Prostate MRI: Evaluating Tumor Volume and Apparent Diffusion Coefficient as Surrogate Biomarkers for Predicting Tumor Gleason Score

Olivio F. Donati¹,4, Asim Afaq¹,5, Hebert Alberto Vargas¹, Yousef Mazaheri², Junting Zheng³, Chaya S. Moskowitz³, Hedvig Hricak¹, and Oguz Akin¹

Abstract

Purpose: To investigate whether tumor volume derived from apparent diffusion coefficient (ADC) maps (VolumeADC) and tumor mean ADC value (ADCmean) are independent predictors of prostate tumor Gleason score (GS).

Experimental Design: Tumor volume and GS were recorded from whole-mount histopathology for 131 men (median age, 60 years) who underwent endorectal diffusion-weighted MRI for local staging of prostate cancer before prostatectomy. VolumeADC and ADCmean were derived from ADC maps and correlated with histopathologic tumor volume and GS. Univariate and multivariate analyses were performed to evaluate prediction of tumor aggressiveness. Areas under receiver-operating characteristics curves (AUC) were calculated to evaluate the performance of VolumeADC and ADCmean in discriminating tumors of GS 6 and GS ≥7.

Results: Histopathology identified 116 tumor foci >0.5 mL. VolumeADC correlated significantly with histopathologic tumor volume ($r = 0.683$). The correlation increased with increasing GS ($r = 0.453$ for GS 6 tumors; $r = 0.643$ for GS 7 tumors; $r = 0.980$ for GS ≥8 tumors). Both VolumeADC ($r = 0.286$) and ADCmean ($r = -0.309$) correlated with GS. At univariate analysis, both VolumeADC ($P = 0.0325$) and ADCmean ($P = 0.0033$) could differentiate GS = 6 from GS ≥7 tumor foci. However, at multivariate analysis, only ADCmean ($P = 0.0156$) was a significant predictor of tumor aggressiveness (i.e., GS 6 vs. GS ≥7). For differentiating GS 6 from GS ≥7 tumors, AUCs were 0.644 and 0.704 for VolumeADC and ADCmean, respectively, and 0.749 for both parameters combined.

Conclusion: In patients with prostate cancer, ADCmean is an independent predictor of tumor aggressiveness, but VolumeADC is not. The latter parameter adds little to the ADCmean in predicting tumor GS.

Introduction

It was estimated that 30% to 50% of the approximately 238,590 American men diagnosed with prostate cancer in 2013 would have an indolent form of the disease unlikely to become life-threatening. These men could potentially take advantage of an increasing spectrum of patient-tailored disease management options—including active surveillance and various forms of focal ablation—that are designed to minimize adverse treatment-related effects (1–3). However, to ensure that patients are indeed suited for such conservative management approaches, it is essential not only to detect and localize prostate cancer, but also to assess its aggressive potential—a task that remains challenging. Clinical, biochemical and pathologic features are typically used to triage patients according to the likelihood of rapid disease progression (4–8).

Recently, diffusion-weighted MRI (DWI) has garnered interest for its potential to noninvasively characterize prostate cancer aggressiveness. DWI probes variations in free water movement within tissues, which tends to be more restricted in the presence of tumor due to changes in cell number, size, and architecture. On DWI images, variations in water diffusion manifest as changes in signal intensity, and degrees of diffusion restriction can be assessed quantitatively by means of the apparent diffusion coefficient (ADC). A relatively simple metric, the ADC can be calculated on a pixel-by-pixel basis with clinical MRI platforms.

Note: Supplementary data for this article are available at Clinical Cancer Research Online (http://clincancerres.aacrjournals.org/).

Corresponding Author: Oguz Akin, Department of Radiology, Memorial Sloan-Kettering Cancer Center, 1275 York Avenue, New York, NY 10065. Phone: 212-639-3458; Fax: 1 212-794-4010; E-mail: akinod@mskcc.org
doi: 10.1158/1078-0432.CCR-14-0044
©2014 American Association for Cancer Research.
Clinical Cancer Research

Volume derived from ADC maps (VolumeADC) are indistinguishable tumors with GSs of 6 from those with GSs of 7 or higher. Independent predictors of tumor GS and can be used to distinguish tumors of GS 6 from tumors of GS 7. Our findings indicate that independent of the tumor volume, ADCmean correlated with tumor GS, but on multivariate analysis only ADCmean independently distinguished tumors of GS 6 from tumors of GS ≥7. The purpose of our study was to investigate the potential synergy of these two parameters in evaluating prostate cancer aggressiveness.

A number of studies have shown an inverse correlation between ADC values on DWI and prostate cancer Gleason scores (GS; refs. 9–14). However, the ADC values of prostate cancer foci with different GS overlap, and no method has been developed to determine the GS unequivocally based on ADC analysis alone (9–14).

Pathology studies have shown that higher tumor volumes are associated with higher GSs and worse outcomes (15, 16). Tumor volume measured on DWI correlates well with histopathologic tumor volume, and the strength of the correlation increased with the tumor GS. Both VolumeADC and ADCmean correlated with tumor GS, but on multivariate analysis only ADCmean independently distinguished tumors of GS 6 from tumors of GS ≥7. Our findings indicate that independent of the tumor volume, ADCmean could serve as a biomarker to predict prostate cancer aggressiveness.

Translational Relevance

Many prostate cancers diagnosed today are likely indolent, but better means of assessing prostate cancer prognosis are needed to identify the appropriate, patient-specific treatment option. Distinguishing tumors of Gleason score (GS) 6 from tumors of GS ≥7 is especially critical for assessing eligibility for active surveillance. In patients who underwent diffusion-weighted MRI before radical prostatectomy, we assessed the value of the mean apparent diffusion coefficient (ADCmean) and the tumor volume measured from ADC maps (VolumeADC) for predicting two important prognostic factors: tumor volume and tumor GS on histopathology. VolumeADC correlated well with histopathologic tumor volume, and the strength of the correlation increased with the tumor GS. Both VolumeADC and ADCmean correlated with tumor GS, but on multivariate analysis only ADCmean independently distinguished tumors of GS 6 from tumors of GS ≥7. Our findings indicate that independent of the tumor volume, ADCmean could serve as a biomarker to predict prostate cancer aggressiveness.

Materials and Methods

The IRB approved our retrospective study and waived the informed consent requirement. Our study was compliant with the Health Insurance Portability and Accountability Act.

Patients

Patients who underwent MRI of the prostate including DWI between July 2008 and April 2010 were considered to be peripheral zone tumors (9). All images were acquired on a 1.5-Tesla MRI system (GE Healthcare Technologies). A body coil was used for excitation; a pelvic four-channel phased-array coil and an endorectal coil (Medrad) were used for signal reception. T1-weighted, T2-weighted, and DWI sequences were acquired but only DWI sequence was used for analysis in this study. DWI was performed using a single-shot spin-echo echoplanar imaging sequence with b = 0, 1,000 seconds/mm² (TR/TE, 1,200–6,800 ms/40–113 ms; section thickness, 3–4 mm; no intersection gap; field of view, 12–16 cm; matrix, 96 × 96–128 × 128). Parametric maps of ADC values were calculated using a designated workstation (Advanced Workstation, GE Medical Systems).

MRI acquisition

Histopathologic preparation.

After prostatectomy, specimens were submitted to histopathology, where they were sliced from apex to base at 3 to 4 mm intervals. Microslices were placed on glass slides and stained with hematoxylin and eosin after paraffin embedding. For each patient, one of two dedicated genitourinary pathologists at our institution with more than 30 years of combined experience verified and assigned a GS for each tumor outlined on the histology slides.

Measurement of histopathologic tumor volume.

Tumor volume on pathology slices was measured in consensus by two of the authors using software (ImageJ, version 1.47a; NIH, Bethesda, MD). If a lesion extended into more than one pathologic slice, the areas of tumor foci on all slices were summed to obtain an estimate of the histopathologic volume of the whole lesion. Tumors that covered both zones, the transition zone as well as the peripheral zone, were considered to be transition zone tumors if more than 70% of the tumor was in the transition zone (20); all others were considered to be peripheral zone tumors (9).

Correlation of lesions on MRI and histopathology.

Working in consensus, three radiologists (with 1, 1, and 9 years of experience in interpreting prostate MRI) correlated MR images with whole-mount pathology maps to establish the locations of tumors on MRI. Using software (ImageJ, version 1.47a; NIH), the radiologists drew a freehand region of interest (ROI) around the discernible tumor tissue on the ADC maps (19). If a tumor was depicted on more than one slice, all traced ROIs corresponding to that...
tumor were included in the estimation of the tumor volume (VolumeADC) and the calculation of the mean ADC value (ADCmean; ref. 19). On each slice containing tumor, the area of the tumor focus was determined on a voxel basis by considering the acquisition matrix, reconstruction matrix as well as the FOV. VolumeADC (mL) was calculated as [sum of all tumor areas on the slices (cm²) × slice thickness (cm)].

**Statistical analysis**

The correlation between VolumeADC and volume derived from histopathology as well as the correlations of VolumeADC and ADCmean with tumor GS were assessed using Spearman correlation coefficient (ρ). The between-subject correlation coefficient proposed by Bland and Altman (21) was calculated and tested to take into account multiple lesions per patient.

To evaluate whether VolumeADC and ADCmean could differentiate a GS of 6 from a GS ≥7, a generalized linear regression and generalized estimating equations method was used with an independent correlation structure and robust covariance matrix, to take into account multiple lesions per patient. Univariate and multivariate analyses with both VolumeADC and ADCmean as covariates were performed. The OR describing the likelihood of a tumor performance of VolumeADC and ADCmean in discriminating tumors with both VolumeADC and ADCmean as covariates were assessed from histopathology as well as the correlations of volumeADC, and ADCmean with tumor GS were assessed using Spearman correlation coefficient (ρ). The between-subject correlation coefficient proposed by Bland and Altman (21) was calculated and tested to take into account multiple lesions per patient.

In a univariate analysis including all lesions (peripheral zone and transition zone), both VolumeADC and ADCmean could differentiate tumors of GS 6 from those with a GS ≥7 (OR, 1.73 for VolumeADC and 0.64 for ADCmean; P = 0.0325 and 0.0033, respectively; Table 2). In a subanalysis considering only tumors originating in the peripheral zone, ADCmean could differentiate between tumors of GS 6 and those with a GS ≥7 (P = 0.0025), but VolumeADC could not (P = 0.2709; Table 2). The number of lesions originating in the transition zone was too small to permit a subanalysis.

In a multivariate analysis, after adjustments were made for the influence of VolumeADC, ADCmean independently discriminated between tumors of GS 6 and tumors of GS ≥7 (P = 0.0156; Fig. 2). However, after adjustments were made for the influence of ADCmean, VolumeADC could not independently differentiate between these two tumor GS categories (P = 0.0733; Table 2; Fig. 3).

Accuracy in discriminating tumors of GS 6 from those with a GS ≥7 was slightly lower for VolumeADC than for ADCmean (AUC = 0.644 and AUC = 0.704, respectively; P = 0.3262). Combining these variables as covariates in a

Correlation of volumeADC and histopathologic tumor volume

The Spearman correlation coefficient for VolumeADC and histopathologic tumor volume in lesions >0.5 mL was ρ = 0.683 (P < 0.0001; Fig. 1). The correlation coefficient increased as the tumor GS increased, rising from ρ = 0.453 (P = 0.1042) for tumors with a GS of 6 (3+3) to ρ = 0.643 (P < 0.0001) for tumors with a GS of 7 (3+4 or 4+3) and ρ = 0.980 (P < 0.0001) for tumors with a GS ≥8 (Table 1). The correlation between histopathologic tumor volume and VolumeADC was highest for tumors of GS ≥8.

**Correlations of histopathologic tumor volume, volumeADC and ADCmean with GS**

Histopathologic tumor volume and VolumeADC both correlated positively with tumor GS [P = 0.336 (P = 0.0017) and P = 0.286 (P = 0.0081), respectively], whereas ADCmean correlated negatively with tumor GS [P = −0.309 (P = 0.0087)].

**Differentiation of tumor aggressiveness by volumeADC and ADCmean**

In a univariate analysis including all lesions (peripheral zone and transition zone), both VolumeADC and ADCmean could differentiate tumors of GS 6 from those with a GS ≥7 (OR, 1.73 for VolumeADC and 0.64 for ADCmean; P = 0.0325 and 0.0033, respectively; Table 2). In a subanalysis considering only tumors originating in the peripheral zone, ADCmean could differentiate between tumors of GS 6 and those with a GS ≥7 (P = 0.0025), but VolumeADC could not (P = 0.2709; Table 2). The number of lesions originating in the transition zone was too small to permit a subanalysis.

In a multivariate analysis, after adjustments were made for the influence of VolumeADC, ADCmean independently discriminated between tumors of GS 6 and tumors of GS ≥7 (P = 0.0156; Fig. 2). However, after adjustments were made for the influence of ADCmean, VolumeADC could not independently differentiate between these two tumor GS categories (P = 0.0733; Table 2; Fig. 3).

Accuracy in discriminating tumors of GS 6 from those with a GS ≥7 was slightly lower for VolumeADC than for ADCmean (AUC = 0.644 and AUC = 0.704, respectively; P = 0.3262). Combining these variables as covariates in a
multivariate model resulted in a minor increase in AUC (to 0.749; Supplementary Fig. 1).

Discussion

In our study, tumor volume measured on ADC maps correlated with tumor volume on histopathology, and the strength of the correlation increased with the tumor GS. In addition, the ADCmean but not the VolumeADC independently differentiated tumors of GS 6 from those of GS 7 or above. The VolumeADC added little to the ADCmean in predicting the tumor GS.

The correlation between VolumeADC and histopathologic tumor volume in our study ($r = 0.68$) was very similar to that reported by Isebaert and colleagues ($r = 0.75$; ref. 17), and it was slightly higher than the correlation between tumor volume on T2-weighted MRI and histopathologic tumor volume reported by Turkbey and colleagues ($r = 0.63$; ref. 23). The difference between our result and that of Turkbey and colleagues is consistent with an earlier study by Mazaheri and colleagues, which found that prostate cancer tumor volume measurements based on ADC maps correlated better with histopathologic tumor volumes than did measurements based on T2-weighted MRI (17, 18). Furthermore, the correlation between imaging- and histopathology-derived tumor volumes may have been stronger in our study because, unlike Turkbey and colleagues, we used a pixel-based calculation to determine imaging and histopathologic tumor volumes, outlining tumor borders instead of using the ellipsoid formula, which is based on linear measurements and does not take into account the irregular shapes of prostate cancer foci.

In keeping with the existing literature, we demonstrated that ADC-based tumor volume and histopathologic tumor volume correlate better in prostate cancer foci with higher GSs. This may be explained by the fact that tumors with higher GSs are better depicted on ADC maps because they contrast more strongly with benign tissue (10, 13); this makes it easier to trace the borders of the lesions and likely results in more accurate representations of the actual areas of tumor on ADC maps.

The correlation between VolumeADC and GS in our study ($r = 0.29$) was similar to that recently reported by Verma and colleagues ($r = 0.35$; ref. 24). Likewise, the correlation between ADCmean and GS in our patient cohort ($r = 0.31$) was within the range of such correlations reported in recent studies ($r = 0.26$ to $0.38$; refs. 12, 14, 24). At multivariate analysis, ADCmean was the only parameter that independently predicted the category of the tumor GS (GS 6 vs.

### Table 1. Correlation between tumor volume on histopathology and tumor volume on ADC maps

<table>
<thead>
<tr>
<th></th>
<th>$r$</th>
<th>$P$</th>
<th>Lesions</th>
<th>Patients</th>
</tr>
</thead>
<tbody>
<tr>
<td>All tumors</td>
<td>0.683</td>
<td>$&lt;$0.0001</td>
<td>116</td>
<td>85</td>
</tr>
<tr>
<td>PZ tumors</td>
<td>0.706</td>
<td>$&lt;$0.0001</td>
<td>89</td>
<td>73</td>
</tr>
<tr>
<td>TZ tumors</td>
<td>0.677</td>
<td>0.0003</td>
<td>27</td>
<td>24</td>
</tr>
<tr>
<td>GS 6 tumors</td>
<td>0.453</td>
<td>0.1042</td>
<td>16</td>
<td>14</td>
</tr>
<tr>
<td>GS 7 tumors</td>
<td>0.643</td>
<td>$&lt;$0.0001</td>
<td>92</td>
<td>71</td>
</tr>
<tr>
<td>GS $\geq$8 tumors</td>
<td>0.980</td>
<td>$&lt;$0.0001</td>
<td>8</td>
<td>8</td>
</tr>
</tbody>
</table>

NOTE: 95% CI was estimated using the bootstrapping method of resampling patients.

Abbreviations: PZ, peripheral zone; TZ, transition zone.

### Table 2. Results of univariate and multivariate analyses for prediction of tumor GS $\geq$7 by ADCmean and VolumeADC

<table>
<thead>
<tr>
<th></th>
<th>All lesions $&gt;0.5$ mL</th>
<th>P</th>
<th>PZ lesions $&gt;0.5$ mL</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>OR (95% CI)</strong></td>
<td><strong>P</strong></td>
<td><strong>OR (95% CI)</strong></td>
<td><strong>P</strong></td>
<td></td>
</tr>
<tr>
<td>Univariate analysis</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>ADCmean (100-unit increment)</td>
<td>0.64 (0.47–0.86)</td>
<td>0.0033</td>
<td>0.50 (0.32–0.78)</td>
<td>0.0025</td>
</tr>
<tr>
<td>VolumeADC (0.5 mL increment)</td>
<td>1.73 (1.05–2.87)</td>
<td>0.0325</td>
<td>1.50 (0.73–3.07)</td>
<td>0.2709</td>
</tr>
<tr>
<td>Multivariate analysis</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>ADCmean (100-unit increment)</td>
<td>0.68 (0.50–0.93)</td>
<td>0.0156</td>
<td>0.51 (0.33–0.79)</td>
<td>0.0025</td>
</tr>
<tr>
<td>VolumeADC (0.5 mL increment)</td>
<td>1.57 (0.96–2.59)</td>
<td>0.0733</td>
<td>1.14 (0.66–1.98)</td>
<td>0.6359</td>
</tr>
</tbody>
</table>

NOTE: OR interpretation: the tumors are less likely to have a GS of 7 or above as the ADCmean increases (OR for 100-unit increase = 0.68, 95% CI, 0.50–0.93), controlling for VolumeADC.

Transition zone tumors were not assessed separately due to the small number of lesions in the transition zone.

Abbreviation: PZ, peripheral zone.
GS ≥ 7). It seems that though the predictive value of ADC_mean for tumor aggressiveness is independent of tumor size, when ADC_mean is not clearly predictive, Volume_ADC cannot be used to resolve the ambiguity. These results contrast with those of a recent study by Verma and colleagues, in which both mean ADC value and Volume_ADC were identified as significant predictors of tumor aggressiveness in the peripheral zone at multivariate analysis (24). There are several possible reasons for the discrepancy. First, different statistical methods were used for the multivariate analyses of the two studies. Second, our measurements were based on whole-mount step-section pathology slides instead of recreated histologic maps. Third, the \( b \) values used to create the ADC maps in our patient cohort (\( b = 0, 1,000 \) seconds/mm\(^2\)) differed from those used in the other study (\( b = 0, 600 \) seconds/mm\(^2\); ref. 24). (ADC values are dependent on the chosen \( b \) values; ref. 25, and therefore ADC measurements cannot be compared between protocols using different \( b \) values. However, as long as the imaging parameters, including the \( b \) value, are kept constant, ADC values measured in the abdomen may be comparable across different scanners and field strengths; ref. 26). Fourth, in the study by Verma and colleagues, only the voxels of the most central slice were used to calculate ADC parameters. Although the results of the multivariate analyses differed, accuracy levels in identifying prostate cancer foci of GS ≥ 7 by combining ADC_mean and Volume_ADC were similar in the two studies (24).

We acknowledge the following limitations of our study: First, so that we would be able to correlate imaging findings with histopathology, we only included patients who underwent radical prostatectomy, causing a selection bias. Therefore, our results may not apply to a broader population of patients with newly diagnosed prostate cancer, especially because there is a trend for increasing use of active surveillance of low-risk prostate cancer (27). However, this selection bias is inherent to every study that uses whole-mount step-section histopathology specimens as a reference standard for evaluating imaging variables. Second, an endorectal coil was used for acquisition of MRI, potentially deforming the prostate gland and the tumor foci. However, the use of an endorectal coil provides a higher signal-to-noise ratio (28) and may therefore be preferable for quantitative ADC analysis. Third, our approach of retrospectively delineating the prostate cancer foci on ADC maps using the histopathology maps as a guide does not represent the sequence of events in the clinical setting, where histopathology maps would not be available at the time of MRI.

Figure 2. Top row, whole-mount histopathology slice (A) and ADC map (\( b = 0, 1,000 \) seconds/mm\(^2\); B) demonstrating a prostate cancer focus with a GS of 3+3 (arrows), a total tumor volume of 0.65 mL and a mean ADC of 1,165.2 \( \times 10^{-3} \) mm\(^2\)/second. The bottom row shows the whole-mount histopathology slice (C) and ADC map (\( b = 0, 1,000 \) seconds/mm\(^2\); D) of a prostate cancer focus with a GS of 3+4 (arrows). Despite having a histopathologic volume (0.68 mL) similar to that of the tumor focus in A, the tumor focus shown in C and D has a lower mean ADC (964.2 \( \times 10^{-3} \) mm\(^2\)/second). Note: Images A and C show only one representative slice out of 12 and eight total histopathology slices, respectively. The tumors in A and C were both present on seven contiguous slices.

www.aacrjournals.org Clin Cancer Res; 20(14) July 15, 2014 3709

Published OnlineFirst May 21, 2014; DOI: 10.1158/1078-0432.CCR-14-0044
Therefore, we are not able to provide information on the accuracy of prostate cancer detection in this study or on the effect that potentially missed lesions might have had on our results. Furthermore, using histopathology slices for identification of tumor foci may have introduced a potential bias in the evaluation of lesion volume on ADC maps as the location of tumors was available to the radiologists encircling the tumor foci. Although, the ROI drawn by the radiologists for the purposes of this study only contained clearly discernible tumor tissue on ADC maps (e.g., voxels that were visually darker than the surrounding healthy tissue), we acknowledge that the correlations reported in this study would be influenced by the diagnostic accuracy of prostate cancer detection in routine clinical practice.

In summary, our results suggest that although $\text{Volume}_{\text{ADC}}$ is useful to predict true tumor volume, $\text{ADC}_{\text{mean}}$ is the more useful parameter for distinguishing between GS 6 and higher GS tumors, a distinction that is critical for identifying suitable candidates for active surveillance.

**Disclosure of Potential Conflicts of Interest**

No potential conflicts of interest were disclosed.

**Authors’ Contributions**

**Conception and design:** O.F. Donati, A. Afaq, H. Hricak, O. Akin

**Development of methodology:** O.F. Donati, A. Afaq, Y. Mazaheri, O. Akin, H.A. Vargas, H. Hricak, O. Akin

**Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.):** A. Afaq, Y. Mazaheri, O. Akin

**Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis):** O.F. Donati, Y. Mazaheri, H.A. Vargas, J. Zheng, C.S. Moskowitz, O. Akin

**Writing, review, and/or revision of the manuscript:** O.F. Donati, A. Afaq, H.A. Vargas, J. Zheng, H. Hricak, O. Akin

**Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases):** O.F. Donati, A. Afaq, O. Akin

**Study supervision:** O. Akin

**Acknowledgments**

The authors thank Ada Muellner for editing the article and Victor E. Reuter and Samson W. Fine for providing and verifying the histopathology slides.

**Grant Support**

This work was supported by the Swiss National Science Foundation and the Swiss Radiological Society (to O.F. Donati) and the Peter Michael Foundation (to O.F. Donati and A. Afaq).

The costs of publication of this article were defrayed in part by the payment of page charges. This article must therefore be hereby marked advertisement in accordance with 18 U.S.C. Section 1734 solely to indicate this fact.

Received January 9, 2014; revised May 5, 2014; accepted May 8, 2014; published OnlineFirst May 21, 2014.
References


