

Examining Improvement in Risk Prediction

Dave Glidden
24 November 2009

Risk Model

- Existing set of risk factors for disease
- Novel marker or test
genetic, proteomic, imaging, biomarker
- New marker add predictive information
- How much better are predictions?
do they aid in clinical decision making

Examples

- Breast density for breast cancer risk
Tice et al. 2008
- C-reactive protein for CV risk
Cook et al. 2006
- PSA for prostate cancer risk

The Easy Question

and least interesting

- Is there a statistically significant contribution of the new measure/marker?
- Fit a logistic/Cox regression model
- Examine p-value for new measure adjusting for other factors
- Gives p-value
- Implies better prediction

Why not interesting?

- Turns out to be an easy threshold
- Many factors can reach this
- Always need to know how much better we do?
- Relevant summary needed

Measure of Prediction

- TPF = true positive fraction
proportion of case “positive”
aka: sensitivity
- FPF = false positive fraction
proportion of control “positive”
- Trade-off
- Two kinds of errors

Context is Important

- Prostate cancer screening:
FPF = 10%, TPF = 50%,
prevalence = 5% => PVP = 21%
- Metastatic disease in breast cancer
TPF = 92%, FPF = 42%
- What are implications of false positive, false negative?

Odds Ratio

- Suppose FPF = 10%
good specificity (90%)
- Suppose OR = 3.0
pretty hefty OR
- Implies TPF = 25%
very low specificity = 25%

Odds Ratio

- FPF = 0.10
90% specificity
- TPF = 0.80
80% sensitivity
- $OR = (0.80/0.20) / (0.10/0.90) = 36.0$
- Huge OR associated with good discrimination

Dist of Marker and OR

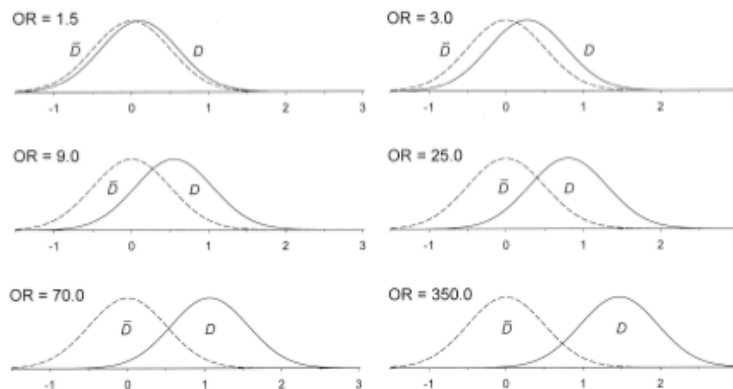


FIGURE 2. Probability distributions of a marker, X , in cases (solid curves) and controls (dashed curves) consistent with the logistic model $\log\{P(D=1|X)\} = \alpha + \beta X$. It has been assumed that X has a mean of 0 and a standard deviation of 0.5 in controls so that a unit increase represents the difference between the 84th and 16th percentiles of X in controls. The marker is normally distributed, with the same variance in cases. The odds ratio (OR) per unit increase in X is shown.

ROC Curves

- Can be problematic for risk
- Don't display absolute risk values
- Two tests graphed =>
FPF v. TPF (varying threshold)
- C-statistic doesn't relate to clinical decisions
measure ability to classify

Tice at al 2008

- Breast cancer risk
- SEER data: 1,095,484 women
- age, race/ethnicity, family hx, biopsy hx
- added value of breast density
HR 4.2 (less than 65), 2.2 (65 or older)
statistically significant
- 5 year risk model (Cox model)
- How much does density add to risk assessment?

Another Approach

- Based on results of a model, calculate risks
- Group risks in context dependent way
Tice: <1%, 1-1.6%, 1.6-2.5%, > 2.5%
- Examine calibration
- Compare between models

Model w/o Density

Strata	<1%	1.0-1.66%	1.67-2.5%	> 2.5%
N	215,402	201,972	148,795	63,105
% event	0.7%	1.3%	1.8%	2.9%

Model with Density

Strata	<1%	1.0-1.66%	1.67-2.5%	> 2.5%
N	249,959	186,106	124,420	68,744
% event	0.7%	1.3%	2.0%	3.1%

Calibration

- % events by category
- Both models well calibrated
- % events observed falls within expected
- Calibration has subjective aspect
*if model predicts 2% but is really 3%
close enough?*

NR Cook

- Compare two *nested* prediction models one includes extra predictor like breast density
- Cross-tabulate risk predictions based on model
- Examine who gets reclassified
- What % and in which direction?

Table 1. Five-Year Risks for Breast Cancer as Predicted by Models That Do and Do Not Include Breast Density*

5-Year Risk from Model without Breast Density	5-Year Risk from Model with Breast Density				Total
	0% to <1%	1% to 1.66%	1.67% to 2.5%	>2.5%	
0% to <1%					
Women, <i>n</i>	176 831	38 500	71	0	215 402
Events, <i>n</i>	1161	415	0	0	1576
Nonevents, <i>n</i>	175 670	38 085	71	0	213 826
Proportion of women with events	0.7	1.1	0.0	–	0.7
1% to 1.66%					
Women, <i>n</i>	64 297	99 456	37 149	1025	201 927
Events, <i>n</i>	526	1328	754	32	2640
Nonevents, <i>n</i>	63 771	98 128	36 395	993	199 287
Proportion of women with events	0.8	1.3	2.0	3.1	1.3
1.67% to 2.5%					
Women, <i>n</i>	8741	45 478	71 309	23 267	148 795
Events, <i>n</i>	74	609	1419	621	2723
Nonevents, <i>n</i>	8667	44 869	69 890	22 646	146 072
Proportion of women with events	0.9	1.3	2.0	2.7	1.8
>2.5%					
Women, <i>n</i>	90	2672	15 891	44 452	63 105
Events, <i>n</i>	0	38	340	1467	1845
Nonevents, <i>n</i>	90	2634	15 551	42 985	61 260
Proportion of women with events	0.0	1.4	2.1	3.3	2.9
Total					
Women, <i>n</i>	249 959	186 106	124 420	68 744	629 229
Events, <i>n</i>	1761	2390	2513	2120	8784
Nonevents, <i>n</i>	248 198	183 716	121 907	66 624	620 445
Proportion of women with events	0.7	1.3	2.0	3.1	1.4

Reclassification Index

with breast cancer

	<1%	1.0-1.66%	1.67-2.5%	> 2.5%
<1%	1161	415	0	0
1.0-1.66%	526	1328	754	32
1.67-2.5%	74	609	1419	612
> 2.5%	0	38	340	1467

Reclassification

- Risk strata same: 5375 (61%)
- Risk strata lower w/ dens 1587 (18%)
wrong direction
- Risk strata higher w/ dens 1813 (21%)
correct direction
- Net reclassification = 21%-18% = 3%
in the right direction

Reclassification Index

no breast cancer

	<1%	1.0-1.66%	1.67-2.5%	> 2.5%
<1%	175,670	38,085	71	0
1.0-1.66%	63,771	98,128	36,395	993
1.67-2.5%	8,667	44,869	69,890	22,646
> 2.5%	90	2,634	1,551	42,985

Reclassification

- Risk strata same: 386,673 (64%)
- Risk strata lower w/ dens 121,582 (20%)
right direction
- Risk strata higher w/ dens 98,190 (16%)
wrong direction
- Net reclassification = 20% - 16% = 3.8%
in the right direction

Plots

- Can avoid cutpoints
- x axis: pred. prob with breast density
- y axis: pred. prob w/o breast density
- plot separately for cases and non-cases

Cautions

- Method doesn't work well in non-nested models
- Nested: one model has all predictors in other
- Can't be applied to case-control data
not possible to estimate "risk"

Risk Reclassification

- Requires risk strata
4 strata is typical
arrived in a context-dependent way
- Shows calibration
- Direct measure of change due to adding marker
- See proportion reclassified

Diagnostic Likelihood Ratios

- DLR: familiar from clinical epi
- Describes how risk of disease is modified by new information
- $\text{pr}(\text{disease} | x) / \text{pr}(\text{disease})$
x is some diagnostic test information
DLR is function of x
- $\text{DLR} > 1$, risk increased
- $\text{DLR} < 1$, risk decreased

Our scenario

- D: disease variable (D=1, has disease)
- X: an established risk factor for disease or even a risk score
- Y: a novel factor
how much does it add?
- Extend the concept of a DLR

DLR

- $DLR(Y) = P(D=1 | X, Y) / P(D=1, X)$
- the ratio of risk given X + Y to X alone
- a function of Y
- How does this differ from an odd ratio for Y?

Gu and Pepe

- $\log(p|X) = \alpha_0 + \alpha_1 * X$
- $\log(p|X,Y) = \beta_0 + \beta_1 * X + \beta_2 * Y$
- Then
 $\log\{DLR(Y)\} = (\beta_0 - \alpha_0) + (\beta_1 - \alpha_1) * X + \beta_2 * Y$
- CI for this is complicated
- But easily graphed

Physician's Health Study

- Randomized clinical trial in 1980s
- 430 men developed prostate cancer
- Nested case control study on stored specimens
- Total PSA and % free PSA

Total PSA alone

```
. logistic d tpsa, coef
```

```
Logistic regression                Number of obs   =      683
                                   LR chi2(1)         =      187.10
                                   Prob > chi2        =      0.0000
Log likelihood = -342.10876         Pseudo R2      =      0.2147
```

d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
tpspsa	.3533952	.0399274	8.85	0.000	.2751389	.4316514
_cons	-1.886684	.1481591	-12.73	0.000	-2.177071	-1.596298

Total PSA alone

```
. logistic d fpsa tpsa, coef
```

```
Logistic regression                Number of obs   =      683
                                   LR chi2(2)        =      189.43
                                   Prob > chi2        =      0.0000
Log likelihood = -340.94405         Pseudo R2      =      0.2174
```

d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
fpspsa	-.3493467	.0829424	-4.21	0.000	-.5119108	-.1867826
tpspsa	.4037308	.0471778	8.56	0.000	.311264	.4961976
_cons	-1.843289	.1438261	-12.82	0.000	-2.125183	-1.561395

log DLR

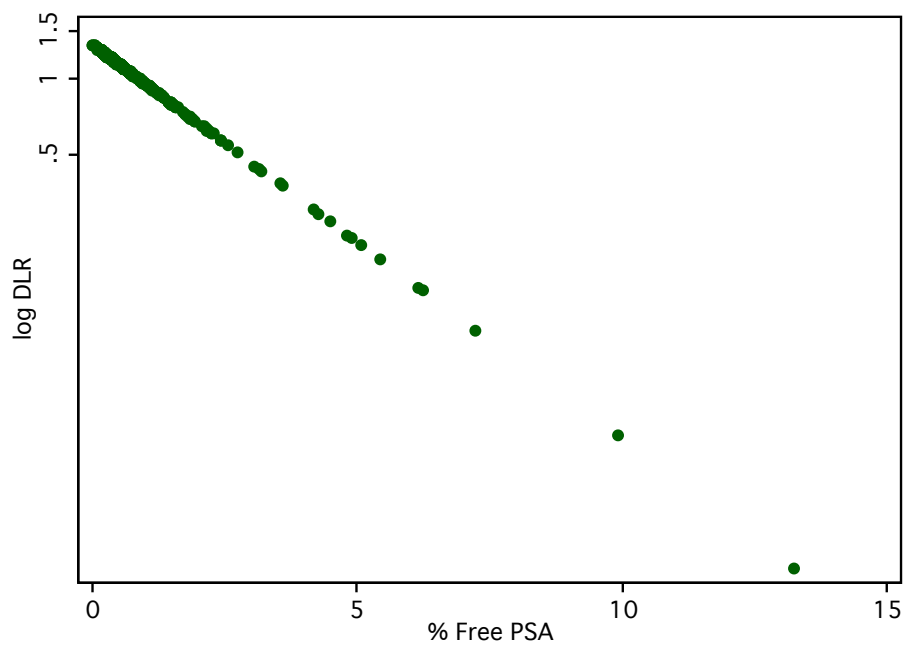
$$\log\{\text{DLR}(Y)\} = (\beta_0 - \alpha_0) + (\beta_1 - \alpha_1)*X + \beta_2*Y = \\ 0.05 + 0.05I*TPSA - 0.35*FPSA$$

Graph this at average TPSA =4.663

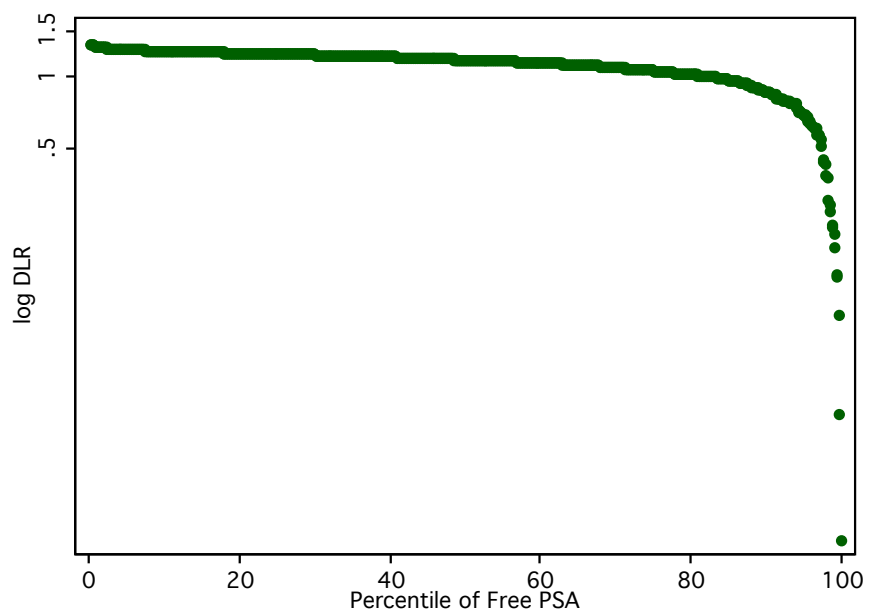
Stata

- `gen log_dlr=0.05+0.05I*4.663-.35*fpsa`
- `gen dlr = exp(dlr)`
- `twoway (scatter dlr fpsa), ytitle(log DLR) yscale(log) xtitle(% Free PSA)`

DLR by % FPSA



DLR by % FPSA



Valued added

- High values indicate decreased risk
- 81% have $DLR > 1$
lowest 81% of FPSA indicate increased risk
- High FPSA can indicated much lower likelihood of cancer (at a given PSA)
- Maybe helpful for ruling out disease in borderline PSA

Interactions

- Value of FPSA (or any marker)
- May depend on configuration of other predictors
- Can examine this using interactions in logistic models

OR

- Risk factor may have elevated OR
- Individual risk is modified by risk factor
- Population risk may not be greatly modified
- Especially true if risk factors cluster
- Both embody some level of truth

C-statistic

- Very difficult to budge
- Measure pure ability to classify
- However, strict classification is rarely of interest
- More interested in risk stratification

Summary

- These kinds of assessments very difficult
- Must weight very difficult trade offs
- Difficult to balance false positives and false negatives
- Utilities may be very different for different people