Predictive Analytics for Optimal Detection of Metastatic Prostate Cancer

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May 16, 2018



Imaging is used to detect bone and lymph node metastasis

Bone Scan (BS)

- Time-consuming procedure (3-4 hours)
- Costs \$600 \$1,000

Computed Tomography (CT)

- Exposes patient to 60 times more radiation than an x-ray
- Costs \$300 \$1,500



Significant harms associated with missing a case of metastatic cancer

 Negative health outcomes due to delays in chemotherapy



 Missed diagnoses subject patients to unnecessary treatments (e.g., radical prostatectomy) that cause serious side effects Avoidance of imaging in low-risk PCa top priority for AUA "Choosing Wisely" campaign

 Potentially harmful radiation exposure



An initiative of the ABIM Foundation

- Incidental findings that require painful and risky follow-up procedures (e.g., bone biopsy)
- Blocks access to imaging resources for other patients and unnecessarily increases healthcare costs

Conflicting imaging guidelines for PCa staging

European Urological Association (EAU)

- American Urological Association (AUA)
- National Comprehensive Cancer Network (NCCN)
- Briganti's classification and regression tree (CART)*

*A. Briganti, N. Passoni et al. "When to perform bone scan in newly diagnosed prostate cancer: External validation of the currently available guidelines and proposal of a novel risk stratification tool" *European Urology*, 57 (4), 2010.

Research objective

To determine which patients should receive a BS and/or a CT scan and which patients can safely avoid imaging on the basis of individual risk factors.





Physician-led, statewide collaborative

- 43 urology practices from throughout Michigan (> 90% of urologists in the state)
- Complete preoperative data for men with newly-diagnosed PCa were retrospectively reviewed

Methodology

The methodological approach consists of

- Development and validation of predictive models
- Correction for the bias due to missing data
- Classification modeling for the detection of metastasis



Risk Prediction Models for Metastatic Prostate Cancer

Can we develop predictive models that are-well calibrated to provide reliable predictions for newly-diagnosed MUSIC patients?

Predictive modeling

Multivariate logistic regression determines the probability of a positive BS and CT scan as a function of several covariates:

Age

- Prostate-specific antigen (PSA) (ng/ml)
- Clinical tumor stage (e.g., T1a/b/c, T2a/b/c and T3/4)
- Gleason score (GS)
- Percentage of biopsy positive cores over total number of cores taken

Development sample characteristics

For BS:

- March 2012 June 2013
- 416 patients received BS, 48~(11.5%) were positive

For CT scan:

- March 2012 September 2013
- 643 patients received CT scan, 62 (9.6%) were positive

Predictors of metastatic disease

	BS Univariable logistic regression model		CT scan Univariable logistic regression model	
Variables	OR (95% CI)	p-value	OR (95% CI)	p-value
Age at diagnosis (year) In(PSA+1), ng/mL	$\begin{array}{c} 1.04 \ (1.01 - 1.08) \\ 2.25 \ (1.76 - 2.88) \end{array}$	0.01 < 0.0001	$\begin{array}{c} 1.02 \; (0.99 - 1.05) \\ 2.79 \; (2.21 - 3.54) \end{array}$	0.02 < 0.0001
Biopsy Gleason score, No. (%)				
$\leq 3 + 4$	Reference		Reference	
4 + 3	5.04(0.90 - 28.31)	0.07	15.49(1.84 - 130.48)	0.01
8 - 10	16.05(3.82 - 67.45)	0.0002	50.69(6.96 - 369.16)	< 0.0001
Clinical T stage, No. (%)				
T1	Reference		Reference	
T2	2.64(1.31 - 5.33)	0.007	2.05(1.09 - 3.86)	0.03
T3/4	9.19(3.51 - 24.03)	< 0.0001	21.05(9.52 - 46.56)	< 0.0001
Positive cores, %	13.32(4.26 - 41.72)	< 0.0001	35.08(12.06 - 102.03)	< 0.0001

Statistical validation

Internal validation

 Boot strapping using the development sample to estimate optimism

External validation

 Independent datasets were used to validate the predictive models

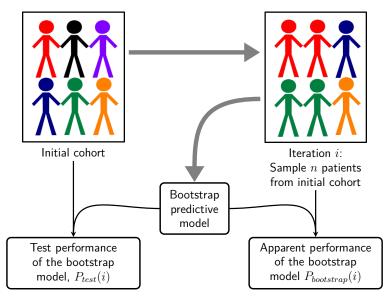
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Iteration i: Sample n patients from initial cohort



 $\begin{array}{l} \mbox{Estimate the optimism:} \\ o(i) = P_{bootstrap}(i) - P_{test}(i) \end{array} \end{array}$

Using bootstrapping to correct for optimism bias

After m iterations of bootstrapping, we can estimate the expected optimism:

$$\mathsf{Optimism} = \frac{\sum\limits_{i=1}^{m} o(i)}{m}$$

This optimism estimate can update the apparent performance of our model:

$$P_{validated} = P_{apparent} - \mathsf{Optimism}$$

Performance measures

Discrimination

 How well can the model differentiate between patients with positive and negative imaging results?

Calibration

- How reliable are the predicted risks?

Calibration

- Calibration slope is equal to one in the development sample
- In an external dataset, the calibration slope (β_{calibration}) is estimated using a logistic regression model with the linear predictor as the only explanatory variable:

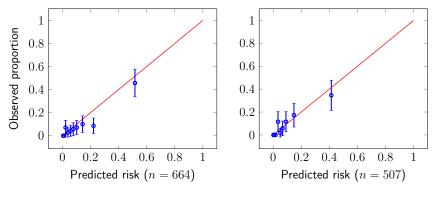
$$\log \frac{\mathbb{P}(\text{Disease present})}{\mathbb{P}(\text{Disease not present})} = \alpha + \beta_{calibration} \operatorname{LP}_{i}$$

Agreement between internal and external validation estimates

	Internal Validation		External Validation	
	Bone scan	CT scan	Bone scan	CT scan
	(n = 416)	(n = 643)	(n = 664)	(n = 507)
ROC area	0.82	0.87	0.81	0.86
Brier score	0.080	0.060	0.068	0.061
Calibration slope	0.86	0.90	0.99	0.94

Performance measures were found by applying the predictive models fit in the development samples to the external validation samples.

Predictive models are well-calibrated to external data sets



BS model

CT scan model

Bias-corrected Performance of Imaging Guidelines

How can we account for the systematic bias as not all men with newly-diagnosed prostate cancer received imaging?

Verification bias

- G+ and G- indicate whether a patient is recommended to receive imaging or not based on guideline G
- Unadjusted sensitivity and specificity are estimated based only on patients who received imaging tests:

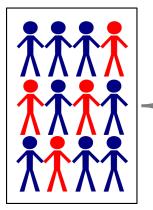
Sensitivity = $\mathbb{P}(G+ | \text{Disease present})$

Specificity = $\mathbb{P}(G - | \text{Disease not present})$

 Not all patients in our cohort received imaging, which leads to verification bias

Verification bias

Entire patient population



Patients who received imaging



Patients who did not receive imaging



Verification bias correction

We used our logistic regression model to estimate sensitivity and specificity based on the entire patient population:

$$\mathbb{P}(G+ \mid \text{Disease present}) = \frac{\mathbb{P}(\text{Disease present} \mid G+)\mathbb{P}(G+)}{\mathbb{P}(\text{Disease present})}$$
$$\mathbb{P}(G- \mid \text{Disease not present}) = \frac{\mathbb{P}(\text{Disease not present} \mid G-)\mathbb{P}(G-)}{\mathbb{P}(\text{Disease not present})}$$

Main assumption: factors considered by the guideline are the only factors that influence imaging decisions

Begg, C. B., Greenes, R. A. Assessment of diagnostic tests when disease verification is subject to selection bias. *Biometrics*, 39:207, 1983.

Verification bias greatly impacts apparent performance of imaging guidelines

	Uncorrected		Bias-corrected	
Clinical guidelines	Sensitivity	Specificity	Sensitivity	Specificity
Bone scan				
EAU	97.9	33.4	84.5	75.7
AUA	97 .9	43.5	81.2	82.0
NCCN	97.9	40.8	82.3	80.9
Briganti's CART	89.6	45.4	79.3	83.3
CT scan				
EAU	98.4	36.5	89.9	74.4
AUA	96 .8	49.2	87.2	82.5

The numbers are the percentages. EAU: European Urological Association; AUA: American Urological Association; NCCN: National Comprehensive Cancer Network; CART: classification and regression tree.

Classification Modeling for Metastatic Cancer Detection

Can we design imaging guidelines using machine learning methods that can outperform the published guidelines?

Classification models

- Two important challenges:
 - Learning from unlabeled data
 - In practice not all patients receive imaging at diagnosis
 - Learning from imbalanced data
 - A minority of patients has metastatic cancer
- To address these challenges, two machine learning paradigms are combined:
 - Semi-supervised learning
 - Cost-sensitive learning

Cost-sensitive Laplacian Kernel Logistic Regression (Cos-LapKLR)

$$f^* = \operatorname{argmin}_{f \in \mathcal{H}} \frac{1}{l} \sum_{i=1}^{l} \delta \mathbb{1}_{\{y_i=1\}} \log \left(1 + e^{-f(\mathbf{x}_i)}\right) + (1 - \delta) \mathbb{1}_{\{y_i=-1\}} \log \left(1 + e^{f(\mathbf{x}_i)}\right)$$
$$+ \gamma_{\mathcal{H}} \|f\|_{\mathcal{H}}^2 + \gamma_{\mathcal{M}} \mathbf{f}^{\mathbf{T}} \mathbf{L} \mathbf{f}$$

where f is the decision function, $f^*(\mathbf{x}) = \sum_{i=1}^{l+u} \alpha_i^* \mathbf{K}(\mathbf{x}_i, \mathbf{x})$, u the number of unimaged patients, \mathbf{K} the positive definite kernel function and \mathbf{L} the Laplacian matrix.

Cost-sensitive Laplacian Kernel Logistic
Regression (Cos-LapKLR)

$$f^* = \operatorname{argmin}_{f \in \mathcal{H}} \frac{1}{i} \sum_{i=1}^{l} \delta \mathbb{1}_{\{y_i=1\}} \log (1 + e^{-f(\mathbf{x}_i)}) + (1 - \delta) \mathbb{1}_{\{y_i=-1\}} \log (1 + e^{f(\mathbf{x}_i)})$$

 $+ \gamma_{\mathcal{H}} ||f||_{\mathcal{H}}^2 + \gamma_{\mathcal{M}} \mathbf{f}^{\mathbf{T}} \mathbf{L} \mathbf{f}$

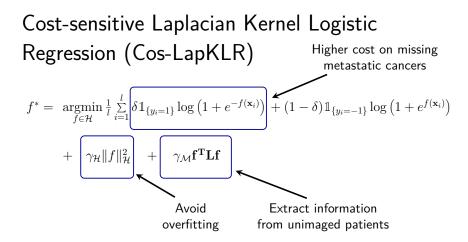
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 $+ \gamma_{\mathcal{H}} ||f||_{\mathcal{H}}^2 + \gamma_{\mathcal{M}} \mathbf{f}^{\mathbf{T}} \mathbf{L} \mathbf{f}$
Avoid
overfitting

where f is the decision function, $f^*(\mathbf{x}) = \sum_{i=1}^{l+u} \alpha_i^* \mathbf{K}(\mathbf{x}_i, \mathbf{x})$, u the number of unimaged patients, \mathbf{K} the positive definite kernel function and \mathbf{L} the Laplacian matrix.

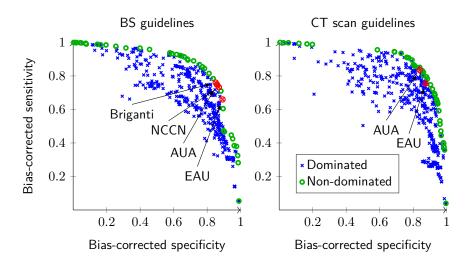


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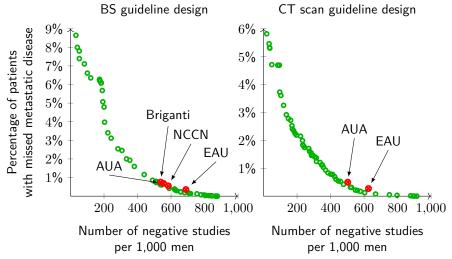
Alternative classification models

- Several other classification models adapted for imbalanced data learning were implemented:
 - Cost-sensitive logistic regression and support vector machines
 - Random forests and AdaBoost combined with advanced resampling techniques
- The diagnostic accuracy of guidelines developed from classification models was corrected for verification bias

Published guidelines are near-Pareto optimal



Published guidelines are close to the efficient frontier with missed metastasis rate <1%



Impact of recommendations if implemented

$\label{eq:Bone scan} \begin{split} & Bone \; scan \\ & PSA > 20 \; \mathrm{or} \; GS > 7 \end{split}$	CT scan PSA > 20 or GS > 7 or clinical T stage \ge T3
 20.7% (prev 27%) of patients would be scanned Of those, 17.0% (prev 12%) would be positive 	 22.6% (prev 27%) of patients would be scanned Of those, 14.3% (prev 10%) would be positive
 Estimated 0.8% of	 Estimated 0.4% of
patients have missed	patients have missed
metastatic disease	metastatic disease
 38% negative scans would	 44% negative scans would
be avoided	be avoided

MUSIC Imaging Appropriateness Criteria instituted across Michigan

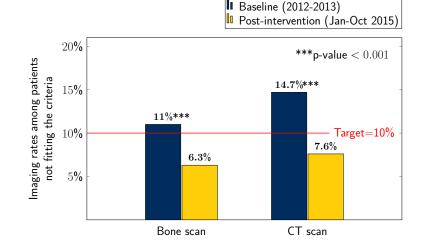
- Statewide goal of performing imaging in > 95% of patients that meet the criteria and in < 10% of those that do not
- MUSIC members were provided with a toolkit including placards with the criteria and explanations for patients



MUSIC Imaging Appropriateness Criteria

	Bone Scan	CT Scan	
	Order Bone Scan If:	Order Bone Scan If:	
PSA	> 20	> 20	
	<u>OR</u>	<u>OR</u>	
Gleason	≥ 8	≥ 8	
		OR	
Clinical T Stage		≥ cT3	
	Imaging Goals		
	 Perform imaging in ≥ 95% of patients meeting criteria Perform imaging in < 10% of patient NOT meeting criteria 		

MUSIC achieved state-wide decrease in utilization of unnecessary imaging tests



Project outcomes

- Reduction in harm to patient health from reduced radiation exposure, fewer unnecessary follow-up procedures, and decreased patient anxiety
- MUSIC collaborative saved more than \$262,000 in 2015 through reducing unnecessary imaging tests
- AskMUSIC web tool for predictive models built from MUSIC data

Project outcomes

Selin Merdan, Christine Barnett, David C. Miller, James E. Montie, Brian T. Denton. Data Analytics for Optimal Detection of Metastatic Prostate Cancer, (preprint available at http://www-personal.umich.edu/~smerdan/).

Selin Merdan, Paul R. Womble, David C. Miller, Christine Barnett, Zhu Ye, Susan M. Linsell, James E. Montie, Brian T. Denton. Toward better use of bone scans among men with early-stage prostate cancer. *Urology*.

Rachel Risko, **Selin Merdan**, Paul R. Womble, Christine Barnett, Zhu Ye, Susan M. Linsell, James E. Montie, David C. Miller, Brian T. Denton. Clinical predictors and recommendations for staging computed tomography scan among men with prostate cancer. *Urology*. Acknowledgments Selin Merdan, PhD Christine Barnett, PhD Susan Linsell, MSHA Karandeep Singh, MD James E. Montie, MD David C. Miller, MD Rachel Risko, BSE Michigan Urological Surgery Improvement Collaborative



Thank you.

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Sensitivity and specificity of imaging tests

Bone scan

- Sensitivity of 86%
- Specificity of 81%
- CT scan
 - Sensitivity of 42%
 - Specificity of 82%

State-wide increase in the utilization of imaging tests for high-risk patients

- An increase in the use of imaging tests in patients that meet the criteria from
 - 82% to 84% for BS
 - 74% to 77% for CT scan
- The MUSIC consortium has made measurable improvements in a short period of time and additional increases are anticipated

Compliance with MUSIC Imaging Criteria

- Our results were presented at collaborative-wide meetings with clinical champions who returned to their practices to present the results to their own practice group
- MUSIC members were provided with a toolkit including placards with the criteria and explanations for patients
- Members received comparative performance feedback that detailed how well their practice patterns correlated with the MUSIC Imaging Criteria

$\mathbb{P}(G+\midDisease \ present)$	=	$\frac{\mathbb{P}(Disease \ present \mid G+)\mathbb{P}(G+)}{\mathbb{P}(Disease \ present)}$
$\mathbb{P}(G-\midDisease not present)$	=	$\frac{\mathbb{P}(Disease not present \mid G-)\mathbb{P}(G-)}{\mathbb{P}(Disease not present)}$

$\mathbb{P}(G+\midDisease \ present)$	=	$\frac{\mathbb{P}(Disease \ present \mid G+)\mathbb{P}(G+)}{\mathbb{P}(Disease \ present)}$
$\mathbb{P}(G-\midDisease not present)$	=	$\frac{\mathbb{P}(Disease not present \mid G-)\mathbb{P}(G-)}{\mathbb{P}(Disease not present)}$

- \blacksquare Separate the entire population into G+ and G-
- Develop logistic regression model among patients who received imaging to estimate the probability of metastatic disease for every patient

$$\begin{split} \mathbb{P}(\mathsf{G}+\mid\mathsf{Disease present}) &= \frac{\mathbb{P}(\mathsf{Disease present}\mid\mathsf{G}+)\mathbb{P}(\mathsf{G}+)}{\mathbb{P}(\mathsf{Disease present})} \\ \mathbb{P}(\mathsf{G}-\mid\mathsf{Disease not present}) &= \frac{\mathbb{P}(\mathsf{Disease not present}\mid\mathsf{G}-)\mathbb{P}(\mathsf{G}-)}{\mathbb{P}(\mathsf{Disease not present})} \end{split}$$

$$\begin{split} \mathbb{P}(\mathsf{G}+\mid\mathsf{Disease present}) &= \frac{\mathbb{P}(\mathsf{Disease present}\mid\mathsf{G}+)\mathbb{P}(\mathsf{G}+)}{\mathbb{P}(\mathsf{Disease present})} \\ \mathbb{P}(\mathsf{G}-\mid\mathsf{Disease not present}) &= \frac{\mathbb{P}(\mathsf{Disease not present}\mid\mathsf{G}-)\mathbb{P}(\mathsf{G}-)}{\mathbb{P}(\mathsf{Disease not present})} \end{split}$$

$$\begin{split} \mathbb{P}(\mathsf{Disease present}) & = & \mathbb{P}(\mathsf{Disease present}|\mathsf{G}+)\mathbb{P}(\mathsf{G}+) + \\ & & \mathbb{P}(\mathsf{Disease present}|\mathsf{G}-)\mathbb{P}(\mathsf{G}-) \end{split}$$

$$\begin{split} \mathbb{P}(\text{Disease not present}) \;\;=\;\; \mathbb{P}(\text{Disease not present}|\mathsf{G}+)\mathbb{P}(\mathsf{G}+)+\\ \mathbb{P}(\text{Disease not present}|\mathsf{G}-)\mathbb{P}(\mathsf{G}-) \end{split}$$