In this study, we have explored the relationship between MR parameters measured in vivo (ADC and $T_2$ intensity), metabolite concentrations (citrate, choline, creatine, and spermine) measured ex vivo with HR-MAS MRS, and histological gland components (percentage area of lumen, nuclei, and stroma) extracted from histopathology with color-based segmentation.
As expected, we find a clear difference in histological features between cancer and non-cancer samples. The decrease in percentage of luminal space and increase in percentage area of nuclei in cancer samples compared to non-cancer samples is in line with previous reported findings by several groups (16, 22, 23). The percentage area of lumen in healthy prostate tissue has previously been reported to be in the range of 20–30% (16, 22), which is higher than we observed. Previous studies have performed color-based segmentation on H&E-stained whole mount sections while we used H&E-stained cryosections. H&E staining of cryosections can be of lower quality than staining of paraffin-embedded tissue, and this might explain some of the discrepancy. Further, our samples were 3 mm in diameter, and luminal space on the edges was omitted in the segmentation, which could also result in lower values. However, we observed a trend toward decreased percentage area of lumen with increased Gleason score, consistent with previous findings (16, 23). The literature is inconsistent regarding stroma content in cancer versus non-cancer tissue (16, 22, 23); however, there was no evidence of differences in percentage area of stroma in our data.

We find a significantly increased percentage area of nuclei in cancer samples compared to non-cancer samples. In cancer samples, we see a weak trend toward increased area of nuclei with increased Gleason score; however, there is no significant difference between different Gleason scores. A trend toward increasing cellularity metrics with increasing Gleason pattern is described in numerous articles, but most of them lack significant differences between Gleason groups (16, 17, 22–24). This has often been attributed to the limited range of Gleason scores in the study cohorts. However, in our study, we have tissue samples ranging from non-cancer to Gleason score 9 with approximately equal number of samples in each group and still find no significant differences. This might be a result of the nature of the Gleason scoring system, where each pattern is based on tissue architecture rather than cellularity per se. This finding might also explain the lack of significant differences in choline between different Gleason scores.
would therefore make sense to calculate a metabolite ratio where citrate and spermine are added. Kobus et al. (22) focused on the metabolite ratio citrate, spermine, and creatine-to-choline and found a positive correlation with lumen with correlation coefficient of 0.5. We find a correlation in the same range between this ratio and percentage area of lumen ($\rho = 0.45$), even though the individual metabolites are weaker correlated with lumen.

This study has some limitations. The low number of tissue samples hampered subdivision into peripheral zone and transition zone samples. There are also some inherent difficulties in the matching between \textit{in vivo} MR images and tissue samples resected for HR-MAS NMR experiments and histopathology; however, the tissue harvesting method used in this study contributes to minimize the matching uncertainty (19). Previous studies have shown that there is a good correlation between \textit{in vivo} and \textit{ex vivo} measured metabolite ratios by using the harvesting and matching technique used in this study (5).

The acquired $T_2$-weighted MR images did not allow for quantitative $T_2$ measurements, and we therefore used $T_2$ intensity, rather than $T_2$ relaxation, in our analysis. Since $T_2$ intensity can be affected by other factors than pure $T_2$ relaxation, we chose to only do brief descriptive statistics of the $T_2$ intensity. Further for ADC calculations, we used a monoexponential diffusion model, since this is the model which is available for ADC calculation in our clinical scanners. The monoexponential diffusion model assumes Gaussian diffusion conditions. This could affect the correlation between ADC and percentage area lumen, since the lumen contains freely diffusing liquid. Bourne et al. suggest that the distinct stromal and glandular diffusion compartments are the origin of the biexponential diffusion decay seen \textit{in vivo} (6). Therefore, application of a biexponential diffusion model might have resulted in a higher correlation of ADC with area of nuclei or stroma.

All samples in this study were taken from patients with prostate cancer. The non-cancer samples were taken from a transversal prostate slice that contained prostate cancer elsewhere, and it is not yet fully understood how the metabolism in normal-appearing tissue is affected by the adjacent cancer (field effect). The citrate production in the histopathologically normal-appearing areas could already have been altered and therefore preclude the expected correlation between citrate and percentage of lumen in non-cancer samples.

**CONCLUSION**

This study adds to the literature of associations between alterations in tissue composition, metabolism, and observed MR imaging parameters. The microstructures that are observed by histopathology are linked to MR characteristics in prostate cancer, and ADC appears to mainly reflect luminal space rather than dense tumor structures.

**AUTHOR CONTRIBUTIONS**

KS: study design, data collection, data analysis and interpretation, drafting the manuscript, and critical revision of the manuscript. RV: data analysis and interpretation, drafting the manuscript, and critical revision of the manuscript. HB: data collection and

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critical revision of the manuscript. AW: data analysis and interpretation and critical revision of the manuscript. AH and AA: study design and critical revision of the manuscript. M-BT: study design, data collection, data analysis and interpretation, and critical revision of the manuscript. TB: study design, data analysis and interpretation, and critical revision of the manuscript.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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