RESEARCH LETTER

Higher prevalence of lymph node metastasis in prostate cancer in patients with diabetes

Dear Editor,

Prostate cancer (PCa) and type 2 diabetes mellitus are among the most frequent diseases in men with a tremendous impact on morbidity and mortality (Giovannucci et al. 2010). Incidence of many common types of cancer is known to be higher in diabetes (Giovannucci et al. 2010). However, studies reported that incidence of PCa is not increased in men with type 2 diabetes, some studies even found reduced prevalence (Kasper et al. 2009). In contrast, PCa survival is markedly reduced in patients with coincident type 2 diabetes (Chen et al. 2017). The underlying molecular mechanisms for shortened survival are still under ongoing debate and not fully understood. They might include altered insulin or IGF-1 signaling and enhanced androgen receptor activity. Of note, diabetes and PCa share numerous risk factors, most importantly the nonmodifiable risk factor age and the modifiable risk factor obesity (Giovannucci et al. 2010). Although PCa incidence is not elevated in obese men, patients with excess bodyweight are reported to display higher cancer-related mortality (Ma et al. 2008). Carefully adjusted studies for these risk factors investigating the impact of diabetes on PCa outcomes and aggressiveness are sparse. To better understand why PCa-related survival is shortened in men with concomitant diabetes, we evaluated the relation of diabetes with TNM staging and an established PCa risk score (Mohler et al. 2016), independent of age and body weight.

Therefore, we studied 624 patients who were diagnosed with PCa and underwent a radical prostatectomy between June 2004 and September 2007 in the Department of Urology, University of Tübingen. Their diabetes status was evaluated from the hospital records or by patients’ questionnaires, after patients gave consent. Of these, 74 men had type 2 diabetes. Information on the antidiabetic therapy is presented in Table 1. As their age and BMI were higher than the mean of the patients without diabetes, we performed matching for these variables. Patients with history of another malignant tumor and type 1 diabetes were excluded. Two patients without diabetes were matched to each patient with diabetes applying propensity score matching, using the nearest match method of the MatchIt package in R (R Project for Statistical Computing, Vienna, Austria). Thus, 222 participants were included (74 patients with diabetes and 148 men without diabetes). As intended, they were comparable in age and BMI (median age (year) (interquartile range): 65 (49–75) and 65 (51–75); median BMI (kg/m²): 28.4 (22.5–57.5) and 28.4 (22.6–46.1), respectively). Gleason score and TNM status were carefully ascertained by an experienced pathologist. The Ethics Committee of the University of Tübingen approved the protocol.

The risk classification tool risk score based on the T-stage, Gleason score, PSA level (ng/mL) and N-stage was used as proposed by the National Comprehensive Cancer Network (NCCN) (Mohler et al. 2016). According to this, patients were stratified into 4 groups: low- (T1-T2a or Gleason score ≤6 or PSA <10 ng/mL), intermediate- (T2b-T2c or Gleason score 7 or PSA 10–20 ng/mL), high- (T3a or Gleason score 8–10 or PSA >20 ng/mL), and very high- (T3b-T4 or N1) risk groups.

Non-normally distributed parameters were log-transformed prior to analyses. Groups were compared by χ² or unpaired two-tailed t-tests. P values <0.05 were considered significant. The statistical software package JMP 11.0 (SAS Institute Inc, Cary, NC, USA) was used.

We found that patients with diabetes displayed no significant difference in serum PSA levels or Gleason scores compared to patients without diabetes (all P>0.2).

The entire distribution of the PCa risk score (NCCN) was not significantly different between men with and without diabetes (P=0.7). However, the proportion of patients in the very high-risk group was significantly greater among patients with diabetes (P=0.028, 20% vs 9%, Fig. 1A).
While the T-stage was not different between the two groups ($P=0.3$), analysis of the N-score revealed significant difference. Only 1% (2/148) of the non-diabetic patients had lymph node metastasis, while 10% (7/74) of men with diabetes already had infiltrated lymph nodes at the time of operation (Fig. 1B). In our data set, this results in a calculated odds ratio of diabetes for lymph node metastasis of 7.63.

Thus, we detected more aggressive tumors in men with diabetes, independent of age and body weight, as indicated by higher risk score and larger proportion of lymph node metastasis.

Abdollah and coworkers previously reported higher Gleason scores in diabetes (Abdollah et al. 2011). We could not reproduce this finding, possibly due to our careful matching for age and BMI, that has not been performed previously. Nevertheless, also in our study, two other parameters indicated more aggressive PCa in type 2 diabetes, namely the NCCN risk score and the N-stage. Though, N-stage is one criterion to calculate NCCN risk score. At least the latter one was suggested to be higher in diabetes in a previous report (Kronig et al. 2017). However, as obesity is a risk factor for lymph node metastasis in PCa (Schnoeller et al. 2015), higher body weight in diabetes could have confounded earlier analyses. We now excluded this possibility and ascertained more frequent lymph node infiltration in diabetes independent of body weight. Thus, our findings may indicate that known shorter PCa survival in men with diabetes could be due to more aggressive tumors. This should further be tested in prospective studies.

A number of molecular mechanisms can underlie these findings: As PCa is an androgen-dependent tumor, circulating and in particular intraprostatically synthesized androgens are in focus of current investigations. Very recently, we demonstrated an augmented gene expression machinery of the androgen receptor (AR) and downstream target genes underscoring activity of this signaling cascade.
in patients with diabetes (Lutz et al. 2017). Of note, the activity of the androgen chain in diabetes was detected despite of the known reduction of testosterone in this patient group.

It is a well-known phenomenon that insulin levels are elevated in type 2 diabetes, either to compensate insulin resistance or because of insulin-elevating therapies. We recently showed altered composition of insulin or IGF-1 receptors that favor the mitogenic isoforms in PCa (Heni et al. 2012). Hence, elevated insulin might promote tumor growth in diabetes.

Furthermore, hyperglycemia is a hallmark of diabetes. Men with higher blood glucose levels have more aggressive prostate cancer (Kim et al. 2010). Thus, more glucose availability may represent another pathomechanistic way toward more aggressive tumors and worse outcome.

One limitation of the study is that we have no evaluation of glycemic control or diabetes duration, i.e. time from diabetes diagnosis to surgery, of the included patients. Furthermore, no systematic follow-up of patient history was available. The applied NCCN risk score depends on status of lymph node metastasis and is therefore unable to address tumor aggressiveness independent of lymph node involvement. We can furthermore not exclude that different time of diagnosis in patients with and without diabetes might have influenced our results, even if comparable PSA levels argue against this possibility.

In conclusion, we report higher prevalence of lymph node metastasis and more high-grade tumors in men with PCa and coexistent type 2 diabetes. These findings may in part explain the worse outcome in patients with type 2 diabetes. Further studies should test the implication of our results for the antidiabetic as well as the antitumor therapies of patients with PCa and type 2 diabetes. They may pave the way toward more personalized treatment of this malignancy.

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Declaration of interest
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Author contribution statement
The study was designed by M H, A F and H U H. Data acquisition was performed by J H, M O S and M H. Data analysis and interpretation was done by R W, J M F, S Z L, T T, H U H, P M and M H. S Z L drafted the manuscript. R W, J H, A F, T T, M H, A S and H U H contributed to the discussion. All authors revised the manuscript and approved the final version to be published.

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