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Postoperative prostate-specific antigen nadir improves accuracy for predicting biochemical recurrence after radical prostatectomy: Results from the Shared Equal Access Regional Cancer Hospital (SEARCH) and Duke Prostate Center databases

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Abstract

Objectives—We previously showed that prostate-specific antigen (PSA) nadir after radical prostatectomy (RP) significantly predicts biochemical recurrence (BCR). Herein, we sought to explore the effect of including PSA nadir into commonly used models on their accuracy to predict BCR after RP.

Methods—This was a retrospective analysis of 943 and 1792 subjects from the Shared Equal Access Regional Cancer Hospital (SEARCH) and Duke Prostate Cancer (DPC) databases, respectively. The discrimination accuracy for BCR of seven previously published models was assessed using concordance index and compared with and without adding PSA nadir level in SEARCH. Using data from SEARCH, we developed a new nomogram incorporating PSA nadir to other known predictors (preoperative PSA, pathological Gleason score, PSA nadir level, surgical findings, prostate weight, body mass index and race) of BCR and externally validated it in the DPC.

Results—In SEARCH, the mean concordance index across all seven nomograms was 0.687. After the inclusion of PSA nadir, the concordance index increased by nearly 7% (mean = 0.753). The concordance index of the new nomogram in SEARCH was 0.779 (bias-corrected = 0.767), which was 5% better than the next best model. In DPC, the new nomogram yielded a concordance index of 0.778.

Conclusion—The addition of postoperative PSA nadir to commonly used nomograms increased their accuracies by nearly 7%. Based upon this, we developed and externally validated a new nomogram, which was well calibrated and highly accurate, and is a potentially valuable tool for patients and physicians to predict BCR after RP.

Keywords

disease-free survival; nomograms; prostate cancer; prostate-specific antigen; prostatectomy; validation studies

Introduction

Despite contemporary advances in prostate cancer, such as widespread disease screening, better surgical candidate selection and improved operative techniques, 15–40% of patients treated with radical prostatectomy (RP) will experience biochemical recurrence (BCR).¹ After recurrence, these patients are at a higher risk for disease progression and death.² Several studies have evaluated the use of early secondary treatments after surgery to decrease BCR and overall mortality. Unfortunately, not all men necessarily benefit from such therapies.^{3–6} Some patients may never develop prostate-specific antigen (PSA) elevation and many others may have an indolent recurrence not requiring treatment. It is critical to properly select patients for adjuvant therapies in order to minimize side-effects and maximize their benefits.⁷ Currently there are several predictive models to risk stratify patients after surgery.⁸ The large majority of these models include preoperative and perioperative variables (e.g. PSA before surgery, pathological Gleason score and tumor stage). However, given adjuvant therapy is usually started a few months after RP and by that time, postoperative PSA levels have reached nadir, it seems reasonable to include postoperative PSA in the decision of whether or not to start secondary treatments. Previously, we showed postoperative PSA nadir was highly correlated with risk of BCR⁹ and overall mortality.¹⁰ Therefore, we sought to explore the effect of including PSA nadir into commonly used models and nomograms on their accuracy to predict BCR after RP among subjects of the Shared Equal Access Regional Cancer Hospital (SEARCH) database. Additionally, we developed a new nomogram incorporating nadir and other well-established predictors of BCR and externally validated it in the Duke Prostate Cancer (DPC) database.

Methods

Study population

After obtaining institutional review board approval from each institution, data from patients undergoing RP between 1998 and 2008 at four Veteran Affairs Medical Centers (West Los Angeles and Palo Alto, CA; Augusta, GA; and Durham, NC) were combined into the SEARCH database (development cohort).¹⁰ The database includes information on patient age at surgery, race, height, weight, clinical stage, cancer grade on diagnostic biopsies, preoperative PSA, surgical specimen pathology (specimen weight, tumor grade, stage and surgical margin status) and follow-up PSA. We restricted our analyses to patients treated between 1998 and 2008 because prior to this time patients were not followed with ultrasensitive PSA assays (sensitivity of 0.01 ng/mL). Data from patients treated at Duke University Medical Center between 1998 and 2007 were abstracted into the DPC Database

(validation cohort).¹¹ The database includes similar information to SEARCH; however, patients were followed with non-ultrasensitive PSA (sensitivity of 0.1 ng/mL). External validation on a non-ultrasensitive PSA cohort was chosen given it allows an assessment of the generalizability of our results even when ultrasensitive PSA is unavailable. Patients treated with preoperative hormonal therapy or radiotherapy were excluded from both cohorts. Of 1300 patients in SEARCH, we excluded eight (<1%) due to start of adjuvant therapy before PSA nadir, 130 (10%) due to missing nadir within 1–6 months postoperatively, 114 (9%) whose nadir was ≥ 0.2 ng/mL (i.e. they recurred based upon nadir PSA value and thus are not at risk for future recurrence), 101 (8%) due to missing follow-up data after nadir and four (<1%) due to positive lymph nodes. This resulted in a study population of 943 subjects in SEARCH. Of 3130 patients in DPC, we excluded 856 (27%) due to missing nadir within 1–6 months postoperatively, 142 (4%) whose nadir was ≥ 0.2 ng/mL, 343 (11%) due to missing follow up after nadir and 17 (<1%) due to positive lymph nodes. This resulted in a population of 1792 subjects in DPC.

All patients were followed with serial PSA and clinical visits at intervals according to attending physician discretion. PSA nadir was defined as the lowest PSA level within 1–6 months postoperatively and before BCR.⁹ Additional postoperative treatments were at the judgment of the patient and treating physician. In both SEARCH and DPC, BCR was defined as a postoperative PSA level ≥ 0.2 ng/mL or secondary treatment for an elevated PSA.¹² For our study purposes, the definition adopted, which slightly differs from the American Urologic Association (AUA) definition,¹³ provides a better classification for patients with rising PSA who received early secondary treatment. The AUA definition, which has PSA-only criteria, would categorize these patients as not having developed biochemical recurrence. We believe this creates a bias in that men at very high risk of recurrence (i.e. PSA of 0.19 ng/mL who received salvage radiation) are censored. This could artificially make the recurrence rates look better than in actuality. Thus, we believe our composite definition based upon PSA and clinical practice (i.e. treatment for an elevated PSA) is better suited to reduce this potential bias.

Statistical analysis

The predictive discrimination of the following seven models was assessed in SEARCH: (i) the risk classification published by D'Amico *et al.*;¹⁴ (ii) the pre-¹⁵ and (iii) postoperative¹⁶ nomograms of Kattan *et al.*; (iv) the DPC nomogram;¹² (v) the Cancer of the Prostate Risk Assessment (CAPRA) model;¹⁷ (vi) the Center for Prostate Disease Research (CPDR/CaPSURE) model;¹⁸ and (vii) the Johns Hopkins Hospital model.¹⁹ The overall discrimination for BCR of the various models was determined using Harrell's concordance index (c-index).²⁰ In exploratory analysis, we calculated the c-index for each model with and without PSA nadir level to determine the changes in concordance by adding nadir. To obtain the accuracy of the models with nadir, we calculated the linear predictor for all subjects based on each nomogram and then used this value along with PSA nadir in a proportional hazard model.

For nomogram development, we considered the following variables for the model: age at surgery (in years, continuous), race (white or non-white), body mass index (BMI in kg/m², continuous and log-transformed), year of surgery (in years, continuous), preoperative PSA (in ng/mL, continuous and log-transformed), PSA nadir level (in ng/mL, continuous and log-transformed), clinical T stage (T₁, T₂₋₃), pathological Gleason score (2–6, 3+4, $\geq 4+3$), surgical findings (organ-confined and margin-negative [OC and M–], positive surgical margins [PSM] and/or extracapsular extension [ECE], seminal vesicle invasion [SVI]) and prostate weight (in grams, continuous and log-transformed). Stepwise variable selection was employed. We used the Cox proportional hazards to analyze the association between the above covariates and time to BCR. The nomogram was created based on the Cox regression

model and designed to predict the BCR-free probability at 1, 3 and 5 years postoperatively. We internally validated the nomogram in SEARCH by determining the bias-corrected c-index using bootstrapping. To examine the effect of PSA nadir in the accuracy of the new model, we internally validated it separately in men with and without detectable nadir values and then compared it to the performance of the other nomograms. External validation of the new nomogram was done in DPC using calibration plots and c-index. All statistical analyses were performed using Stata 10.1 (StataCorp, College Station, TX, USA) and R 2.9.0 (R Foundation for Statistical Computing, Vienna, Austria). $P < 0.05$ was considered statistically significant.

Results

In DPC, 95% had an undetectable PSA nadir vs only 68% in SEARCH (Table 1). In both datasets, there was a similar percentage (3–5%) with nadir values ranging from 0.1 to 0.19 ng/mL. Overall, patients in SEARCH had a higher risk of BCR compared to those from DPC (Fig. 1).

In SEARCH, the mean c-index for BCR across all seven nomograms was 0.687, ranging from 0.650 to 0.716 (Table 2). After including PSA nadir level to all seven models, the mean c-index increased by nearly 7% (mean = 0.753; range 0.737–0.772). Nadir level alone as a predictor of BCR yielded a c-index of 0.661, which was higher than any other individual variable alone (data not shown). The independent predictors of BCR in SEARCH were higher PSA nadir level, higher BMI, non-white race, higher pathological Gleason score, and adverse pathological findings (Table 3). The following variables were included in the nomogram: preoperative PSA, pathological Gleason score, PSA nadir level, surgical findings, prostate weight, BMI and race (Fig. 2). The c-index of the new nomogram in SEARCH was 0.779 with a bias-corrected c-index of 0.767. Of note, this compares to a bias-corrected c-index of 0.718 for the same multivariate model without nadir. Thus, nadir increased the accuracy of our ability to risk stratify men for BCR by 5%. In patients with an undetectable nadir, the new nomogram had a c-index similar to other nomograms (concordance index [CI] = 0.720, Table 4). However, among patients with non-zero nadir, the new nomogram had considerably better performance (CI = 0.791).

To externally validate our results among men followed with non-ultrasensitive PSA, we determined the c-index and calibration in DPC. Compared to SEARCH, patients treated in DPC had similar age, BMI, preoperative PSA and prostate weights. However, subjects in DPC were more likely to be white, had higher ECE and SVI and yet had lower postoperative Gleason scores and lower PSM (Table 1). The predicted risk of BCR within 5 years postoperatively in DPC based on the SEARCH nomogram was calculated across quartiles of risk and compared with the actuarial risk (Fig. 3). Overall, while the nomogram was relatively accurate, it slightly overpredicted disease recurrence in DPC (approximately 5% above the actuarial risk). In the lower risk groups the overprediction was about 3% while in the highest risk groups the predicted risk was 7% higher than the actuarial risk. Nevertheless, except for the highest risk quartile, the ideal prediction line fell within the predicted 95% confidence intervals and the overall shape of the calibration curve was linear, indicating that higher predicted risk did entail a higher actual risk. Finally, even in a cohort of men followed with non-ultrasensitive PSA and when very few men had detectable PSA (i.e. only 5%), the new nomogram had a comparable performance with other well-established nomograms (Table 5).

Discussion

Several studies have examined the predictive accuracy of new markers (i.e. cytokines, genes and pathological parameters) with mixed results.^{21–26} Most of these new markers are associated with additional costs and labors. Consequently, there is no single marker or set of biomarkers regularly employed in prostate cancer prognostication today, except for PSA. PSA is inexpensive and readily available and is thus used routinely in several prostate cancer settings, such as screening, risk stratification and follow up. We have previously shown that postoperative PSA nadir is highly associated with disease recurrence⁹ and overall survival,¹⁰ being potentially useful in risk assessment. However, to date, only a few limited studies evaluated the predictive properties of postoperative PSA nadir.^{27,28} Therefore, we sought to explore the predictive characteristics of PSA nadir and the effect of including nadir into commonly used models and nomograms on their accuracy to predict BCR after RP.

In SEARCH, the currently available models to predict BCR after RP yielded a mean c-index of 0.687. After including postoperative PSA nadir, the c-index increased by nearly 7%. Although the clinical meaning of this improvement could not be determined in the present study, this improvement was equivalent to improvements seen by the use of the best new biomarkers currently under investigation.²³ However, in contrast to new markers, no extra cost or labor is involved with PSA nadir. Additionally, nadir alone yielded a c-index of 0.661, which is comparable to the accuracy of preoperative models and higher than any other single variable alone. In patients with a detectable nadir, the c-index of nadir alone was 0.745, which was higher than the accuracy of currently available nomograms in our population. Therefore, these results reinforce our previous findings that PSA nadir is useful in risk stratification.^{9,10}

We then developed a new nomogram including traditional predictors of BCR (i.e. surgical findings, grade, and pre-operative PSA) along with PSA nadir. We also included some non-traditional variables (i.e. race, BMI and prostate weight) given previous studies showed they can improve nomogram accuracy.¹² In SEARCH, the new nomogram yielded a c-index of 0.770, which was nearly 8% higher than the average of the other nomograms and over 5% higher than the next best model. We then sought to validate it using an external dataset, the DPC cohort. Therein, we found the nomogram performed equally well with a c-index of 0.778. Thus, even in a cohort where ultrasensitive PSA is unavailable, our model appears to be a highly reliable, validated nomogram to predict disease recurrence after RP.

Among men with an undetectable nadir, the new nomogram had a comparable performance to other nomograms (CI = 0.720). We also found similar results in DPC, wherein 95% of patients had undetectable PSA nadirs; our nomogram had a similar performance to other well-established nomograms. However, within SEARCH, the c-index was considerably higher when the nomogram was applied to men with a detectable PSA nadir (CI = 0.790). Unfortunately, such an analysis was not possible in DPC given patients were followed with non-ultrasensitive PSA (with only one decimal digit) and there were only two possible nadir values: undetectable and 0.1 ng/mL. This lack of data granularity decreases the model's capability of stratifying patients given men with considerably different prognoses are grouped into the same category. For example, in DPC, men whose actual nadir value varied from 0 to 0.09 ng/mL were determined to have an “undetectable” nadir as their PSA values were below the limits of detection of the assay. However, we previously showed that patients with a nadir of 0.05–0.09 ng/mL are 12 times more likely to recur than those with an undetectable nadir using an ultrasensitive assay.⁹ As such, this absence of granular data on the exact nadir value likely reduced the performance of our nomogram in DPC. Nevertheless, we found that even when used with non-ultrasensitive PSA data, our

nomogram had a performance that was equivalent to other well-established nomograms. However, when used with ultrasensitive PSA data (i.e. SEARCH), our model clearly had a superior performance. Thus, the strengths of our nomogram include the use of readily available variables without adding further costs or labor, it applies to both ultra-sensitive and non-ultrasensitive PSA populations, it is highly-accurate, validated and easy-to-use. However, it is necessary to further validate our nomogram in datasets with ultrasensitive PSA nadir data in order to confirm the dramatic improvements in accuracy seen within SEARCH.

The present study is limited by the retrospective nature of our cohort. Additionally, patients at very high risk (i.e. those whose nadir is ≥ 0.2 ng/mL) were not evaluated in the study because those patients were not at risk for BCR given that they already met the criteria for recurrence based upon their nadir PSA value. Moreover, the differences in demographics, pathological features and prognoses between SEARCH and DPC may have influenced the accuracy of our nomogram in external validation. However, these significant variations are in fact the strengths of our study because they allow for the assessment of generalizability of our model and transportability of our results among considerably dissimilar populations. Although recurrence has been shown to correlate with progression to metastatic disease²⁹ and higher prostate-cancer specific mortality,³⁰ models to predict long-term outcomes are still needed to further improve patient selection for adjuvant treatments. Finally, the use of the nomogram in medical practice is necessary to determine its clinical relevance.

In conclusion, the addition of postoperative PSA nadir to commonly used nomograms to predict BCR after RP considerably increased their predictive accuracies by nearly 7%. Based upon this, we developed and externally validated a new nomogram, which was well calibrated and highly accurate and, in our opinion, is a valuable tool for patients and physicians to predict BCR after RP.

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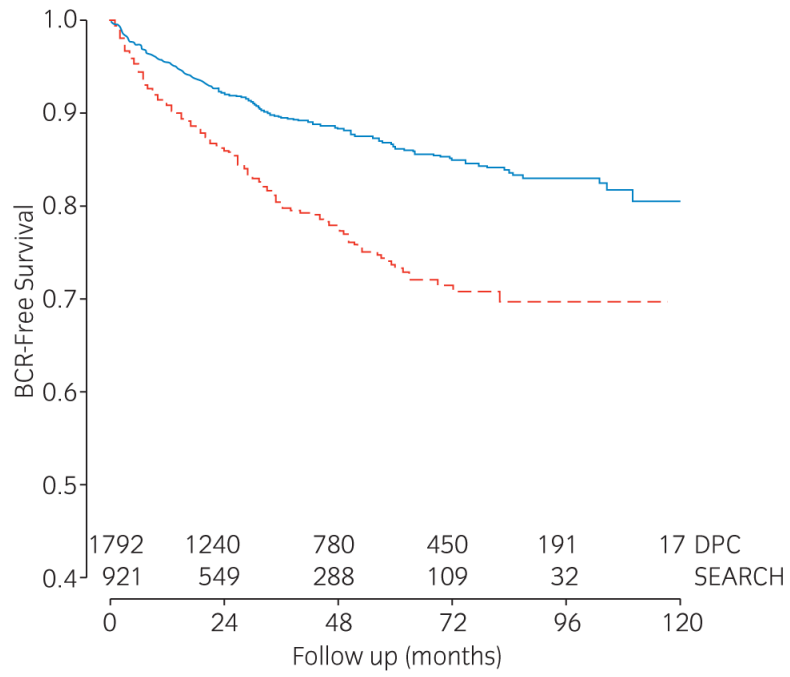


Fig. 1. Kaplan-Meier plot depicting biochemical recurrence (BCR)-free survival after radical prostatectomy in the (---) Shared Equal Access Regional Cancer Hospital (SEARCH) cohort ($n = 921$) and the (—) Duke Prostate Cancer (DPC) ($n = 1792$) cohort.

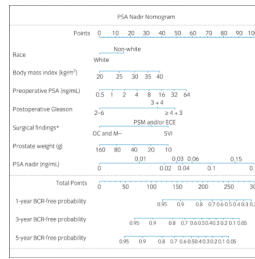


Fig. 2. Prostate-specific antigen (PSA) nadir nomogram. BCR, biochemical recurrence; ECE, extracapsular extension; *OC and M-, organ-confined and margin-negative; PSM, positive surgical margins; SVI, seminal vesicle invasion.

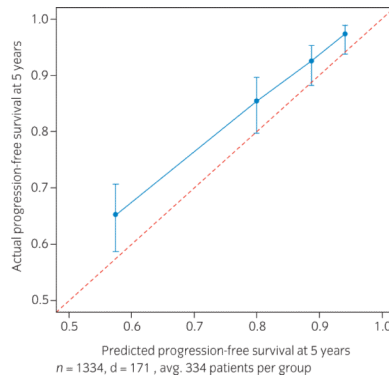


Fig. 3. Calibration plot depicting predicted biochemical recurrence (BCR) compared to actuarial BCR at 5 years after radical prostatectomy in the Duke Prostate Cancer cohort.

Table 1

Baseline characteristics

Variable	SEARCH (Development) No. (%)	DPC (Validation) No. (%)
Total number of patients	943	1792
PSA nadir (ng/mL)		
Undetectable	642 (68)	1709 (95)
0.01	87 (9)	–
0.02	67 (7)	–
0.03	43 (4)	–
0.04	25 (3)	–
0.05–0.09	49 (5)	–
0.10–0.19	31 (3)	83 (5)
Age at surgery (years)		
Mean (SD)	60 (6.4)	61 (7.3)
Median (IQR)	60 (56–65)	61 (56–66)
Veteran Affairs Medical Center		
1	320 (34)	–
2	72 (8)	–
3	237 (25)	–
4	314 (33)	–
Race		
White	444 (48)	1451 (81)
Non-white	475 (52)	337 (19)
Body mass index (kg/m ²)		
Mean (SD)	28.0 (4.7)	28.3 (5.5)
Median (IQR)	27.7 (25.0–30.4)	27.6 (25.5–30.4)
Biopsy Gleason score		
2–6	554 (61)	1222 (73)
3 + 4	215 (24)	244 (14)
≥4 + 3	137 (15)	215 (13)
Preoperative stage:		
T ₁	573 (64)	1285 (80)
T ₂ –T ₃	324 (36)	329 (20)
Preoperative PSA (ng/mL)		
Mean (SD)	7.9 (7.5)	6.7 (5.2)
Median (IQR)	6.2 (4.5–9.0)	5.5 (4.2–7.5)
Postoperative Gleason score		
2–6	360 (40)	816 (46)
3 + 4	389 (43)	621 (35)
≥4 + 3	159 (17)	349 (19)
Extracapsular extension	165 (18)	434 (24)
Positive margins	365 (40)	564 (31)

Variable	SEARCH (Development) No. (%)	DPC (Validation) No. (%)
Seminal vesicle invasion	40 (4)	108 (6)
Prostate weight (g)		
Mean (SD)	44 (24)	44 (20)
Median (IQR)	38 (31–51)	40 (32–52)
Follow up (months)		
Median (IQR)	41 (21–65)	45 (22–76)

IQR, interquartile range; PSA, prostate-specific antigen; SD, standard deviation.

Table 2

Concordance indexes of various models for predicting biochemical recurrence in the Shared Equal Access Regional Cancer Hospital (SEARCH) cohort with and without prostate-specific antigen (PSA) nadir

Model	Concordance index		Difference	Patients	Recurrences
	Without PSA nadir	With PSA nadir			
Preoperative models					
d'Amico	0.650	0.737	0.087	866	169
CAPRA	0.696	0.755	0.059	790	152
Preoperative Kattan	0.677	0.745	0.068	843	166
<i>Average</i>	<i>0.674</i>	<i>0.746</i>	<i>0.07</i>	<i>833</i>	<i>162</i>
Postoperative models					
Postoperative Kattan	0.705	0.764	0.059	812	152
DPC	0.716	0.772	0.056	829	159
CPDR	0.674	0.6740	0.066	859	167
JHH	0.691	0.756	0.065	827	156
<i>Average</i>	<i>0.697</i>	<i>0.758</i>	<i>0.062</i>	<i>832</i>	<i>159</i>
<i>Average</i>	<i>0.687</i>	<i>0.753</i>	<i>0.066</i>	<i>832</i>	<i>160</i>

CAPRA, Cancer of the Prostate Risk Assessment; CPDR, Center for Prostate Disease Research; DPC, Duke Prostate Center; JHH, Johns Hopkins Hospital.

Table 3

Multivariable predictors of biochemical recurrence in the Shared Equal Access Regional Cancer Hospital (SEARCH) cohort

Variable	HR	95%CI	P
PSA nadir (per 0.01 ng/mL)	1.14	1.10–1.18	<0.001
Age (years)	1.01	0.99–1.04	0.382
Race			
White	<i>ref</i>	–	–
Non-white	1.37	1.02–1.97	0.035
Body mass index (log[kg/m ²])	3.02	1.15–7.92	0.025
Year of surgery (years)	0.96	0.89–1.03	0.221
Preoperative PSA (log[ng/mL])	1.25	0.99–1.59	0.066
Clinical T stage			
T ₁	<i>ref</i>	–	–
T ₂ –T ₃	1.17	0.85–1.61	0.333
Pathological Gleason score			
2–6	<i>ref</i>	–	–
3 + 4	2.04	1.33–3.15	0.001
≥4 + 3	2.75	1.71–4.43	<0.001
Surgical findings			
OC and M-	<i>ref</i>	–	–
PSM or ECE	1.99	1.39–2.84	<0.001
SVI	2.68	1.44–4.97	0.002
Prostate weight (log[g])	0.72	0.47–1.11	0.137

CI, confidence interval; ECE, extracapsular extension; HR, hazard ratio; OC and M-, organ-confined and margin-negative; PSA, prostate-specific antigen; PSM, positive surgical margins; SVI, seminal vesicle invasion.

Table 4

Concordance indexes of various models for predicting biochemical recurrence among patients with undetectable and detectable prostate-specific antigen (PSA) nadir in the Shared Equal Access Regional Cancer Hospital (SEARCH) cohort

Model	Undetectable PSA nadir		Detectable PSA nadir			
	CI	N	R	CI	N	R
Preoperative models						
d'Amico	0.635	607	96	0.618	275	78
CAPRA	0.687	562	85	0.671	242	72
Preoperative Kattan	0.660	593	94	0.649	266	78
Postoperative models						
PSA nadir alone	–	–	–	0.745	289	69
Postoperative Kattan	0.729	572	87	0.708	254	68
DPC	0.706	591	92	0.731	252	70
CPDR	0.666	607	97	0.684	267	74
JHH	0.695	577	88	0.694	265	72
PSA nadir nomogram	0.720	632	89	0.790	289	69

CAPRA, Cancer of the Prostate Risk Assessment; CI, concordance index; CPDR, Center for Prostate Disease Research; DPC, Duke Prostate Center; JHH, Johns Hopkins Hospital; N, number of subjects; R, number of recurrences.

Table 5

Concordance indexes of various models for predicting biochemical recurrence in the Duke Prostate Cancer (DPC) cohort

Model	Concordance Index	Patients	Recurrences
Preoperative models			
d'Amico	0.658	1603	188
CAPRA	0.659	1076	103
Preoperative Kattan	0.700	1478	173
Postoperative models			
Postoperative Kattan	0.788	805	108
DPC	0.770	1647	195
CPDR	0.725	1663	197
JHH	0.752	860	113
PSA nadir nomogram	0.778	1334	171

CAPRA, Cancer of the Prostate Risk Assessment; CPDR, Center for Prostate Disease Research; DPC, Duke Prostate Center; JHH, Johns Hopkins Hospital.