

Lecture 5

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Table of contents

- 1 Table of contents
- 2 Outline
- 3 Conditional probability
- 4 Conditional densities
- 5 Bayes' Rule
- 6 Diagnostic tests
- 7 DLRs
- 8 2×2 tables
- 9 ROC and AUC

Outline

- 1 Define conditional probabilities
- 2 Define conditional mass functions and densities
- 3 Motivate the conditional density
- 4 Bayes' rule
- 5 Applications of Bayes' rule to diagnostic testing

Conditional probability, examples

- What is the probability for a 30 year old woman to develop breast cancer within 10 years?
- X is “develop cancer within the next 10 years”
- We would like to calculate probabilities of the type

$$P(X = 1 | \text{sex} = 1, \text{age} = 30)$$

- What happens if $\text{age} = 50$?
- What happens if the person is a man $\text{sex} = 0$?
- What one conditions on is crucial

Conditional probability, examples

- What is the probability of surviving more than 1 year for a man who is 50 years old and has an estimated glomerular filtration rate (eGFR) equal to 15?
- X is surviving time
- We would like to calculate probabilities of the type

$$P(X > 1 | \text{sex} = 0, \text{age} = 50, \text{eGFR} = 15)$$

Conditional probability, motivation

- The probability of getting a one when rolling a (standard) die is usually assumed to be one sixth
- Suppose you were given the extra information that the die roll was an odd number (hence 1, 3 or 5)
- *conditional on this new information*, the probability of a one is now one third

Conditional probability, definition

- Let B be an event so that $P(B) > 0$
- Then the conditional probability of an event A given that B has occurred is

$$P(A \mid B) = \frac{P(A \cap B)}{P(B)}$$

- Notice that if A and B are independent, then

$$P(A \mid B) = \frac{P(A)P(B)}{P(B)} = P(A)$$

Example

- Consider our die roll example
- $B = \{1, 3, 5\}$
- $A = \{1\}$

$$\begin{aligned}P(\text{one given that roll is odd}) &= P(A \mid B) \\&= \frac{P(A \cap B)}{P(B)} \\&= \frac{P(A)}{P(B)} \\&= \frac{1/6}{3/6} = \frac{1}{3}\end{aligned}$$

Conditional densities and mass functions

- Conditional densities or mass functions of one variable conditional on the value of another
- Let $f(x, y)$ be a bivariate density or mass function for random variables X and Y
- Let $f(x)$ and $f(y)$ be the associated marginal mass function or densities disregarding the other variables

$$f(y) = \int f(x, y) dx \quad \text{or} \quad f(y) = \sum_x f(x, y)$$

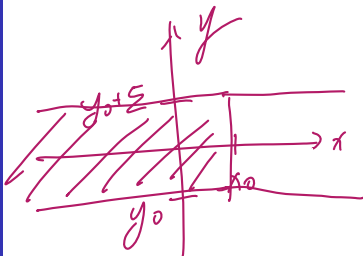
- Then the **conditional** density or mass function *given that* $Y = y$ is given by

$$f(x | y) = f(x, y) / f(y)$$

- It is easy to see that, in the discrete case, the definition of conditional probability is exactly as in the definition for conditional events where $A =$ the event that $X = x_0$ and $B =$ the event that $Y = y_0$
- The continuous definition is harder to motivate, since the events $X = x_0$ and $Y = y_0$ each have probability 0
- However, a useful motivation can be performed by taking the appropriate limits as follows
- Define $A = \{X \leq x_0\}$ while $B = \{Y \in [y_0, y_0 + \epsilon]\}$

Continued

$$P(X \leq x_0 \mid Y \in [y_0, y_0 + \epsilon]) = P(A \mid B) = \frac{P(A \cap B)}{P(B)}$$



$$= \frac{P(X \leq x_0, Y \in [y_0, y_0 + \epsilon])}{P(Y \in [y_0, y_0 + \epsilon])}$$

$$= \frac{\int_{y_0}^{y_0 + \epsilon} \int_{-\infty}^{x_0} f(x, y) dx dy}{\int_{y_0}^{y_0 + \epsilon} f(y) dy}$$

$$= \frac{\epsilon \int_{y_0}^{y_0 + \epsilon} \int_{-\infty}^{x_0} f(x, y) dx dy}{\epsilon \int_{y_0}^{y_0 + \epsilon} f(y) dy}$$

$$\begin{aligned}
 &= \frac{\int_{-\infty}^{y_0+\epsilon} \int_{-\infty}^{x_0} f(x,y) dx dy - \int_{-\infty}^{y_0} \int_{-\infty}^{x_0} f(x,y) dx dy}{\epsilon} \\
 &= \frac{\int_{-\infty}^{y_0+\epsilon} f(y) dy - \int_{-\infty}^{y_0} f(y) dy}{\epsilon} \\
 &= \frac{g_1(y_0+\epsilon) - g_1(y_0)}{\epsilon} \\
 &= \frac{g_2(y_0+\epsilon) - g_2(y_0)}{\epsilon}
 \end{aligned}$$

where

$$g_1(y_0) = \int_{-\infty}^{y_0} \int_{-\infty}^{x_0} f(x,y) dx dy \quad \text{and} \quad g_2(y_0) = \int_{-\infty}^{y_0} f(y) dy.$$

- Notice that the limit of the numerator and denominator tends to g'_1 and g'_2 as ϵ gets smaller and smaller
- Hence we have that the conditional distribution function is

$$P(X \leq x_0 \mid Y = y_0) = \frac{\int_{-\infty}^{x_0} f(x, y_0) dx}{f(y_0)}.$$

- Now, taking the derivative with respect to x yields the conditional density

$$f(x_0 \mid y_0) = \frac{f(x_0, y_0)}{f(y_0)}$$

for every x_0 and y_0 and subscript can now be dropped

Geometry

- Geometrically, the conditional density is obtained by taking the relevant slice of the joint density and appropriately renormalizing it
- This idea extends to any other linear or non-linear function

Example

- Let $f(x, y) = ye^{-xy-y}$ for $0 \leq x$ and $0 \leq y$
- Then note

$$f(y) = \int_0^{\infty} f(x, y) dx = e^{-y} \int_0^{\infty} ye^{-xy} dx = e^{-y}$$

- Therefore

$$f(x | y) = f(x, y) / f(y) = \frac{ye^{-xy-y}}{e^{-y}} = ye^{-xy}$$

- Calculate $P(X \geq 5 | Y = 3)$

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Table of
contents

Outline

Conditional
probability

**Conditional
densities**

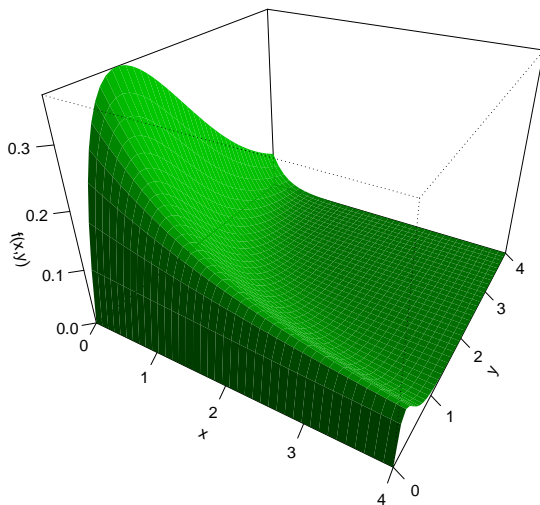
Bayes' Rule

Diagnostic
tests

DLRs

2×2 tables

ROC and AUC



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Table of
contents

Outline

Conditional
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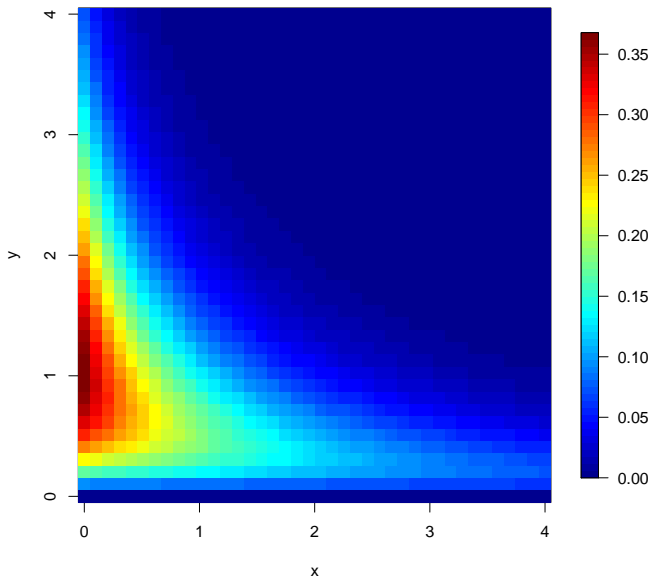
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ROC and AUC



- Check out the R functions `persp`, `image.plot`, `plot3D`, `surface3d`
- Useful packages: `rgl`, `fields`

Example

- Let $f(x, y) = 1/\pi r^2$ for $x^2 + y^2 \leq r^2$
- X and Y are uniform on a disk with radius r
- What is the conditional density of X given that $Y = 0$?
- Probably easiest to think geometrically

$$f(x \mid y = 0) \propto 1 \quad \text{for} \quad -r \leq x \leq r$$

- Therefore

$$f(x \mid y = 0) = \frac{1}{2r} \quad \text{for} \quad -r \leq x \leq r$$

Bayes' rule

Table of
contents

Outline

Conditional
probabilityConditional
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2 × 2 tables

ROC and AUC

- Let $f(x | y)$ be the conditional density or mass function for X given that $Y = y$
- Let $f(y)$ be the marginal distribution for y
- Then if y is continuous

$$f(y | x) = \frac{f(x | y)f(y)}{\int f(x | t)f(t)dt}$$

- If y is discrete

$$f(y | x) = \frac{f(x | y)f(y)}{\sum_t f(x | t)f(t)}$$

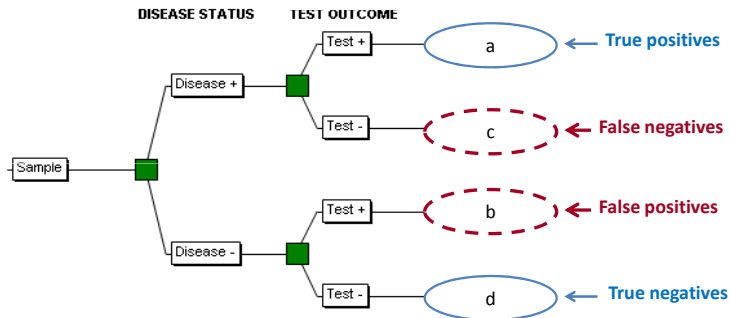
- Bayes' rule relates the conditional density of $f(y | x)$ to the conditional density $f(x | y)$ and the marginal density $f(y)$
- A special case of this kind relationship is for two sets A and B , which yields that

$$P(B | A) = \frac{P(A | B)P(B)}{P(A | B)P(B) + P(A | B^c)P(B^c)}.$$

Proof:

- Let X be an indicator that event A has occurred
- Let Y be an indicator that event B has occurred
- Plug into the discrete version of Bayes' rule

Example: diagnostic tests



Example: diagnostic tests

- Let $+$ and $-$ be the events that the result of a diagnostic test is positive or negative, respectively
- Let D and D^c be the event that the subject of the test has or does not have the disease respectively
- The **sensitivity** is the probability that the test is positive given that the subject actually has the disease, $P(+ \mid D)$
- The **specificity** is the probability that the test is negative given that the subject does not have the disease, $P(- \mid D^c)$

More definitions

- The **positive predictive value** is the probability that the subject has the disease given that the test is positive, $P(D \mid +)$
- The **negative predictive value** is the probability that the subject does not have the disease given that the test is negative, $P(D^c \mid -)$
- The **prevalence of the disease** is the marginal probability of disease, $P(D)$

More definitions

- The **diagnostic likelihood ratio of a positive test**, labeled DLR_+ , is $P(+ | D)/P(+ | D^c)$, which is the

$$sensitivity / (1 - specificity)$$

- The **diagnostic likelihood ratio of a negative test**, labeled DLR_- , is $P(- | D)/P(- | D^c)$, which is the

$$(1 - sensitivity) / specificity$$

	T^+	T^-
D^+	99.7	0.3
D^-	1.5	98.5

$$P(D) = 0.001 \quad \text{Example}$$

$$= \frac{P(D^+)}{P(D^+) + P(D^-)}$$

- A study comparing the efficacy of HIV tests, reports on an experiment which concluded that HIV antibody tests have a sensitivity of 99.7% and a specificity of 98.5%
- Suppose that a subject, from a population with a .1% prevalence of HIV, receives a positive test result. What is the probability that this subject has HIV?
- Mathematically, we want $P(D | +)$ given the sensitivity, $P(+ | D) = .997$, the specificity, $P(- | D^c) = .985$, and the prevalence $P(D) = .001$

$$\frac{0.001 \times 0.997}{0.001 \times 0.997 + 0.999 \times 0.015}$$

$$\text{已知 } T^+ \rightarrow \frac{P(D^+)}{P(D^+) + P(D^-)}$$

Using Bayes' formula

$$\begin{aligned}
 \textcircled{P(D \mid +)} &= \frac{P(+ \mid D)P(D)}{P(+ \mid D)P(D) + P(+ \mid D^c)P(D^c)} \\
 &= \frac{P(+ \mid D)P(D)}{P(+ \mid D)P(D) + \{1 - P(- \mid D^c)\}\{1 - P(D)\}} \\
 &= \frac{.997 \times .001}{.997 \times .001 + .015 \times .999} \\
 &= .062
 \end{aligned}$$

Handwritten notes in red:
 $\frac{P(+)}{P(+)}$ (above the first equation)
 $P(D \mid +)$ (circled next to the first equation)

- In this population a positive test result only suggests a 6% probability that the subject has the disease
- (The positive predictive value is 6% for this test)

More on this example

- The low positive predictive value is due to low prevalence of disease and the somewhat modest specificity
- Suppose it was known that the subject was an intravenous drug user and routinely had intercourse with an HIV infected partner
- Notice that the evidence implied by a positive test result does not change because of the prevalence of disease in the subject's population, only our interpretation of that evidence changes

Likelihood ratios

- Using Bayes rule, we have

$$P(D \mid +) = \frac{P(+ \mid D)P(D)}{P(+ \mid D)P(D) + P(+ \mid D^c)P(D^c)}$$

and

$$P(D^c \mid +) = \frac{P(+ \mid D^c)P(D^c)}{P(+ \mid D)P(D) + P(+ \mid D^c)P(D^c)}.$$

- Therefore

$$\frac{P(D \mid +)}{P(D^c \mid +)} = \frac{P(+ \mid D)}{P(+ \mid D^c)} \times \frac{P(D)}{P(D^c)}$$

ie

post-test odds of $D = DLR_+ \times$ pre-test odds of D

- Similarly, DLR_- relates the decrease in the odds of the disease after a negative test result to the odds of disease prior to the test.

HIV example revisited

- Suppose a subject has a positive HIV test
- $DLR_+ = .997 / (1 - .985) \approx 66$
- The result of the positive test is that the odds of disease is now 66 times the pretest odds
- Or, equivalently, the hypothesis of disease is 66 times more supported by the data than the hypothesis of no disease

HIV example revisited

- Suppose that a subject has a negative test result
- $DLR_- = (1 - .997)/.985 \approx .003$
- Therefore, the post-test odds of disease is now .3% of the pretest odds given the negative test.
- Or, the hypothesis of disease is supported .003 times that of the hypothesis of absence of disease given the negative test result

Comparing two tests

- Test 1: $DLR_+ = a$, Test 2: $DLR_+ = b$
- Test 1: a is the factor that multiplies the pre-test odds to obtain the post-test odds

$$\frac{P(D|T_1 = +)}{P(D_C|T_1 = +)} = a \times \frac{P(D)}{P(D_C)}$$

- Test 2: b is the factor that multiplies the pre-test odds to obtain the post-test odds

$$\begin{aligned} O(D|T_1 = +, T_2 = +) &= b \times O(D|T_1 = +) \\ &= a \times b \times O(D) \end{aligned}$$

Tests and 2×2 tables

A particularly interesting and important question today is that of testing for drugs. Suppose it is assumed that about 5% of the general population uses drugs. You employ a test that is 95% accurate, which well say means that if the individual is a user, the test will be positive 95% of the time, and if the individual is a nonuser, the test will be negative 95% of the time. A person is selected at random and is given the test. Its positive. What does such a result suggest? Would you conclude that the individual is a drug user? What is the probability that the person is a drug user?

The 2 × 2 table



	Disease +	Disease -	Total
Test +	a	b	$a + b$
Test -	c	d	$c + d$
Total	$a + c$	$b + d$	$a + b + c + d$

$PPV = P(D | +) = \frac{a}{a+b}$
 $NPV = P(\bar{D} | -) = \frac{d}{c+d}$
 $Sens = P(+ | D) = \frac{a}{a+c}$
 $Spec = P(- | \bar{D}) = \frac{d}{b+d}$

The 2×2 table: example

	Disease +	Disease -	Total	
Test +	48	47	95	PPV = 51%
Test -	2	903	905	NPV = 99%
Total	50	950	1000	

The 2×2 table: example

	Disease +	Disease -	Total	
Test +	190	40	230	PPV = 83%
Test -	10	760	770	NPV = 99%
Total	200	800	1000	

Point: PPV depends on **prior probability** of disease in the population

Prediction with binary outcomes

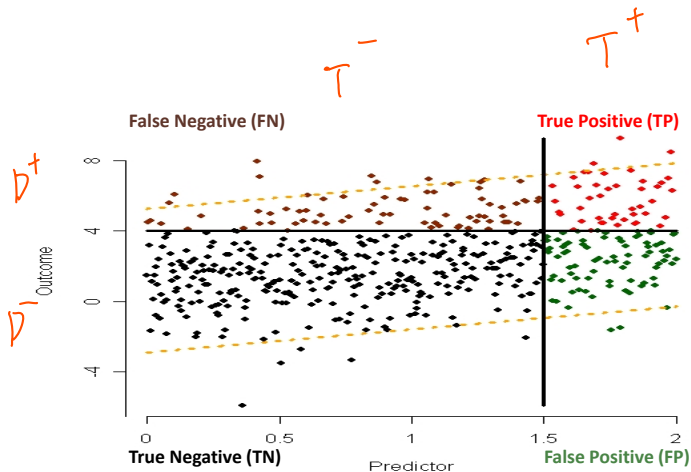
- Outcome is 0/1
- Examples
 - Non-diseased/diseased
 - Alive/Dead
 - Failure/Success (procedure)
- Continuous predictor
- Examples
 - Outcome of a diagnostic test
 - Prediction score (based on multiple characteristics)
 - Clinical score (e.g. SOFA score in ICU)

The ROC

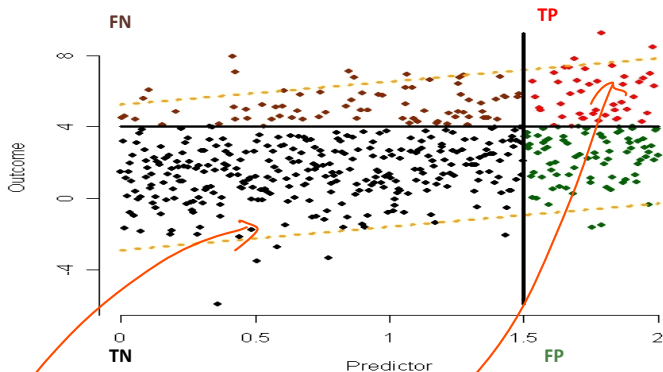
- Outcome $D \in \{0, 1\}$, X scalar predictor
- For every threshold t predict $\hat{D} = 1$ if $X > t$
- $\text{Sens}(t) = P(X > t | D = 1)$, $\text{Spec}(t) = P(X \leq t | D = 0)$
- The receiver operatic characteristic (ROC) function is

$$\{1 - \text{Spec}(t), \text{Sens}(t)\} \text{ for all } t$$

Luck, error and randomness



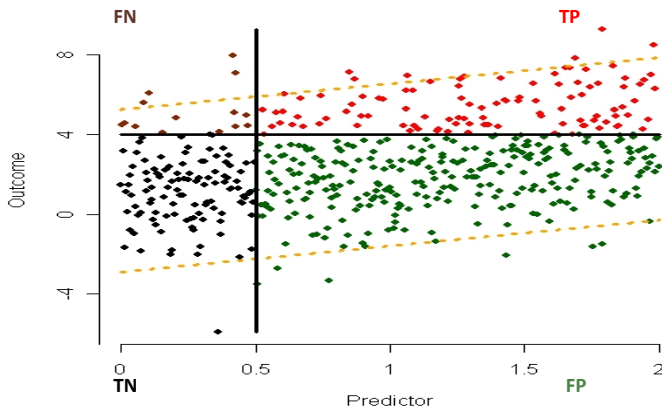
Dependence on the threshold



$$\text{Spec} = \frac{\text{TN}}{\text{TN} + \text{FP}} = \frac{301}{301 + 81} = 0.79$$

$$\text{Sens} = \frac{\text{TP}}{\text{TP} + \text{FN}} = 0.37$$

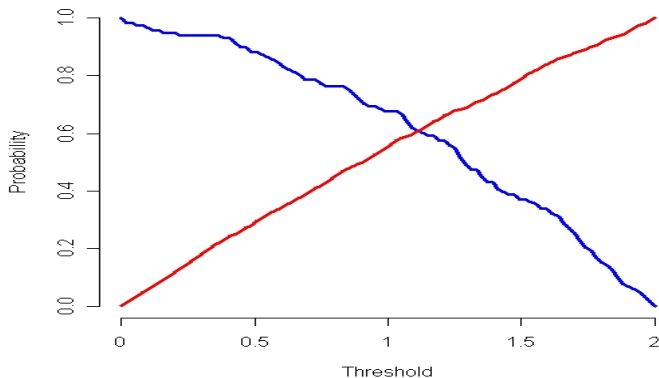
Dependence on the threshold

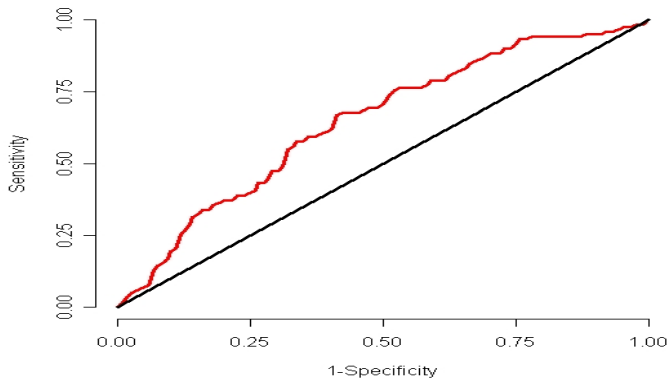


$$\text{Spec} = \frac{\text{TN}}{\text{TN} + \text{FP}} = \frac{111}{111 + 271} = 0.29$$

$$\text{Sens} = \frac{\text{TP}}{\text{TP} + \text{FN}} = 0.88$$

Sensitivity and Specificity curves



[Table of
contents](#)[Outline](#)[Conditional
probability](#)[Conditional
densities](#)[Bayes' Rule](#)[Diagnostic
tests](#)[DLRs](#)[2 × 2 tables](#)[ROC and AUC](#)

- Area under the ROC curve is denoted by AUC
- Probability that the model will assign a higher probability of an event to the subject who will experience the event than to the one who will not experience the event
- AUC is one of the main criteria for assessing discrimination accuracy
- $AUC=0.68$ in the example

AUC interpretation proof

$$\text{Sens}(t) = S(t) = P(X > t | D = 1) = \int_t^1 f(x | D = 1) dx$$

$$1 - \text{Spec}(t) = P(t) = P(X > t | D = 0) = \int_t^1 f(x | D = 0) dx$$

$$\text{AUC} = \int_0^1 S(t) \frac{d}{dt} P(t) dt = \int_0^1 S(t) f(t | D = 0) dt$$

$$= P(X_i > X_j | D_i = 1, D_j = 0)$$

Note that $f(x_i, x_j | D_i = 1, D_j = 0) = f(x_i | D_i = 1) f(x_j | D_j = 0)$

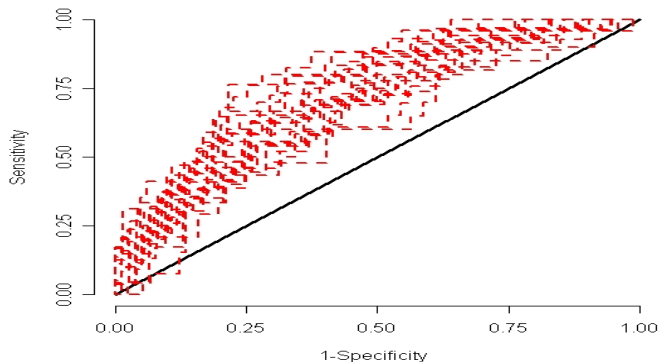
Some comments

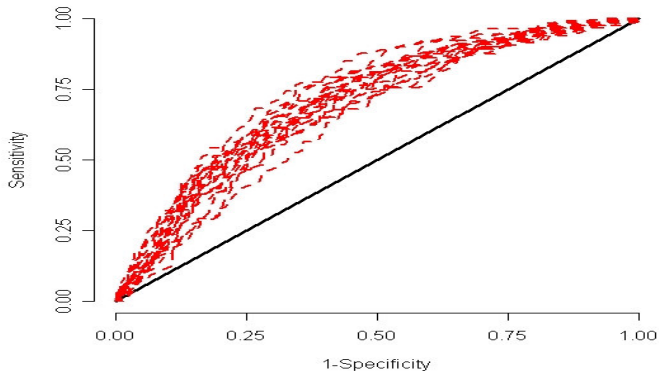
- ROC, AUC are never observed
- They are estimated based on a data set
- They have statistical variability
- Variability is controlled by the amount of data
- Important fact: more data improves the precision of the ROC and AUC estimators. It does not improve prediction!

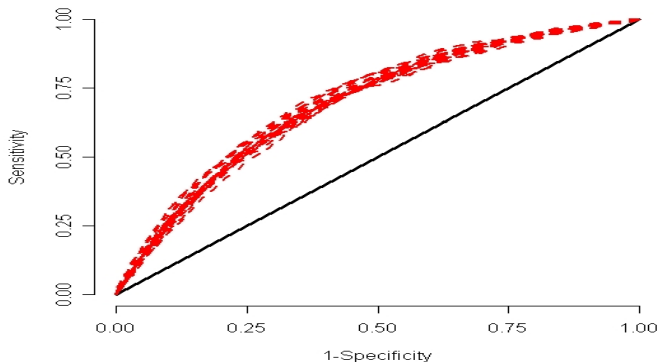
Bootstrapping ROCs and AUCs

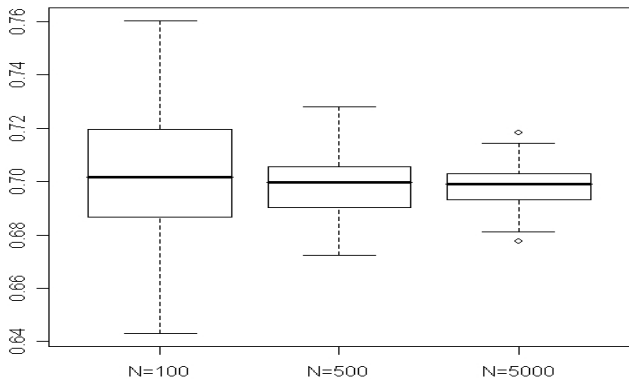
- Have a method for estimating ROC, AUC from data
- Bootstrap subjects nonparametrically (say 10,000 times)
- Repeat the estimation procedure for each data set
- Report the bootstrap distribution of ROCs and AUCs

```
for (i in 1:10000)  
  {boot<-sample(n,replace=TRUE)}
```

[Table of
contents](#)[Outline](#)[Conditional
probability](#)[Conditional
densities](#)[Bayes' Rule](#)[Diagnostic
tests](#)[DLRs](#)[2 × 2 tables](#)[ROC and AUC](#)

[Table of
contents](#)[Outline](#)[Conditional
probability](#)[Conditional
densities](#)[Bayes' Rule](#)[Diagnostic
tests](#)[DLRs](#)[2 × 2 tables](#)[ROC and AUC](#)

[Table of
contents](#)[Outline](#)[Conditional
probability](#)[Conditional
densities](#)[Bayes' Rule](#)[Diagnostic
tests](#)[DLRs](#)[2 × 2 tables](#)[ROC and AUC](#)



- Variability can be very large even for large data sets
- Variability can be mistaken for signal
- This can lead to spurious, irreproducible results

“As reviewer of grants dedicated to discovery of novel biomarkers, I cannot believe how often the emphasis is on p-values (statistical significance) and not on predictive measures (predictive performance)”