Novel Prognostic Models for Patients With Penile Carcinoma

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Abstract

Although penile carcinoma is a rare malignancy, there is still an unmet need to identify prognostic factors associated with poor survival. In this study, we utilized demographic and clinical information to identify the most informative variables associated with overall survival in patients with penile cancer. From a full model including all covariates found to be statistically significant in univariable analyses, we identified a parsimonious reduced model containing tumor site (penis glans: hazard ratio [HR] = 0.48; 95% CI: 0.28-0.85 and penis not otherwise specified: HR = 0.45; 95% CI: 0.25-0.84), undetermined tumor differentiation (HR = 0.48; 95% CI: 0.27-0.86), and TNM stage III/IV (HR = 2.83; 95% CI: 1.68-4.75). When all of the covariates from the full model were subjected to classification and regression tree analysis, we identified 6 novel risk groups. Of particular interest, we found marriage was associated with substantial improvement in survival among men with the same stage and disease site. Specifically, among single/widowed/divorced men with TNM stage 0-II and prepuce/penis corpus/overlapping lesions had worse survival (5-year survival = 18.2%) versus married men (5-year survival = 62.5%). Since marital status is linked to social support, these findings warrant a deeper investigation into the relationships between disease prognosis and social support in patients with penile carcinoma.

Keywords

penile cancer, penile cancer survival, prognostic factors, epidemiology, cancer survival

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Introduction

Penile cancer is rate accounting for less than 1% of cancers in men in the United States. In 2019, in the United States, approximately 2,080 new cases of penile carcinoma are expected to be diagnosed and 410 deaths are estimated to occur.¹ The overall 5-year relative survival rate for localized disease is 83% but drops to 48% with regional disease that has spread to the regional lymph nodes.² Squamous cell carcinoma (SCC), which presents as a painless lump, ulcer, or irregular mass,³ is the most common subtype comprising over 90% of all invasive penile carcinomas in the United States.¹ Although surgical resection with a partial or radical penectomy remains the gold standard therapy for advanced invasive penile carcinoma, the management of penile carcinoma has evolved in recent years resulting in a paradigm shift toward minimally invasive surgical treatment.⁴

Although penile cancer is rare and the overall prognosis is modestly optimistic, identifying prognostic factors is important to identify high-risk patients associated with poor outcomes. Previous studies have shown that presence and extent of regional inguinal lymph node metastases is an important prognostic factor for individuals with penile carcinoma.⁴⁻⁸ As a precursor to

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lymph node metastasis, the existence of lymphovascular invasion (LVI) has been shown to be associated with reduced survival among those without lymph node metastasis.⁹ Yet differing outcomes between studies warrant validation of the role of LVI as an independent prognostic indicator of reduced survival in patients with penile carcinoma.⁹ Studies have demonstrated clinicopathological variables, such as inguinal lymph node metastases, perineural invasion, tumor stage, and grade, to be predictors of poor survival¹⁰⁻¹²; however, there are socioeconomic and psychosocial factors that influence survival risk. Demographic and socioeconomic predictors, such as lower education attainment. African-American. and Hispanic ethnicity, have been shown to be associated with poor survival.¹³ Based on patient experiences including feelings of embarrassment and denial, there are psychosocial factors that may contribute to poor outcomes.⁴ In addition, men with penile cancer have expressed the importance of social support in coping with treatment and that social support helps to improve their mental well-being.¹⁴ There is a need for identifying men with penile cancer at high risk for poor survival, including assessment of sociodemographic factors with clinical factors. Thus, in this study, we utilized demographic and clinical data and a rigorous analytical approach to identify parsimonious models for identifying patients who are at risk of poor outcomes.

Patients and Methods

Study Population

This retrospective study included all 230 patients who were diagnosed with penile carcinoma at the H. Lee Moffitt Cancer Center & Research Institute from 1986 to 2013. There were no inclusion or exclusion criteria. Because penile cancer is a rare disease, this analysis included all available cases. The protocol for this study was approved by Advarra (IRB#Pro00014743; approval no. CR#00150095).

Cancer Registry Data

Patient data were obtained from electronic medical records and institutional databases including age at diagnosis, race/ethnicity, marital status, tobacco/alcohol use, and clinical covariates (tumor site, stage, histology, summary of first course of treatment, T stage, N stage, M stage, tumor size, surgical margins, and systemic surgery sequence). Pathologic staging was utilized and if these data were not available clinical staging was used. TNM staging was derived from the SEER Collaborative Staging which included site-specific factors SSF10 (involvement of corpus spongiosum or corpus cavernosum), SSF17 (extranodal extension of regional lymph nodes), and tumor size, extension, lymph nodes, and metastases. Moffitt's Cancer Registry collects vital status through active (ie, chart review and contacting patients, relatives, and other medical providers) and passive methods (ie, mortality records). Summary of first course of treatment was defined as all treatment methods recorded in the treatment plan and administered to the patient before disease progression, recurrence, or death. Tobacco use was categorized as self-reported ever, never, or unknown. Alcohol use was categorized as self-reported ever or never drinker. Overall survival was the dependent variable and was defined as date of diagnosis to date of death or last follow-up and was right censored at 5 years.

Statistical Analysis

The log-rank test was used to test for differences in overall survival (censored at 12, 36, and 60 months) by demographic and clinical variables. We utilized a model building approach to identify the most informative demographic and clinical variables associated with overall survival. Overall survival was defined as date of cancer diagnosis to date of event or date of last follow-up. Among patients without an event, censoring occurred at either 5-years or date of last follow-up if it was less than 5 years.

Univariable Cox proportional hazards regression models were used to identify which demographic and clinical variables were significantly associated with overall survival. All covariates that were found to be statistically significant in the univariable analyses were then included into a single full multivariable model. To reduce the full model to the most informative covariates, 2 separate analyses were conducted: backward elimination analysis and classification and regression tree (CART) analysis. Using the covariates from the full model, a backward elimination approach was utilized using 0.05 as the pre-specified P value for removal from the model. Classification and regression tree analysis, adapted for survival time, was also employed to identify novel patient risk groups (nodes) based on the most informative covariates from the full models for model 1 based on TNM group stage and for model 2 based on T, N, and M stages. Classification and regression tree is a nonparametric tree-building technique that can reveal complex interactions between predictors and the outcome of interest (survival).¹⁵ Survival curves of the risk groups identified by CART were estimated using Kaplan-Meier method, and differences in survival time were tested for significance using the log-rank test. Receiver operating characteristics (ROC) curves were used to assess performance of TNM stage alone versus the CART models. P value <.05 was considered statistically significant. All statistical analyses were performed using Stata/ MP 14.2 (StataCorp LP).

Results

Survival Time by Demographic and Clinical Characteristic

Overall survival rates, censored at 12, 36, and 60 months, were compared by demographic and clinical variables (Table 1). There were significant differences for the survival rates by marital status, tumor site, differentiation, stage of disease, T stage, N stage, M stage, tumor size, surgical margins, and systemic/surgery sequence. Specifically, single (61.8%) and widowed/divorced/unknown (45.4%) had lower 5-year

Reyes et al

| Characteristics | N (%) | 12 months % | 36 months % | 60 months % |
|---|-------------------------------------|--------------------------|--------------------------|---------------|
| Year of diagnosis | | | | |
| <2005 | 112 (48.7) | 86.5 | 68.3 | 59.9 |
| | 118 (51.3) | 92.6 | 75.9 | 66.0 |
| P value | | .207 | .288 | .291 |
| Age at diagnosis, years | | | | |
| Mean (SD), 60 (13.7) | | | | |
| <55 | 75 (32.6) | 94.2 | 70.8 | 67.1 |
| 56-69 | 92 (40.0) | 83.5 | 65.0 | 63.4 |
| >70 | 63 (27.4) | 91.8 | 81.2 | 55.6 |
| P value | | .103 | .107 | .785 |
| Race | | | | |
| White | 211 (91.7) | 88.3 | 71.0 | 62.2 |
| Other | 19 (8.3) | 100.0 | 73.3 | 73.3 |
| P value | | .144 | .495 | .370 |
| Ethnicity | | | | |
| Non-Hispanic | 199 (86.9) | 90.3 | 73.5 | 65.3 |
| Hispanic | 30 (13.1) | 81.9 | 57.6 | 46.1 |
| Unknown | 1.0 (0.4) | 0.0 | 0.0 | 0.0 |
| P value | () | .362 | .179 | .101 |
| Marital status | | | , | |
| Married | 149 (64 8) | 90.8 | 76.8 | 68.7 |
| Single | 32 (13.9) | 93 | 67.9 | 61.8 |
| Widowed/divorced/missing | 49 (21 3) | 82.5 | 56.8 | 45.4 |
| P value | 17 (21.3) | 308 | 061 | 029 |
| | | .500 | .001 | .027 |
| Never | 121 (52.6) | 89.4 | 77 0 | 71.2 |
| Ever | 93 (40 4) | 88.2 | 65.9 | 56.4 |
| Missing | 16 (7 0) | 93.3 | 71 1 | 53.3 |
| P value | 10 (7.0) | 788 | 292 | 168 |
| | | ./00 | .272 | .100 |
| Never/unknown | 74 (32 2) | 90.2 | 72 4 | 62.2 |
| Ever | 156 (67.8) | 88.7 | 70.9 | 63 1 |
| P value | 130 (07.0) | 785 | 867 | 946 |
| Tumor site | | ./05 | .007 | .710 |
| Prepuse/penis corpus/overlapping lesion | 46 (20.0) | 81.7 | 53.2 | 41.6 |
| Penis glans | 94 (40 9) | 90.9 | 72.9 | 66.5 |
| Penis NOS | 90 (39 1) | 914 | 80 1 | 71.2 |
| P value | 70 (37.1) | 186 | 013 | 005 |
| Histology | | .100 | .015 | .005 |
| Squamous cell carcinoma | 204 (99 7) | 99 4 | 70.7 | 63 1 |
| Nonsquamous cell carcinoma | 207 (00.7) | 07. 1 97.9 | 70.7 75 A | 617 |
| P value | 20 (11.5) | 884 | 73. 4 674 | 988 |
| Tumor differentiation | | .004 | .074 | .700 |
| Poorly and undifferentiated | 38 (14 5) | 79 | 41.2 | 41.2 |
| Moderately and well differentiated | 94 (40 9) | 95 A | 42.5 | 56.4 |
| Indetermined | | 94.4 | 02.5 | 75 9 |
| P volue | 70 (42.0) | 70.0 00E | 67.3 < 001 | / 3.7 |
| TVIM staging | | .005 | \. 001 | \.001 |
| ^ | (0 (20 () | 00.2 | 5 50 | 77.0 |
| 0 | 60 (27.6) 62 (27.4) | 70.3 | 72.7 | 77.0 |
| 1 | 03 (27. 4) 29 (17.0) | 71.0 05.2 | 72.0 | 70.4 |
| 2 | 37 (17.0) | 82 A | /1.7 <u>/</u> 10 | ۵U.۶ ۸۱۹ |
| 5 A | J7 (10.1) | 50.0 | דו.7 <u>24</u> א | 11.7 74 A |
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| | 50 (32.7) | /4.7 | JÖ.۷ م ا | 38.2 |
| r value | | .010 | \.UUI | <.UUI |

(continued)

Table I. (continued)

| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | Characteristics | N (%) | 12 months % | 36 months % | 60 months % |
|---|--|---------------------|--------------|-------------|-------------|
| $\begin{array}{l c c c c c } \hline Or ClS & 62 (26.9) & 98.1 & 95.9 & 63.1 \\ c c c c c c c c c c c c c c c c c c $ | Pathologic T stage | | | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | Overall | | | | |
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| IA 9 (3.9) 100.0 100.0 100.0 IB 8 (3.5) 85.7 85.7 0.0 2 34 (14.8) 81.6 65.3 58.0 3 A 1 (0.4) 100.0 100.0 100.0 4 4 (1.7) 50.0 50.0 50.0 4A 1 (0.4) 100.0 100.0 00.0 0.0 Missing 19 (8.3) 100.0 71.3 44.4 Collapsed 0/CIS 62 (2.9) 98.1 95. 66.5 64.3 1 72 (31.3) 89.5 66.5 64.3 35.2 44.4 5.2.2 60.0 60.0 30.0 30.0 30.0 30.0 30.0 30.0 30.0 30.0 30.0 30.0 30.0 31.4 44.4 5.2.2 6.00 60.0 30.0 30.0 31.4 44.4 5.2.2 6.00 60.0 30.0 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 72.9 <td>1</td> <td>55 (23.9)</td> <td>88.7</td> <td>62.9</td> <td>60.8</td> | 1 | 55 (23.9) | 88.7 | 62.9 | 60.8 |
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| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | IB | 8 (3.5) | 85.7 | 85.7 | 0.0 |
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| 4 4 (1.7) 50.0 50.0 50.0 Mising 19 (8.3) 100.0 71.3 44.4 Collapsed 72 (31.3) 89.5 66.5 64.3 1 72 (31.3) 89.5 66.5 64.3 2 34 (14.8) 81.6 65.3 58.0 3 38 (16.5) 79.5 47.6 47.6 4 5 (2.2) 60.0 60.0 30.0 Mising 19 (8.3) 100.0 71.3 44.4 Pathologic N stage 002 <001 | 3A | l (0.4) | 100.0 | 100.0 | 100.0 |
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| Pathologic M stage0213 (92.6)88.872.264.812 (0.9)50.00.00.0Missing15 (6.5)100.071.448.9P value.033<.001 | P value | | <.001 | <.001 | <.001 |
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| Missing15 (6.5)100.071.448.9P value.033<.001 | I | 2 (0.9) | 50.0 | 0.0 | 0.0 |
| P value .033 <.001 <.001 Tumor size, mm Mean (SD), 29.8 (22.3) <25 (median) | Missing | 15 (6.5) | 100.0 | 71.4 | 48.9 |
| Tumor size, mm Mean (SD), 29.8 (22.3) <25 (median) | P value | | .033 | <.001 | <.001 |
| Mean (SD), 29.8 (22.3)<25 (median) | Tumor size, mm | | | | |
| <25 (median)43 (40.6)94.377.571.9≥2563 (59.4)83.447.945.2Missing124 (53.9)90.780.368.7P value.131<.001 | Mean (SD), 29.8 (22.3) | | | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | <25 (median) | 43 (40.6) | 94.3 | 77.5 | 71.9 |
| Missing124 (53.9)90.780.368.7P value.131<.001 | >25 | 63 (59.4) | 83.4 | 47.9 | 45.2 |
| P value.131<.001<.001Surgical marginNo primary surgery70 (30.4)86.658.747.5No residual tumor114 (49.6)91.477.568.8Microscopic/macroscopic/residual tumor, NOS14 (6.1)85.761.261.2Not evaluable/missing32 (13.9)90.081.676.8P value.676.031.015Summary of first course treatment.72 (74.8)89.973.863.9Combination/other58 (25.2)87.264.061.1P value.473.088.221Systemic and/or no surgery89 (38.7)93.674.966.3Surgery + neoadjuvant35 (15.2)96.981.775.8Surgery + neoadjuvant4 (1.7)100.00.00.0Surgery + neoadjuvant + adjuvant2 (0.9)50.050.00.0Missing100 (43.5)85.066.857.4P value.045.345.316 | Missing | 124 (53.9) | 90.7 | 80.3 | 68.7 |
| Surgical margin No primary surgery 70 (30.4) 86.6 58.7 47.5 No residual tumor 114 (49.6) 91.4 77.5 68.8 Microscopic/macroscopic/residual tumor, NOS 14 (6.1) 85.7 61.2 61.2 Not evaluable/missing 32 (13.9) 90.0 81.6 76.8 P value .676 .031 .015 Summary of first course treatment .676 .031 .015 Surgery only 172 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 .00 .00 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 .00 <tr< td=""><td>P value</td><td>()</td><td>.131</td><td><.001</td><td><.001</td></tr<> | P value | () | .131 | <.001 | <.001 |
| No primary surgery70 (30.4)86.658.747.5No residual tumor114 (49.6)91.477.568.8Microscopic/macroscopic/residual tumor, NOS14 (6.1)85.761.261.2Not evaluable/missing32 (13.9)90.081.676.8 P value.676.031.015Summary of first course treatment | Surgical margin | | | | |
| No primery origery 114 (49.6) 91.4 77.5 68.8 Microscopic/macroscopic/residual tumor, NOS 14 (6.1) 85.7 61.2 61.2 Not evaluable/missing 32 (13.9) 90.0 81.6 76.8 P value .676 .031 .015 Summary of first course treatment .676 .031 .015 Surgery only 172 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | No primary surgery | 70 (30 4) | 86.6 | 587 | 47 5 |
| Microscopic/macroscopic/residual tumor, NOS 14 (6.1) 85.7 61.2 61.2 Not evaluable/missing 32 (13.9) 90.0 81.6 76.8 P value .676 .031 .015 Surgery only 172 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic/surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | No residual tumor | 114 (49.6) | 914 | 77 5 | 68.8 |
| Not evaluable/missing 32 (13.9) 90.0 81.6 76.8 P value .676 .031 .015 Summary of first course treatment .676 .031 .015 Surgery only 172 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic/surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 .345 | Microscopic/macroscopic/residual tumor NOS | 14 (6 1) | 85.7 | 61.2 | 61.2 |
| P value .676 .031 .015 Surgery only .676 .031 .015 Surgery only .676 .031 .015 Combination/other .58 (25.2) .87.2 .64.0 .61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Surgery + adjuvant .35 (15.2) 96.9 .81.7 .75.8 Surgery + neoadjuvant .4 (1.7) .00.0 .0.0 .0.0 Surgery + neoadjuvant + adjuvant .2 (0.9) .50.0 .50.0 .0.0 Missing .00 (43.5) .85.0 .66.8 .57.4 P value .045 .345 .316 | Not evaluable/missing | 32 (13 9) | 90.0 | 81.6 | 76.8 |
| F value 1.010 1.011 Surgery only 172 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic/surgery sequence .550 96.9 81.7 75.8 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | P voluo | 52 (15.7) | 70.0 474 | 01.0 | 0.0 |
| Surgery only I72 (74.8) 89.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | r value | | .070 | .031 | .015 |
| Surgery only 172 (74.8) 87.9 73.8 63.9 Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Summary of first course treatment | 172 (74 0) | 00.0 | 72.0 | (2.0 |
| Combination/other 58 (25.2) 87.2 64.0 61.1 P value .473 .088 .221 Systemic/surgery sequence .473 .088 .221 No systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Surgery only | 172 (74.8) | 89.9 | / 3.8 | 63.7 |
| P value .4/3 .088 .221 Systemic/surgery sequence .088 .221 No systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Combination/other | 58 (25.2) | 87.2 | 64.0 | 61.1 |
| Systemic/surgery sequence No systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | P value | | .4/3 | .088 | .221 |
| No systemic and/or no surgery 89 (38.7) 93.6 74.9 66.3 Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Systemic/surgery sequence | | | | |
| Surgery + adjuvant 35 (15.2) 96.9 81.7 75.8 Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | No systemic and/or no surgery | 89 (38.7) | 93.6 | 74.9 | 66.3 |
| Surgery + neoadjuvant 4 (1.7) 100.0 0.0 0.0 Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Surgery + adjuvant | 35 (15.2) | 96.9 | 81.7 | 75.8 |
| Surgery + neoadjuvant + adjuvant 2 (0.9) 50.0 50.0 0.0 Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Surgery + neoadjuvant | 4 (1.7) | 100.0 | 0.0 | 0.0 |
| Missing 100 (43.5) 85.0 66.8 57.4 P value .045 .345 .316 | Surgery + neoadjuvant + adjuvant | 2 (0.9) | 50.0 | 50.0 | 0.0 |
| P value .045 .345 .316 | Missing | 100 (43.5) | 85.0 | 66.8 | 57.4 |
| | P value | · · / | .045 | .345 | .316 |

Abbreviations: NOS, not otherwise specified; SD, standard deviation. ^aBold font indicates statistically significant *P* values ^bTNM stages: Unknown values were excluded from the analysis.

Table 2. Univariable Analysis.^{a,b}

| Characteristics | HR (95% CI) |
|---|-------------------|
| Year of diagnosis | |
| ≤2005 [°] | Ref. |
| ≥2006 | 0.77 (0.47-1.26) |
| Age at diagnosis | |
| ≤55 | Ref. |
| 56-69 | 1.22 (0.69-2.16) |
| ≥70 | 1.12 (0.60-2.08) |
| Race | |
| White | Ref. |
| Other | 0.59 (0.19-1.89) |
| Ethnicity | |
| Non-Hispanic ^c | Ref. |
| Hispanic | 1.91 (1.04-3.50) |
| Marital status | |
| Married | Ref. |
| Single/widowed/divorced/missing | 1.71 (1.06-2.75) |
| Alcohol use | |
| Never | Ref. |
| Ever | 1.60 (0.97-2.65) |
| Missing | 1.59 (0.65-3.87) |
| Tobacco use | |
| Never/unknown | Ref. |
| Ever | 0.98 (0.59-1.62) |
| Tumor site | |
| Prepuce/penis corpus/overlapping lesion | Ref. |
| Penis glans | 0.49 (0.28-0.87) |
| Penis, NOS | 0.41 (0.22-0.74) |
| Histology | |
| Squamous cell carcinoma | Ref. |
| Nonsquamous cell carcinoma | 0.99 (0.49–2.01) |
| Tumor differentiation | |
| Poorly and undifferentiated | Ref. |
| Moderately and well differentiated | 0.62 (0.34-1.13) |
| Undetermined | 0.25 (0.13-0.49) |
| TNM stage group | |
| 0/CIS/missing | Ref. |
| 1/11 | 1.62 (0.86-3.03) |
| III/IV | 4.70 (2.48-8.93) |
| Tumor size, mm (median) | |
| <25 | Ref. |
| ≥ 25 | 2.91 (1.32-6.39) |
| Missing | 1.18 (0.54-2.55) |
| Surgical margins | |
| No primary surgery | Ref. |
| No residual tumor | 0.49 (0.29-0.82) |
| Microscopic/macroscopic/residual, NOS | 0.85 (0.30-2.42) |
| Not evaluable/missing | 0.36 (0.15-0.86) |
| T stage (collapsed) | |
| 0/CIS | Ref. |
| | 3.05 (1.30-7.18) |
| 2 | 3.8/ (1.55-9.71) |
| 3 | 5.61 (2.30-13.67) |
| 4 | 9.17 (2.37-35.53) |
| Missing | 4.48 (1.67-12.04) |
| N stage | . . |
| U . | Kef. |
| | 1.07 (0.26-4.45) |
| 2 | 4.09 (2.11-7.91) |
| | |

(continued)

 Table 2. (continued)

| Characteristics | HR (95% CI) |
|---|--------------------|
| 3 | 9.93 (4.36-22.64) |
| Missing | 1.68 (0.83-3.36) |
| M stage | |
| 0 | Ref. |
| I | 15.75 (3.65-67.96) |
| Missing | 1.08 (0.61-1.91) |
| Summary of first course treatment | |
| Surgery only | Ref. |
| Combination/other | 1.39 (0.82-2.39) |
| Systemic/surgery sequence | |
| No systemic treatment and/or no surgery | Ref. |
| Systemic after surgery | 0.73 (0.29-1.83) |
| Systemic before surgery | 2.25 (0.30-17.0) |
| Systemic before and after surgery | 2.66 (0.36-19.9) |
| Missing | 1.46 (0.85-2.51) |
| Year of first contact | |
| ≤ 2005 | Ref. |
| ≥2006 | 0.71 (0.43-1.16) |

Abbreviations: CI, confidence interval; HR, hazards ratio; NOS, not otherwise specified; Ref., reference.

^aN = 230.

^bBold font indicates a statistically significant HR.

^cNon-Hispanic, includes one unknown.

survival rates compared to married men (68.7%; log-rank P value = .029). By site, there were significant differences by 5-year survival between those with prepuce/penis corpus/overlapping lesions (41.6%) compared to other penile tumor sites, penis glans (66.5%), and penis nos (71.2%; log-rank P value = .005). For tumor differentiation, significant differences were observed between poorly and undifferentiated (41.2%) compared to moderately and well-differentiated (56.4%), and undetermined (75.9%) at 5 years (log-rank P value < .001). For TNM staging, there were significant differences at 5 years (log-rank P value < .001) for TNM stage 0 (77.0%), 1 (70.4%), 2 (60.9%), 3 (41.9%), 4 (26.4%), and unknown (55.6%). For pathologic T stage (collapsed), pathologic N stage, and pathologic T stage there were also significant differences for 5-year survival (log-rank P value < .001) and significant differences for 5-year survival for tumor size <25 mm (71.9%) versus ≥ 25 mm (45.2%; log-rank P value < .001). For surgical margins, no primary surgery (47.5%), no residual tumor (68.8%), microscopic/macroscopic/residual tumor, not otherwise specified (NOS; 61.2%), and not evaluable/missing (76.8%) differed significantly for 5-year survival (log-rank P value = .015).

Univariable Analyses

As presented in Table 2, statistically significant univariable hazard ratios (HRs) were observed for Hispanics (HR = 1.91, 95% CI = 1.04-3.50), single/widowed/divorced (HR = 1.71, 95% CI = 1.06-2.75), tumor site (penis glans: HR = 0.49, 95% CI = 0.28-0.87 and penis NOS: HR = 0.41, 95% CI = 0.22-0.74), for undetermined tumor differentiation (HR = 0.25,

95% CI = 0.13-0.49), TNM stage III/IV (HR = 4.70, 95% CI = 2.48-8.93), tumor size \geq 25 mm (HR = 2.91, 95% CI = 1.32-6.39), surgical margins (HR = 0.49, 95% CI = 0.29-0.82), pathologic T stage (collapsed) for stage 1 (HR = 3.05, 95% CI = 1.30-7.18), 2 (HR = 3.87, 95% CI = 1.55-9.71), 3 (HR = 5.61, 95% CI = 2.30-13.67), 4 (HR = 9.17, 95% CI = 2.37-35.53), missing (HR = 4.48, 95% CI = 1.67-12.04), pathologic N stage for stage 2 (HR = 4.09, 95% CI = 2.11-7.91), 3 (HR = 9.93, 95% CI = 4.36-22.64), and pathologic M stage for those positive for metastasis (HR = 15.75, 95% CI = 3.65-67.96).

Full and Reduced Models

The full multivariable models included all covariates found to be statistically significant from the univariable analyses (Table 3). However, model 1 considered TNM group stage (and excluded T stage, N stage, and M stage), while model 2 considered pathologic T stage, N stage, and M stage (and excluded TNM group stage). In the full model for model 1, the only statistically significant covariates were ever alcohol use (HR = 1.80; 95% CI: 1.06-3.05), tumor site (penis glans: HR = 0.53; 95% CI: 0.29-0.96, and penis NOS: HR = 0.42; 95% CI: 0.22-0.80, undetermined tumor differentiation (HR = 0.40; 95% CI: 0.18-0.91), and TNM stage III/IV (HR = 2.64; 95% CI: 1.11-6.30). The full model 1 was subjected to a backward elimination approach to yield a parsimonious model which contained tumor site (penis glans: HR = 0.48; 95% CI: 0.28-0.85 and penis NOS: HR = 0.45; 95% CI: 0.25-0.84), undetermined tumor differentiation (HR = 0.48; 95%CI: 0.27-0.86), and TNM stage III/IV (HR = 2.83; 95% CI: 1.68-4.75).

In the full model for model 2, statistically significant covariates were ever alcohol use (HR = 2.04, 95% CI = 1.19-3.50), tumor site (penis NOS: HR = 0.35, 95% CI = 0.17-0.72), pathologic T stage 4 (HR = 6.44, 95% CI = 1.46-28.38), and N stage (2: HR = 2.93, 95% CI = 1.36-6.33; 3: HR = 12.6, 95% CI = 4.01-39.56). Backward elimination resulted in a parsimonious model containing single/widowed/divorced/missing (HR = 1.70, 95% CI = 1.05-2.76), ever alcohol use (HR = 1.81, 95% CI = 1.08-3.03), penis NOS (HR = 0.54, 95% CI = 0.31-0.94), undetermined tumor differentiation (HR = 0.52, 95% CI = 0.28-0.96), \geq 25 mm tumor size (HR = 1.99, 95% CI = 1.18-3.36), pathologic T stage 4 (HR = 6.28, 95% CI = 1.83-21.55), and pathologic N stage 2 (HR = 2.90, 95% CI = 1.48-5.68) and stage 3 (HR = 13.17, 95% CI = 5.28-32.89).

Classification and Regression Tree Analysis

When all the covariates from the full model 1 were subjected to CART analysis, we identified 6 risk groups (Table 4 and Figure 1A-B) based on 4 covariates. Classification and regression tree model 1 included tumor stage, tumor site, surgical margins, and marital status. Compared to risk group 1 (stage 0-II + penis glans/NOS + macroscopic/microscopic/residual tumor, NOS), men in risk group 4 (stage 0-II, prepuce/penis

corpus/overlapping lesion, single/widowed/divorced) exhibited a 6.5-fold increased risk of death (HR = 6.51; 95% CI = 2.86-14.81). Men in risk group 6 (stages III/IV + prepuce/penis corpus/overlapping lesion + macroscopic/microscopic/residual tumor, NOS) had an elevated risk of death (HR = 10.93; 95% CI = 5.11-23.36) compared to risk-group 1 (HR = 1.00).

When all the covariates from the full model 2 were subjected to CART analysis, we identified 6 risk groups based on 4 covariates. The 4 covariates included in CART model 2 were N stage, tumor differentiation, site, and marital status. Risk group 3 (N stage 0-1 + moderately/well-differentiated/ undetermined + prepuce/penis corpus/overlapping lesion + single/widowed/divorced) had an increased risk of dying (HR = 4.39; 95% CI = 1.76-10.94). Those in risk group 4 (N stage 0 - 1 + poorly and undifferentiated tumors) had an increased risk of dying (HR = 3.79; 95% CI = 1.72-8.35) compared to risk group 1 (HR = 1.00). Risk group 6 (N stage 2-3) had the greatest increase in risk (HR = 7.82; 95% CI = 4.11-14.85).

When CART risk groups for both models (Table 4) were adjusted for alcohol use, ethnicity, and tumor differentiation, the point estimates were slightly somewhat attenuated, but the overall trend was the same. For risk groups identified in CART model 1 (Figure 1A), men with stage 0-II disease and were married (risk group 3) had better survival than single/divorced/widowed (risk group 4; 62.5% 5-year survival and HR = 2.19, 95% 0.89-5.36 for risk group 3 vs 18.0% 5-year survival and HR = 5.29, 95% CI: 2.17-12.90 for risk group 4). For risk groups identified in CART model 2 (Figure 1B), men with N stage 0-1 and married (risk group 2) had better survival than single/divorced/widowed (risk group 3; 67.7% 5-year survival and HR = 1.63, 95% CI: 0.61-4.37 vs 34.3% 5-year survival and HR = 3.61, 95% CI: 1.38-9.47).

Model Performance

Receiver operating characteristics curves (Figure 2) were used to assess performance of TNM stage alone versus the models and revealed both CART model 1 (ROC = 0.722) and CART model 2 (ROC = 0.709) were better predictors than TNM stage alone (ROC = 0.643). Likewise, reduced model 1(AUC = 0.713) and 2 (AUC = 0.744) performed better than TNM stage alone (Table 5).

Discussion

This analysis of a cohort of patients diagnosed with penile cancer identified novel prognostic models containing the most informative covariates significantly associated with overall survival. The most informative covariates using a backward elimination approach were found to be tumor site, tumor differentiation, tumor size, TNM stage, pathologic N stage, and marital status. However, after using a decision tree approach (CART), novel patient risk groups were identified for 2 separate CART models and both showed a beneficial impact on survival among married men. Specifically, among men with

Table 3. Univariable, Multivariable, Reduced Model Analyses.^a

| $ \begin{array}{c c c c c c c c c c c c c c c c c c c $ | | Univariable ^b HR (95% CI) | Full multivariable model I HR (95% CI) | Reduced multivariable model I HR (95% CI) | Full multivariable model 2 HR (95% CI) | Reduced multivariable model 2 |
|---|-----------------------------|---|--|--|--|-------------------------------------|
| | Characteristics | | | | | HR (95% CI) |
| Non-Hispanic ⁴ Ref. | Ethnicity | | | | | |
| Hispanic 1.91 (1.04-3.50) 1.81 (0.94-3.49) - 1.61 (0.80-3.26) - Marital status Ref. R | Non-Hispanic ^c | Ref. | Ref. | Ref. | Ref. | Ref. |
| Mantial statuts Married Ref. Ref. <td>Hispanic</td> <td>1.91 (1.04-3.50)</td> <td>1.81 (0.94-3.49)</td> <td>-</td> <td>1.61 (0.80-3.26)</td> <td>_</td> | Hispanic | 1.91 (1.04-3.50) | 1.81 (0.94-3.49) | - | 1.61 (0.80-3.26) | _ |
| Married Ref. | Marital status | . , | . , | | | |
| Single/widewed/divorced/ missing I.71 (1.06-2.75) I.52 (0.91-2.50) - I.51 (0.88-2.60) I.70 (1.05-2.76) Alcohol use Never Ref. | Married | Ref. | Ref. | Ref. | Ref. | Ref. |
| missing Alcohol use Ref. Ref. </td <td>Single/widowed/divorced/</td> <td>1.71 (1.06-2.75)</td> <td>1.52 (0.91-2.50)</td> <td>-</td> <td>1.51 (0.88-2.60)</td> <td>1.70 (1.05-2.76)</td> | Single/widowed/divorced/ | 1.71 (1.06-2.75) | 1.52 (0.91-2.50) | - | 1.51 (0.88-2.60) | 1.70 (1.05-2.76) |
| Akcohol use Never I. Ref. Ref. Ref. Ref. Ref. Ref. Ref. Ref | missing | . , | | | | . , |
| Never Ref. < | Alcohol use | | | | | |
| Ever 1.60 ($0.97-2.65$) 1.80 ($1.06-3.05$) - 2.26 ($1.19-3.50$) 1.81 ($1.08-3.03$) Tumor site Prepue2/penis corpus/ overtapping lesion Ref. | Never | Ref. | Ref. | Ref. | Ref. | Ref. |
| $\begin{array}{l c c c c c c c c c c c c c c c c c c c$ | Ever | 1.60 (0.97-2.65) | 1.80 (1.06-3.05) | _ | 2.04 (1.19-3.50) | 1.81 (1.08-3.03) |
| Tumor site Prepue/pensito corpus/ overlapping lesion Ref. Obsta (0.32-0.48) 0.35 (0.12-0.72) 0.54 (0.31-0.94) Tumor differentiated Deorly and undifferentiated Ref. | Missing | 1.59 (0.65-3.87) | 2.14 (0.83-5.53) | _ | 2.56 (0.98-6.72) | 2.81 (1.11-7.12) |
| Prepuez/penis corpus/ overlapping lesion Ref. Ref.< | Tumor site | | | | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | Prepuce/penis corpus/ | Ref. | Ref. | Ref. | Ref. | Ref. |
| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | Penis glans | 0 49 (0 28-0 87) | 0 53 (0 29-0 96) | 0 48 (0 28-0 85) | 0.58 (0.30-1.12) | _ |
| Non-residual | Penis NOS | 0.41 (0.22 - 0.07) | 0.33(0.27-0.70) 0.42(0.22-0.80) | 0.45 (0.25-0.83) | 0.35 (0.17-0.72) | 0 54 (0 31-0 94) |
| No number of the second seco | Tumor differentiation | 0.11 (0.22 0.7 1) | 0.12 (0.22 0.00) | 0.15 (0.25 0.01) | 0.55 (0.17 0.72) | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | Poorly and undifferentiated | Ref | Ref | Ref | Ref | Ref |
| Undetermined Undetermined 0.25 (0.13-0.49) 0.40 (0.18-0.91) 0.48 (0.27-0.86) 0.46 (0.19-1.12) 0.52 (0.28-0.96) TNM stage group 0/ClS/missing Ref. Ref. Ref. Nef. | Moderately and well | 0.62 (0.34-1.13) | 0.63 (0.33-1.22) | - | 0.64 (0.32-1.29) | _ |
| Ontechning Crist (crist(crist)) Crist (crist(crist)) Crist (crist(crist)) Crist (crist(crist)) DVCIS/missing Ref. Ref. Ref. Ref. I//I 1.62 (0.86-3.03) 1.05 (0.48-2.31) - III//V 4.70 (2.48-8.93) 2.64 (1.11-6.30) 2.83 (1.68-4.75) Tumor size (mm), median <25 | Indetermined | 0 25 (0 13-0 49) | 0 40 (0 18-0 91) | 0 48 (0 27-0 86) | 0.46(0.19-1.12) | 0 52 (0 28-0 96) |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | TNM stage group | 0.23 (0.13-0.47) | 0.40 (0.10-0.71) | 0.40 (0.27-0.00) | 0.17-1.12) | 0.52 (0.20-0.70) |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | 0/CIS/missing | Pof | Pof | Pof | | |
| 11 1.03 (0.005.30) 1.03 (0.005.31) - 111//V 4.70 (2.48-8.93) 2.64 (1.11-6.30) 2.83 (1.68-4.75) Tumor size (mm), median - - 2.23 (0.94-5.31) 1.99 (1.18-3.36) 25 2.91 (1.32-6.39) 1.76 (0.76-4.04) - 2.23 (0.94-5.31) 1.99 (1.18-3.36) Missing 1.18 (0.54-2.55) 1.42 (0.64-3.15) - 1.48 (0.63-3.50) - Surgical margins No residual tumor 0.49 (0.29-0.82) 1.01 (0.55-1.88) - 0.98 (0.49-1.93) - Microscopic/macroscopic/ 0.85 (0.30-2.42) 1.38 (0.46-4.15) - 1.40 (0.44-4.43) - residual, NOS Not evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - T stage (collapsed) 0/CIS Ref. Ref. Ref. Ref. 0/CIS Ref. Ref. Ref. Ref. - - 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) - - - - - - - - - - - - - - - | | | | itel. | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | | 4 70 (2 48-8 93) | 2.64(1.11-6.30) | - 2 83 (1 68-4 75) | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | Tumor size (mm) median | 4.70 (2.40-0.75) | 2.04 (1.11-0.30) | 2.03 (1.00-4.73) | | |
| $\begin{array}{c c c c c c c c c c c c c c c c c c c $ | <25 | Rof | Rof | Rof | Rof | Rof |
| 2.5 2.7 (1.526.37) 1.70 (0.764.07) - 2.23 (0.74.35) 1.70 (1.763.05) Missing 1.18 (0.54-2.55) 1.42 (0.64-3.15) - 1.48 (0.63-3.50) - Surgical margins No primary surgery Ref. Ref. Ref. Ref. Ref. No residual tumor 0.49 (0.29-0.82) 1.01 (0.55-1.88) - 0.98 (0.49-1.93) - Microscopic/macroscopic/ 0.85 (0.30-2.42) 1.38 (0.46-4.15) - 1.40 (0.44-4.43) - residual, NOS Not evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - T stage (collapsed) 0/CIS Ref. Ref. Ref. Ref. 0/CIS Ref. 1.53 (0.54-4.34) - - 1 3.05 (1.30-7.18) 1.53 (0.54-6.29) - - 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) - - 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) - - 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) N stage 0 Ref. Ref. | >25 | 2 91 (1 32-6 39) | 1.76 (0.76 4.04) | | 2 22 (0 94-5 31) | |
| Ning Ning (0.174.35) Ning (0.174.35) Ning (0.174.35) Surgical margins No residual tumor 0.49 (0.29-0.82) 1.01 (0.55-1.88) – 0.98 (0.49-1.93) – Microscopic/macroscopic/ 0.85 (0.30-2.42) 1.38 (0.46-4.15) – 0.98 (0.44-4.43) – No residual tumor 0.49 (0.29-0.82) 1.01 (0.55-1.88) – 0.98 (0.44-4.43) – Microscopic/macroscopic/ 0.85 (0.30-2.42) 1.38 (0.46-4.15) – 1.40 (0.44-4.43) – Not evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) – 0.86 (0.31-2.38) – T stage (collapsed) 0/CLS Ref. Ref. Ref. Ref. 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) – – 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) – 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) – 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) – N stage 0 Ref. Ref. Ref. 1 1.07 | ≥25 Missing | 1 18 (0.54.2.55) | 1.70 (0.70-4.04) | _ | 1.48 (0.63-3.50) | - |
| No primary surgery Ref. R | Surgical margins | 1.10 (0.31 2.33) | 1.12 (0.01 5.15) | | 1.10 (0.05 5.50) | |
| No residual tumor 0.49 (0.29-0.82) 1.01 (0.55-1.88) - 0.98 (0.49-1.93) - Microscopic/macroscopic/ 0.85 (0.30-2.42) 1.38 (0.46-4.15) - 1.40 (0.44-4.43) - residual, NOS 0.85 (0.30-2.42) 1.38 (0.46-4.15) - 1.40 (0.44-4.43) - Vor evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - Vor evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - Vor evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - Vor evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - Vor evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) - 0.86 (0.31-2.38) - Vor evaluable/missing 0.35 (1.30-7.18) 1.51 (0.45-5.07) - - - 1.3 0.51 (2.30-13.67) 1.51 (0.45-5.07) - - - - 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) - - N stage 0. | No primary surgery | Ref | Ref | Ref | Ref | Ref |
| Not residuality 0.81 (0.20 + 0.02) 1.81 (0.45 + 0.05) 0.00 (0.17 + 0.5) Microscopic/macroscopic/ 0.85 (0.30 - 2.42) 1.38 (0.46 - 4.15) - 1.40 (0.44 - 4.43) - residual, NOS 0.66 (0.15 - 0.86) 0.80 (0.30 - 2.17) - 0.86 (0.31 - 2.38) - T stage (collapsed) 0/CIS Ref. Ref. Ref. Ref. 1 3.05 (1.30 - 7.18) 1.53 (0.54 - 4.34) - 2 3.87 (1.55 - 9.71) 1.89 (0.57 - 6.29) - 3 5.61 (2.30 - 13.67) 1.51 (0.45 - 5.07) - 4 9.17 (2.37 - 35.53) 6.44 (1.46 - 28.38) 6.28 (1.83 - 21.55) Missing 4.48 (1.67 - 12.04) 1.33 (0.31 - 5.66) - N stage 0 Ref. Ref. 0.71 (0.16 - 3.16) - 1 1.07 (0.26 - 4.45) 0.71 (0.16 - 3.16) - 2.93 (1.36 - 6.23) 2.90 (1.48 - 5.68) 3 9.93 (4.36 - 22.64) 12.6 (4.01 - 39.56) 13.17 (5.28 - 32.89) - Missing 1.68 (0.83 - 3.36) 1.68 (0.45 - 6.23) - - M stage 0 Ref. | No residual tumor | 0 49 (0 29-0 82) | 1.01 (0.55-1.88) | _ | 0.98 (0.49-1.93) | _ |
| Introducipation Introducipation Introducipation Introducipation residual, NOS Not evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) – 0.86 (0.31-2.38) – T stage (collapsed) 0/CIS Ref. Ref. Ref. Ref. 1.53 (0.54-4.34) – 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) – – – – 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) – – – – 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) – – – – 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) – – – N stage 0 Ref. Ref. Ref. – – – 0 Ref. 1.07 (0.26-4.45) 0.71 (0.16-3.16) – <t< td=""><td>Microscopic/macroscopic/</td><td>0.85 (0.30-2.42)</td><td>1 38 (0 46-4 15)</td><td>_</td><td>1 40 (0 44-4 43)</td><td>_</td></t<> | Microscopic/macroscopic/ | 0.85 (0.30-2.42) | 1 38 (0 46-4 15) | _ | 1 40 (0 44-4 43) | _ |
| Not evaluable/missing 0.36 (0.15-0.86) 0.80 (0.30-2.17) – 0.86 (0.31-2.38) – T stage (collapsed) 0/CIS Ref. Ref. Ref. Ref. Ref. 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) – 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) – 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) – 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) – N stage 0 Ref. Ref. Ref. 0 Ref. 1.07 (0.26-4.45) 0.71 (0.16-3.16) – 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) – Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) – Missing 1.08 (0.61-1.91) 0.77 (0.10-6.13) – 0 Ref. Ref. Ref. Ref. 1 1.575 (3.65-67.96 | residual NOS | 0.00 (0.00 2.12) | | | | |
| T stage (collapsed) Ref. Ref. Ref. Ref. Ref. 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) - 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) - 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) - 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) - N stage 0 Ref. Ref. Ref. 0 Ref. 0.71 (0.16-3.16) - 1 1.07 (0.26-4.45) 0.71 (0.16-3.16) - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. 0 Ref. Ref. Ref. - 1 15.75 (3.65-67.96) 0.77 (0.10-6.13) - 0 Ref. Ref. Ref. - 1 10.8 (0.61-1.91) 0.97 | Not evaluable/missing | 0 36 (0 15-0 86) | 0 80 (0 30-2 17) | _ | 0.86 (0.31-2.38) | _ |
| Normalization Ref. Ref. Ref. Ref. Ref. Ref. 1 3.05 (1.30-7.18) 1.53 (0.54-4.34) - 2 3.87 (1.55-9.71) 1.89 (0.57-6.29) - 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) - 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) - N stage 0 Ref. Ref. Ref. 0 Ref. Nef. - - 1 1.07 (0.26-4.45) 0.71 (0.16-3.16) - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) - - M stage 0 Ref. Ref. Ref. 0 Ref. Ref. Ref. - 1 15.75 (3.65-67.96) 0.77 (0.10-6.13) - 1 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | T stage (collapsed) | | | | 0.00 (0.01 2.00) | |
| I $3.05 (1.30-7.18)$ $1.53 (0.54-4.34)$ $-$ 2 $3.87 (1.55-9.71)$ $1.89 (0.57-6.29)$ $-$ 3 $5.61 (2.30-13.67)$ $1.51 (0.45-5.07)$ $-$ 4 $9.17 (2.37-35.53)$ $6.44 (1.46-28.38) 6.28 (1.83-21.55)$ Missing $4.48 (1.67-12.04)$ $1.33 (0.31-5.66)$ $-$ N stage 0 Ref.Ref.Ref.0Ref.Ref.Ref. $-$ 2 $4.09 (2.11-7.91)$ $2.93 (1.36-6.33) 2.90 (1.48-5.68)$ 3 $9.93 (4.36-22.64)$ $12.6 (4.01-39.56) 13.17 (5.28-32.89)$ Missing $1.68 (0.83-3.36)$ $1.68 (0.45-6.23)$ $-$ M stage 0 Ref.Ref.Ref.1 $15.75 (3.65-67.96)$ $0.77 (0.10-6.13)$ $-$ Missing $1.08 (0.61-1.91)$ $0.97 (0.23-4.14)$ $-$ | 0/CIS | Ref. | | | Ref. | Ref. |
| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | I | 3.05 (1.30-7.18) | | | 1.53 (0.54-4.34) | _ |
| 3 5.61 (2.30-13.67) 1.51 (0.45-5.07) - 4 9.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55) Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) - N stage 0 Ref. Ref. Ref. 1 1.07 (0.26-4.45) 0.71 (0.16-3.16) - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) - - M stage 0 Ref. Ref. Ref. 0 Ref. Ref. Ref. - M stage 0 1.68 (0.83-3.36) - - M stage 0 Ref. Ref. Ref. 1 15.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | 2 | 3.87 (1.55-9.71) | | | 1.89 (0.57-6.29) | _ |
| 49.17 (2.37-35.53) 6.44 (1.46-28.38) 6.28 (1.83-21.55)Missing4.48 (1.67-12.04)1.33 (0.31-5.66) $-$ N stage 0 Ref.Ref.Ref.0Ref.1.07 (0.26-4.45)0.71 (0.16-3.16) $-$ 24.09 (2.11-7.91)2.93 (1.36-6.33)2.90 (1.48-5.68)39.93 (4.36-22.64)12.6 (4.01-39.56)13.17 (5.28-32.89)Missing1.68 (0.83-3.36)1.68 (0.45-6.23) $-$ M stage 0 Ref.Ref.Ref.115.75 (3.65-67.96)0.77 (0.10-6.13) $-$ Missing1.08 (0.61-1.91)0.97 (0.23-4.14) $-$ | 3 | 5.61 (2.30-13.67) | | | 1.51 (0.45-5.07) | _ |
| Missing 4.48 (1.67-12.04) 1.33 (0.31-5.66) - N stage 0 Ref. Ref. Ref. Ref. 0 Ref. 0.71 (0.16-3.16) - - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. 0 Ref. Ref. Ref. - M stage 0 1.68 (0.61-1.91) 0.97 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | 4 | 9.17 (2.37-35.53) | | | 6.44 (1.46-28.38) | 6.28 (1.83-21.55) |
| N stage Ref. Ref. Ref. - 0 Ref. 0.71 (0.16-3.16) - 1 1.07 (0.26-4.45) 0.71 (0.16-3.16) - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. 0 Ref. Ref. 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | Missing | 4.48 (1.67-12.04) | | | 1.33 (0.31-5.66) | _ |
| No. Ref. Ref. Ref. Ref. Ref. Ref. I | N stage | | | | | |
| I 1.07 (0.26-4.45) 0.71 (0.16-3.16) - 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) I2.6 (4.01-39.56) I3.17 (5.28-32.89) Missing 1.68 (0.83-3.36) I.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. I I5.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | 0 | Ref. | | | Ref. | Ref. |
| 2 4.09 (2.11-7.91) 2.93 (1.36-6.33) 2.90 (1.48-5.68) 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. 1 15.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | I | 1.07 (0.26-4.45) | | | 0.71 (0.16-3.16) | _ |
| 3 9.93 (4.36-22.64) 12.6 (4.01-39.56) 13.17 (5.28-32.89) Missing 1.68 (0.83-3.36) 1.68 (0.45-6.23) - M stage 0 Ref. Ref. Ref. 1 15.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | 2 | 4.09 (2.11-7.91) | | | 2.93 (1.36-6.33) | 2.90 (1.48-5.68) |
| Missing I.68 (0.83-3.36) I.68 (0.45-6.23) - M stage 0 Ref. Ref. 0 Ref. 0.77 (0.10-6.13) - Missing I.08 (0.61-1.91) 0.97 (0.23-4.14) - | 3 | 9.93 (4.36-22.64) | | | 12.6 (4.01-39.56) | 13.17 (5.28-32.89) |
| M stage Ref. | Missing | 1.68 (0.83-3.36) | | | 1.68 (0.45-6.23) | _ |
| O Ref. Ref. Ref. I 15.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing I.08 (0.61-1.91) 0.97 (0.23-4.14) - | M stage | | | | · · · / | |
| I I5.75 (3.65-67.96) 0.77 (0.10-6.13) - Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) - | 0 | Ref. | | | Ref. | Ref. |
| Missing 1.08 (0.61-1.91) 0.97 (0.23-4.14) – | I | 15.75 (3.65-67.96) | | | 0.77 (0.10-6.13) | - |
| | Missing | 1.08 (0.61-1.91) | | | 0.97 (0.23-4.14) | - |

Abbreviations: CI, confidence interval; HR, hazard ratio; Ref., reference.

^aBold font indicates a statistically significant HR.

^bFor comparison to the full and reduced model, these are the significant HRs from the univariable analyses from Table 2.

^cNon-Hispanic includes one unknown.

Table 4. Hazard Ratios for the Risk Groups Identified by CART Analysis.^a

| Risk group | Covariates | 5-year survival rate (%) | Univariable, HR (95% CI) | Multivariable, ^b HR (95% CI) |
|---------------|---|-----------------------------|-----------------------------|--|
| Model | 1 | | | |
| Ι | TNM stage 0-II + Penis glans/NOS + Macroscopic/microscopic/residual tumor, NOS | 81.2 | Ref. | Ref. |
| 2 | TNM stage 0-II + Prepuce/penis corpus/overlapping lesion + No residual tumor, NOS | 58.7 | 2.57 (1.19-5.54) | 1.73 (0.76-3.98) |
| 3 | TNM stage 0-II + Prepuce/penis corpus/overlapping lesion + married | 62.5 | 2.19 (0.90-5.31) | 2.19 (0.89-5.36) |
| 4 | TNM stage 0-II + Prepuce/penis corpus/overlapping lesion + Single/ widowed/divorced | 18.2 | 6.51 (2.86-14.81) | 5.29 (2.17-12.90) |
| 5 | TNM stages III/IV + Prepuce/penis corpus/overlapping lesion + No residual tumor, NOS | 46.5 | 4.52 (2.20-9.31) | 3.35 (1.53-7.35) |
| 6 | TNM stages III/IV + Prepuce/penis corpus/overlapping lesion + Macroscopic/microscopic/residual tumor, NOS | 22.2 | 10.93 (5.11-23.36) | 8.59 (3.71-19.92) |
| Model | 2 | | | |
| Ι | N stage 0-1 + Moderately/well-differentiated/undetermined + Penis glans/penis, NOS | 79.1 | Ref. | Ref. |
| 2 | N stage 0-1 + Moderately/well-differentiated/undetermined + Prepuce/ penis corpus/overlapping lesion + Married | 67.7 | 1.49 (0.56-3.97) | 1.63 (0.61-4.37) |
| 3 | N stage 0-1 + Moderately/well-differentiated/undetermined + Prepuce/ penis corpus/overlapping lesion + Single/widowed/divorced | 34.3 | 4.39 (1.76-10.94) | 3.61 (1.38-9.47) |
| 4 | N stage 0-1 $+$ Poorly and undifferentiated | 45.0 | 3.79 (1.72-8.35) | 3.61 (1.10-11.84) |
| 5 | N stage missing | 54.2 | 2.46 (1.15-5.26) | 2.46 (1.14-5.32) |
| 6 | N stage 2-3 | - | 7.82 (4.11-14.85) | 6.45 (3.16-13.15) |

Abbreviations: Cl, confidence interval; CART, classification and regression tree; HR, hazard ratio; NOS, not otherwise specified; Ref., reference.

^aBold font indicates a statistically significant HR. ^bAdjusted for alcohol use, ethnicity, and tumor differentiation.

TNM stage 0-II and prepuce/penis corpus/overlapping lesions, single/widowed/divorced men had worse survival (5-year survival = 18.2%) versus married men (5-year survival = 62.5%). Moreover, late-stage men with prepuce/penis corpus/overlapping lesion and no residual tumor (5-year survival = 46.5%) and macroscopic/microscopic/residual tumor, NOS (5-year survival = 22.2%) had better survival than single/widowed/ divorced men with early stage and prepuce/penis corpus/overlapping lesions. In addition to clinicopathologic covariates, these data may suggest that social support network has a beneficial impact on outcomes of patients diagnosed with early stage disease.

From the full model which contained all significant univariable covariates (Table 3), we utilized feature reduction analyses which yielded 2 reduced models and 2 CART models. These 4 models yielded 7 covariates: tumor site, staging variables (TNM stage, N stage, and T stage), tumor differentiation, marital status, alcohol use, surgical margins, and tumor size. Staging variables and tumor site were found in all 4 models which are well-established cancer prognostic factors. Consistent with previous reports,^{4-8,16} our analysis found that diagnosis of advanced disease stage significantly associated with decreased patient survival for penile cancer. In support of this, a report by Pandey et al indicated positive inguinal and pelvic nodal metastasis as predictors for shorter survival.⁶ These results are in concordance with our finding that men with TNM stage III/IV have worse outcomes compared to those with TNM stage 0/CIS/I/II. Our analyses also demonstrated that men with tumors located at the penis glans had a reduced risk of mortality compared to those with prepuce/penis corpus/overlapping lesions. About 70% of penile SCCs originate in the mucosal epithelium of the glans¹⁷ and typically tumors at the glans are diagnosed at earlier stages, while corpus is diagnosed at later stages.¹⁸ Men diagnosed with poorly differentiated tumors are associated with having advanced stage and nodal metastasis. There are several studies that have found that those with poorly differentiated tumors are at increased risk of poor prognosis.¹⁹ Tumor size, which is correlated with tumor stage, is also associated with survival outcomes. A prior study found that 5-year survival rate was 56% to 78% for men with tumor thickness of <5 mm and decreases to 48% to 64% for tumors that are >5mm.²⁰ We also found that alcohol use was also significantly associated with survival. Alcohol consumption is a potent modulator of immune function,²¹ associated with an increased risk of penile cancer,²² and classified as a "Group 1 Carcinogen" by the International Agency for Research of Cancer.²³ It is believed that alcohol consumption may impair host defense against viral infections.²⁴ An analysis of alcohol consumption in men revealed that alcohol consumption is associated with an elevated risk for human papillomavirus (HPV) infection.²⁴ Considering that HPV is detected in 48% of penile cancers,²⁵ we can speculate that HPV infections may have played a biological role in the increased risk of mortality observed in our cohort of patients with penile cancer who were alcohol users.

Our novel decision tree approach (i.e., CART analysis) demonstrated that a subset of unmarried early-stage patients



(group 5 and group 6), stage 0 to II was segmented by tumor site forming terminal node for penis glans/NOS and was further segmented by macroscopic/microscopic/residual tumor, NOS (group 1) and no residual tumor (group 2), and prepuce/corpus/overlapping lesion as married (group 3), or single/divorced/widowed (group 4). B, N stage segmented to form terminal nodes for N stage 2 to 3 (group 5 and group 6), and N stage 0 to 1 which further segmented on tumor differentiation, resulting in terminal node for poorly and undifferentiated tumors (group 4). Tumor differentiation further segmented on tumor site with a terminal node for penis glans/NOS (group 1) and marital status segmented to 2 terminal nodes for married (group 2) and single/widowed/divorced (group 3). C, Overall differences in survival between CART groups for model 1 were significant (log rank P < .001). D, Overall differences in survival Figure 1. A, Associations between tumor stage and overall survival were assessed by CART analysis. For model 1, while stage II/IV + surgical margins generated 2 terminal nodes between CART groups for model 2 were significant (log rank P < .001). CART indicates classification and regression tree; NOS, not otherwise specified. 0.722) and CART model 2 (ROC = 0.709) to be more accurate predictors of prognosis than TNM stage (ROC = 0.643). CART indicates classification and regression tree; ROC, receiver operating characteristics.

Table 5. Harrell's C-index and ROC.

| Characteristics | C-index | ROC |
|--|---------|-------|
| TNM stage only Model I (TNM stage included) | 0.674 | 0.643 |
| Full multivariable model | 0.741 | _ |
| Reduced model | 0.713 | _ |
| CART | 0.716 | 0.722 |
| Model 2 (TNM stage included) | | |
| Full multivariable model | 0.756 | _ |
| Reduced model | 0.744 | - |
| CART | 0.707 | 0.709 |

Abbreviations: CART, classification and regression tree; ROC, receiver operating characteristics.

had worse outcomes than late-stage patients (Figure 1C-D). Due to the potentially mutilating surgery that patients may face, patients with penile cancer may experience an increase in psychological distress that requires psychosocial care.²⁶ The emotional stress they experience can result from a multitude of factors, including the diagnosis itself, treatment, and fear of recurrence.²⁶ A study about marital status among men with penile cancer demonstrated marital status to be an independent prognostic factor for early stage but not for late-stage disease.²⁷ Additionally, other studies have also revealed significant associations between marital status and penile cancer. A population-based registry study by Torbrand et al²⁸ showed that married men with penile cancer had lower risk of latestage tumors compared to divorced and never married men. However, the authors did not find an association between marital status and cause-specific mortality. Other studies have revealed an increased risk of mortality for nonmarried men with localized advanced penile cancer²⁹ and an increased risk of invasive penile cancer for nonmarried men.^{29,30} Similar to the results found in our analyses, Rippentrop et al showed that

married men with localized or regional penile carcinoma had significantly longer survival time compared to never married patients. However, among patients with distant disease, there were no significant differences by marital status.³¹ Our study provides further evidence of the importance of marital status on outcomes in patients with penile cancer.

We acknowledge that there are limitations with these analyses. The higher number of non-Hispanic whites from a single comprehensive cancer center may limit the generalizability of our findings. The incidence of penile carcinoma is higher among Hispanics and African American men compared to non-Hispanic whites³² and despite having a small cohort of Hispanics within our study, we observed Hispanics to have an increased risk of dying. The status of other prognostic factors such as HPV type and molecular markers (p16 expression) in our study cohort is not known. As basaloid carcinomas are more often diagnosed over other subtypes among those infected with HPV, a future stratified analysis by HPV status may be possible with a larger cohort of patients with penile cancer. An investigation of whether patients diagnosed with early-stage penile cancer are likely to depend more on psychosocial support and how it effects outcomes is important to understanding the influences of these potential risk factors on prognosis.33,34

In summary, this study identified novel prognostic models for penile cancer that were largely concordant in terms of the covariates that were identified based on the 2 analytical approaches (backward elimination and decision tree analysis). Importantly, our decision tree analyses showed that men diagnosed with early-stage penile cancer and that were not currently married had poorer survival compared to men that were currently married. The coexistence of poor clinicopathologic variables in combination with lack of social support may negatively impact prognosis. It can be concluded from this study that the assessment of clinicopathologic risk factors, including socioeconomic factors, may help to identify those at high risk and that would benefit from psychosocial support.

Authors' Note

The protocol for this study (IRB#Pro00014743) was approved by Advarra (CR# 00150095), and waiver of patient consent was obtained. M.E.R. and H.B. contributed equally.

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Declaration of Conflicting Interests

The author(s) declared the following potential conflicts of interest with respect to the research, authorship, and/or publication of this article: Dr Phillipe E. Spiess serves as Vice-chair of the NCCN Bladder and Penile Cancer Panel.

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Figure 2. Comparison of ROC curves show CART model I (ROC =

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