Sonic Hedgehog Expression is Associated with Lymph Node Invasion in Urothelial Bladder Cancer

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Abstract
Bladder cancer (BC) is a deadly disease characterized by high recurrence rates and frequent progression to an aggressive phenotype. Dysregulation of various signaling pathways have been implicated in BC tumorigenesis, however, the clinical relevance of sonic hedgehog pathway (Shh) remains under investigated. The aim of the current study was to analyze the prognostic value of Shh expression in patients with bladder carcinoma. Immunohistochemical expression of Shh was performed using tissue microarray with 128 specimens from bladder cancer patients. Kaplan-meier survival was analysed and correlation between Shh protein expression and patients’ clinicopathological parameters was examined using Fisher’s exact test. The immuno-staining results revealed that Shh protein exhibits cytoplasmic localization and is expressed in 49% of the analyzed bladder cancer cohort. Our data indicated that high Shh expression significantly correlated with increased lymph node metastasis (p = 0.02), however no association was reported between Shh expression and other clinicopathological parameters. High expression of sonic hedgehog was associated with lymph node invasion which may indicate that Shh might play an important role in progression and metastasis of bladder cancer.

Keywords Bladder cancer · Sonic hedgehog · Expression · Lymph node · Prognosis

Introduction
According to a recent report by Antoni et al. 2017, an estimated 430,000 new bladder cancer (BC) cases have happened in the year 2012 ranking BC the ninth most common type of cancer worldwide [1]. Approximately, 75% of the newly diagnosed BC patients have non-muscle invasive tumours (NMIBC) confined to the mucosa (stages Tis, Ta and T1) and the remaining 25% of patients have muscle invasive (MIBC) or metastatic disease at initial presentation in the clinic [2]. Patients with NMIBC disease have high risk for cancer recurrence and the possibility for progression to an aggressive muscle invasive form is also high which necessitate continuous surveillance and monitoring of BC patients [3]. Kaufman et al. (2009) demonstrated that 50–70% of non-muscle invasive tumours will recur over time and approximately 10–20% will progress to muscle invasive tumours (T2 - T4) [4]. This situation necessitates an urgent need to identify a reliable prognostic biomarker that would assist in clinical setting, with the currently-used grading/ staging system, to predict for early disease development and progression [5, 6]. Thus far, several promising biomarkers have been identified and a number of FDA-approved laboratory tests have been developed for regular cancer monitoring however most of which have not been sufficiently sensitive or specific to predict clinical outcome [7–9].

Sonic Hedgehog (Shh) is a member of the hedgehog (HH) family, which also includes desert hedgehog and Indian hedgehog. Shh was first discovered as a secreted protein...
mediating fundamental cellular processes such as vertebrate development and embryogenesis [10]. Later, it was discovered that SHh plays also important roles in regulating cell proliferation, differentiation and cell fate [11, 12]. Recent studies have demonstrated the role of Shh in mediating the tumorigenic properties of several cancer types including lung, prostate, breast, colon, ovarian, pancreatic and hepatocellular carcinoma [13–19].

Recent data revealed that sonic hedgehog is critical regulator of several cellular processes related bladder carcinogenesis. Constitutive activation of the Shh ligand binding to the trans-membrane protein Patched1 abrogates the inhibitory effect on smoothened (SMO) which undergo phosphorylation at accumulation at the cell surface. Activation of SMO generates downstream signaling cascade leading to nuclear translocation of the transcription factor Gli1 which in turn induces the transcription of several target genes including Gli1 [20–22]. A number of studies have shown that dysregulation of Shh ligand or one of its downstream mediators (Patched1, smoothened or Gli1) has been associated with urothelial carcinoma initiation and progression [23–25] and in regulating cancer stem cells activities [26]. It has been reported that Shh-expressing basal cells give rise to muscle-invasive bladder cancer phenotype [27]. Interestingly, targeted inhibition of sonic hedgehog or one of its signaling components has recently been recognized to be one of the key targets that would have significant clinical implications in novel cancer therapeutics [28, 29]. The aim of the current study was to investigate the expression of sonic hedgehog protein in patients with bladder cancer by immunohistochemistry. Correlation analysis between Shh expression and patients’ clinico-pathological parameters and its prognostic value as a biomarker in bladder cancer was also evaluated.

Materials and Methods

Patients and Samples Collection

Tissue specimens were collected from bladder cancer patients who underwent surgical resections at King Abdulaziz University hospital between 2005 and 2011. A cohort samples from 128 patients were formalin fixed and embedded in paraffin and stored, in the department of pathology within the same institution, until use. This study was ethically approved by the institutional research ethics committee, faculty of medicine, King Abdulaziz University (ref. N. 149–14).

Data Collection

Demographical, clinical and pathological data related to the collected tissue samples were also gathered and maintained in database. The collected information includes patients’ age, gender, marital status, tumour grade, stage, smoking. The mean follow-up was 30 months (range from 1 to 138 months). None of the patients had received chemo/radiation therapy before the surgery.

Tissue Microarray (TMA) Construction

Formalin-fixed paraffin-embedded blocks were used for tissue microarray construction. Two cores (0.6 mm each) from each patient were integrated into a recipient block using a tissue chip microarrayer. A total of 128 blocks of bladder cancer were successfully transferred to construct four TMA slides making the total of 256 tissue cores in the TMA (Table 1). Hematoxylin & Eosin staining was done to obtain representative tumour cores and confirm the presence of tumour.

| Parameters             | Bladder cancer (%) |
|------------------------|--------------------|
| Sex                    | Male 108/131 82.443% |
| Grade                  | High Grade 065/131 49.618% |
|                       | Low Grade 052/131 39.695% |
|                       | Unknown 014/131 10.687% |
| Age                    | < 60 Years 053/131 40.458% |
|                       | ≥ 60 Years 076/131 58.015% |
|                       | Unknown 002/131 01.527% |
| Blood Group            | A+ 022/131 16.794% |
|                       | A− 001/131 00.763% |
|                       | B+ 019/131 14.504% |
|                       | AB+ 003/131 02.290% |
|                       | O+ 038/131 29.008% |
|                       | O− 003/131 02.290% |
|                       | Unknown 045/131 34.351% |
| Type of Cancer         | MIBC 058/131 44.275% |
|                       | NMIBC 049/131 37.405% |
|                       | Undecided 024/131 18.321% |
| Subtypes               | Transitional 122/131 93.12% |
|                       | Squamous 006/131 04.580% |
|                       | Adenocarcinoma 001/131 00.765% |
|                       | Unknown 003/131 02.290% |
| Smoking                | YES 041/131 31.298% |
|                       | NO 026/131 19.847% |
|                       | Unknown 064/131 48.855% |
| Survival               | Died of Disease 038/131 29.008% |
|                       | Alive 091/131 69.466% |
|                       | Unknown 002/131 01.527% |
| Recurrence             | 051/131 38.931% |
Immunohistochemistry Staining

Immunohistochemical staining was performed using the Bench-Mark XT automated staining system (Ventana Medical Systems, Inc., Tucson, AZ, USA) according to the manufacturers’ instructions and as previously described [27]. Briefly, slides were de-waxed using EZprep solution for 30 min at 75 °C. Antigen retrieval was achieved using CC1 cell conditioning buffer for 64 min at 95 °C, then the slides were incubated with the primary rabbit anti-human sonic hedgehog polyclonal antibody (Spring Bioscience, USA) at a dilution of 1:100 for 30 min. Slides were then incubated with DAB chromogen detection kit as follows: application of universal DAB Inhibitor, then, universal DAB chromogen, then universal DAB H₂O₂, and last DAB Copper. Counterstaining was performed with hematoxylin and bluing reagent for 4 min. The sections were then dehydrated with ethanol then xylene and permanently mounted with mounting media and coverslip.

Scoring and Data Analysis

Scoring of the IHC staining was performed using light microscope (X40 objective). Sub-cellular localization and the intensity of the staining were recorded during the scoring process. For the intensity of the cytoplasmic staining, four categories were used on scale 0 to 3: 0 (−) = negative: no detectable staining; 1 (+) = weak, but still detectable staining; 2 (++) = moderate, clearly positive but still weak: and 3 (+++) = strong staining. To calculate the staining index both the intensity of staining and the proportion of positively-stained cells were taken into consideration, using the following formula as previously reported [27]:

Fig. 1 Expression of sonic hedgehog (SHh) in bladder cancer. Immunohistochemical staining of bladder cancer tissue microarray using Shh antibody. a, b and c. No Shh expression. d, e and f. Moderate Shh expression. g, h and i. Strong Shh expression. Images were taken using different objectives (×10, ×20, ×40)
\[ I = 0^*f_0 + 1^*f_1 + 2^*f_2 + 3^*f_3 \], where I is the staining index, \( f_0-f_3 \) are the proportions of the cells showing a defined level of staining intensity (from 0 to 3). The staining index ranges from 0 to 300. For association analysis between the level of SHh expression and patients’ clinical and pathological parameters Fisher’s two-sided exact tests was applied. Correlation between SHh levels and cancer-specific survival, Kaplan–Meier curve was used and \( p \) value was calculated using the log-rank test.

### Statistical Analysis

Statistical analysis was performed using SPSS (version 21). Fisher’s tests were used to analyse the association between sonic hedgehog expression and clinicopathological features and results were considered significant for values of \( p < 0.05 \). Kaplan–Meier curve was used to evaluate the effect of Shh expression on overall patients’ survival; \( p \) value was calculated using the log-rank test.

### Results

#### Patient Characteristics

Our cohort subjects included 128 bladder cancer patients, 105 males (82%) and 23 females (18%). The age of our cohort ranged from 26 to 93 years with a mean value of 61 years. Tumour metastasis was detected in 14 patients and recurrence was observed in 35% of the tested group (Table 1).

| Clinicopathological parameters | Number | Sonic hedgehog expression | \( p \) value |
|--------------------------------|--------|---------------------------|-------------|
|                                |        | Low (%) | High (%) |        |
| Age group (years)              |        |         |         |        |
| < 60                           | 127    | 23 (45.1%) | 28 (54.9%) |        |
| ≥ 60                           |        | 41 (53.9%) | 35 (46.1%) |        |
| Marital status                 | 113    | 3 (50.0%) | 3 (50.0%) |        |
| Single                         |        | 3 (50.0%) | 3 (50.0%) |        |
| Married                        |        | 55 (51.4%) | 52 (48.6%) |        |
| Gender                         | 128    | 51 (48.6%) | 54 (51.4%) |        |
| Male                           |        | 14 (60.9%) | 9 (39.1%) |        |
| Female                         |        | 10 (41.7%) | 11 (43.9%) |        |
| Smoking                        | 65     | 23 (56.1%) | 18 (43.9%) |        |
| Yes                            |        | 14 (58.3%) | 10 (41.7%) |        |
| No                             |        | 14 (58.3%) | 10 (41.7%) |        |
| Type of cancer                 | 110    | 27 (46.6%) | 31 (53.4%) |        |
| MIBC                           |        | 28 (53.8%) | 24 (46.2%) |        |
| NMIBC                          |        | 27 (46.6%) | 31 (53.4%) |        |
| Histological grade            | 113    | 23 (46.0%) | 27 (54.0%) |        |
| Low grade                      |        | 34 (54.0%) | 29 (46.0%) |        |
| High grade                     |        | 34 (54.0%) | 29 (46.0%) |        |
| Family history                 | 65     | 6 (85.7%) | 1 (14.3%) |        |
| Yes                            |        | 30 (51.7%) | 28 (48.3%) |        |
| No                             |        | 30 (51.7%) | 28 (48.3%) |        |
| Lymph node status              | 83     | 43 (59.7%) | 29 (40.3%) |        |
| Negative                       |        | 2 (18.2%) | 9 (81.8%) |        |
| Positive                       |        | 2 (18.2%) | 9 (81.8%) |        |
| Metastasis                     | 80     | 6 (42.9%) | 8 (75.1%) |        |
| Positive                       |        | 40 (60.6%) | 26 (39.4%) |        |
| Negative                       |        | 40 (60.6%) | 26 (39.4%) |        |
| Recurrence                     | 127    | 23 (52.3%) | 21 (47.7%) |        |
| Yes                            |        | 41 (49.4%) | 42 (50.6%) |        |
| No                             |        | 41 (49.4%) | 42 (50.6%) |        |

\( ns \) not significant
Sonic Hedgehog Expression in Bladder Cancer

In order to analyze the expression pattern of sonic hedgehog protein in bladder cancer, immunohistochemical staining of tissue microarrays, containing core biopsies from 128 patients affected with bladder cancer, was performed using hedgehog-targeted antibody. Assessment of the staining pattern revealed that sonic hedgehog protein was predominantly localized in the cytoplasm of the bladder cancer cells, as illustrated in Fig. 1. Variation in intensity of sonic hedgehog expression in the bladder cancer specimens was scored as follows: 0 ($n=33$), 1 ($n=80$), 2 ($n=15$), 3 ($n=0$). The receiver operating characteristic (ROC) curve was used to determine the cutoff that can be used to discriminate between high and low sonic hedgehog expression. Forty nine percent (49%) of the analyzed cohort exhibited high cytoplasmic expression of sonic hedgehog. The intensity and distribution of nuclear staining was not reported as significant while membranous localization of Shh was not seen (Fig. 1).

Association between Sonic Hedgehog Expression and Clinicopathological Parameters

All 128 patients were included in the analysis. Correlation analysis of Shh staining was undertaken to examine the relationship between the protein levels of Shh and any of the patients’ clinicopathological features. Our data indicated that the expression of Shh is significantly associated with lymph node invasion in bladder cancer patients ($p = 0.02$). Statistical analysis indicated no correlation between the expression of Shh and other clinical parameters such as tumor grade, stage, smoking status or gender (Table 2). Similarly, there was no significant difference between negative and positive sonic hedgehog expression status and patients’ overall survival (Fig. 2).

Discussion

Sonic hedgehog is a member of hedgehog family of small secreted proteins, that were originally discovered as important regulator during vertebrates development [28]. It is well documented that Shh is expressed in normal bladder epithelium to maintain the regenerative potential of the epithelium and this expression exhibited different spatial and temporal distribution during abnormal bladder development indicating the important role of Shh in bladder tumorigenesis [30, 31]. Recent findings revealed that deregulation of sonic hedgehog pathway is associated with plethora of malignancies in various tissue-types through mutations in Patched (Ptc1) and/or the G protein-coupled receptor smoothened (SMO) genes [29, 32]. The potential oncogenic role of sonic hedgehog and the components of its signaling pathway on bladder pathogenesis is not well delineated. However several attempts have been made and reports indicated the involvement of Shh in bladder cancer growth and tumorigenicity [22, 33, 34]. Chen et al. (2010) undertook genotyping analysis on 177 single-nucleotide polymorphisms (SNP) using 803 bladder cancer cases and equal number of healthy controls and found that germ-line genetic variations in the Shh pathway predicted clinical outcomes of non-muscle-invasive bladder cancer patients receiving transurethral resection and BCG treatment [35]. In an independent study, Shin et al. 2011 demonstrated increased levels of Shh and Gli1 mRNAs in response to bladder tissue injury suggesting that Shh increase the proliferative potential of bladder stem cells. The latter deemed responsible for bladder cancer recurrence and drug resistance [36].

Our study showed high expression of Shh protein in the cytoplasmic compartment of bladder cancer using immunohistochemistry. The high expression was seen in almost 50% of the tested cohort. Interestingly, the expression of Shh protein significantly correlated with lymph node metastasis of tumours. This was consistent with previous findings by He et al. (2012) and Pignot et al. (2012), both authors reported that the levels of Shh, Ptc1 and Gli1 proteins were significantly elevated in bladder cancer tissues. Over-expression of these proteins correlated with a number of pathological parameters including lymph node metastasis [37, 38]. In an attempt to decipher the mechanism by which activation of Shh pathway mediate the tumorigenic potential and malignant phenotype in bladder cancer, Islam et al. (2016) performed a number of in-vitro and in-vivo experiments and revealed that Shh was able to induce the epithelial-to-mesenchymal

![Fig. 2](image-url) Overall survival of patients with bladder cancer. Kaplan-Meier curve showing no survival difference based on sonic hedgehog expression (log-rank $p = 0.85$)
phenotype (EMT) through activation of the transforming growth factor beta (TGF-β1) pathway [26]. It has recently been found that Shh plays an important role in urothelial bladder cancer by promoting bladder cancer stem cells self-renewal [39–41] which may also explain the elevated rates of tumour recurrence and resistance to conventional chemo/radiotherapies associated with bladder cancer. Involvement of sonic hedgehog in regulating bladder cancer stem cells activities could also add to the burden of the aggressive phenotype of bladder cancer, thus, targeting Shh pathway with curcumin or another chemopreventive agent might be an effective strategy for inhibiting bladder cancer development [42].

In conclusion, there is clear evidence that Shh signaling pathway is abnormally activated in several cancers including urothelial carcinoma of the bladder and the underlying mechanism is yet to be determined. This pathway represents a target for potential anti-cancer therapy and controlling recurrence in urothelial bladder carcinoma.

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Compliance with Ethical Standards

Competing Interests The authors do not have any conflicts of interest associated with this manuscript.

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