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# 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

$$\begin{aligned}
 P(X \leq x_0 \mid Y \in [y_0, y_0 + \epsilon]) &= P(A \mid B) = \frac{P(A \cap B)}{P(B)} \\
 &= \frac{P(X \leq x_0, Y \in [y_0, y_0 + \epsilon])}{P(Y \in [y_0, y_0 + \epsilon])} \\
 &= \frac{\int_{y_0}^{y_0 + \epsilon} \int_{-\infty}^{x_0} f(x, y) dx dy}{\int_{y_0}^{y_0 + \epsilon} f(y) dy} \\
 &= \frac{\epsilon \int_{y_0}^{y_0 + \epsilon} \int_{-\infty}^{x_0} f(x, y) dx dy}{\epsilon \int_{y_0}^{y_0 + \epsilon} f(y) dy}
 \end{aligned}$$

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

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

## Lecture 5

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**Conditional  
densities**

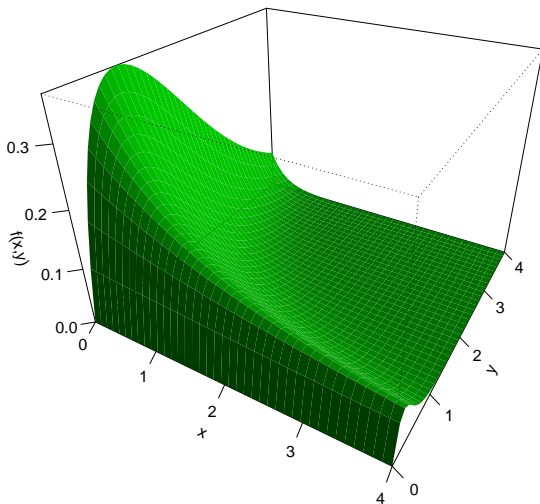
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Diagnostic  
tests

DLRs

$2 \times 2$  tables

ROC and AUC





## Lecture 5

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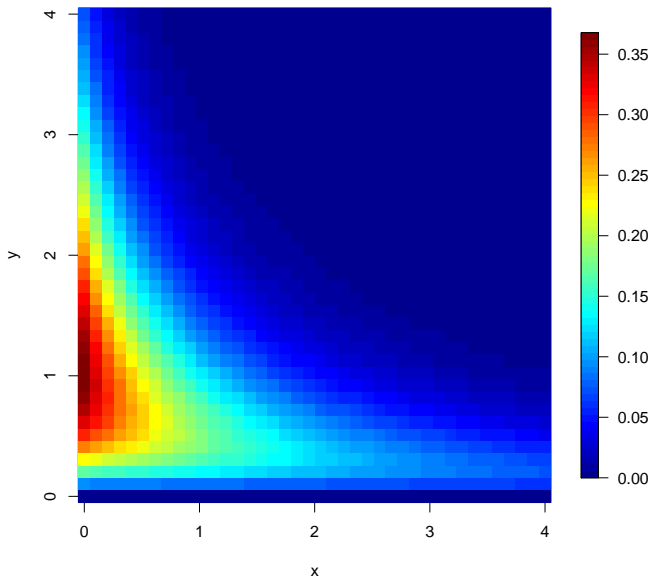
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ROC and AUC



- 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

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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

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

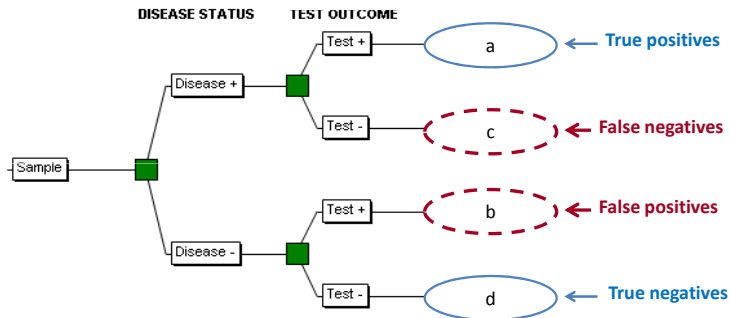
- Bayes' rule relates the conditional density of  $f(y | x)$  to the conditional density  $f(x | y)$  and the marginal density  $f(y)$
- A special case of this kind relationship is for two sets  $A$  and  $B$ , which yields that

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

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

$$(1 - sensitivity) / specificity$$

## Example

- 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  $P(D \mid +)$  given the sensitivity,  $P(+ \mid D) = .997$ , the specificity,  $P(- \mid D^c) = .985$ , and the prevalence  $P(D) = .001$

## Using Bayes' formula

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

- 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 $2 \times 2$ 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 we will 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. It's 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 $2 \times 2$ table

	Disease +	Disease -	Total
Test +	<b>a</b>	<b>b</b>	<b>a + b</b>
Test -	<b>c</b>	<b>d</b>	<b>c + d</b>
Total	<b>a + c</b>	<b>b + d</b>	<b>a + b + c + d</b>

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

The  $2 \times 2$  table: example

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

The  $2 \times 2$  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

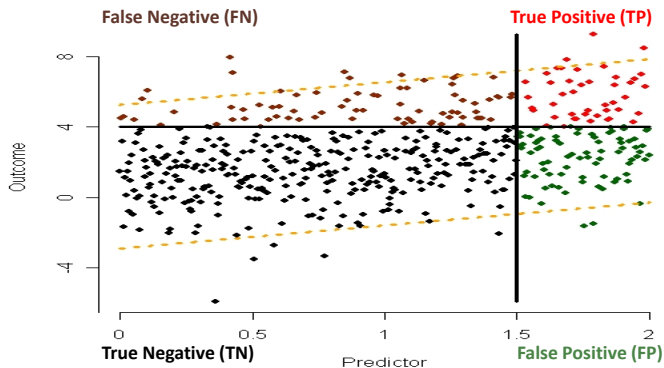
- Outcome is 0/1
- Examples
  - Non-diseased/diseased
  - Alive/Dead
  - Failure/Success (procedure)
- Continuous predictor
- Examples
  - Outcome of a diagnostic test
  - Prediction score (based on multiple characteristics)
  - Clinical score (e.g. SOFA score in ICU)

## The ROC

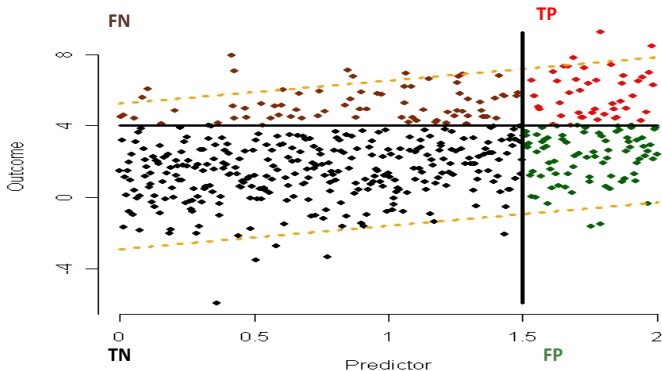
- Outcome  $D \in \{0, 1\}$ ,  $X$  scalar predictor
- For every threshold  $t$  predict  $\hat{D} = 1$  if  $X > t$
- $\text{Sens}(t) = P(X > t | D = 1)$ ,  $\text{Spec}(t) = P(X \leq t | D = 0)$
- The receiver operatic characteristic (ROC) function is

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

# Luck, error and randomness



## Dependence on the threshold

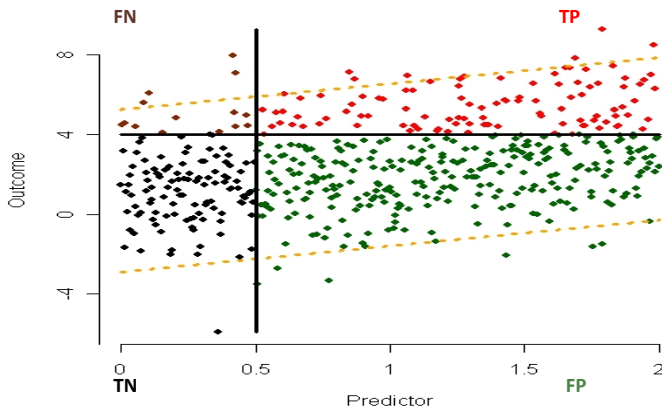


$$\text{Spec} = \frac{\text{TN}}{\text{TN} + \text{FP}} = \frac{301}{301 + 81} = 0.79$$

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



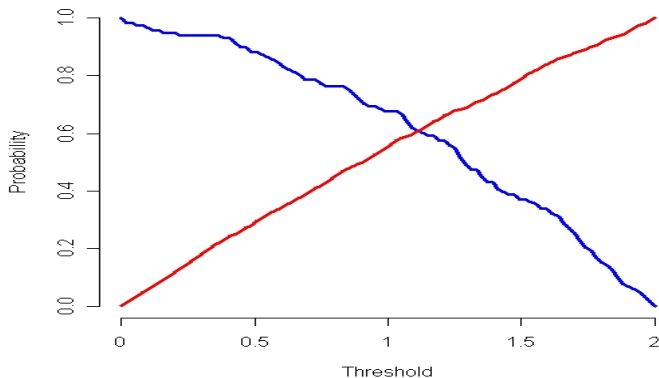
## Dependence on the threshold

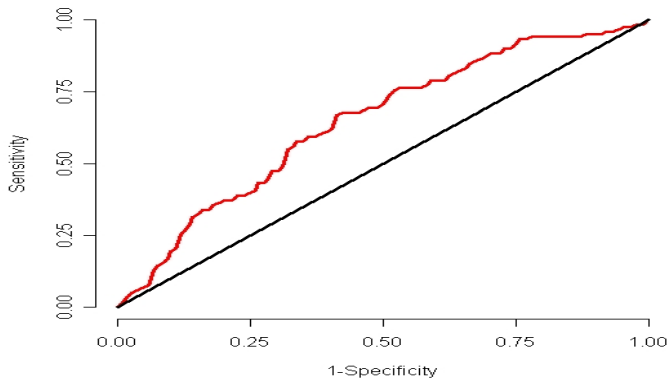


$$\text{Spec} = \frac{\text{TN}}{\text{TN} + \text{FP}} = \frac{111}{111 + 271} = 0.29$$

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

# Sensitivity and Specificity curves



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- 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

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

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

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

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

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

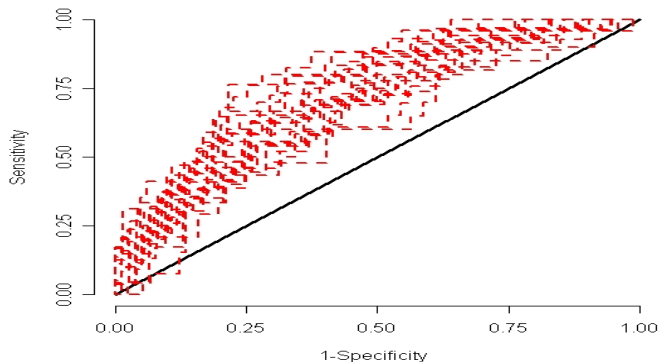
## Some comments

- 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!

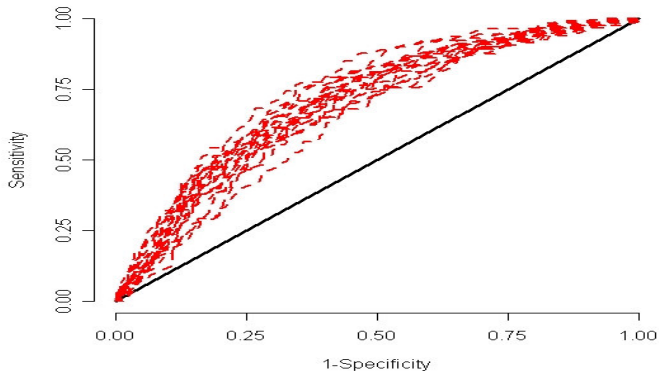
# Bootstrapping ROCs and AUCs

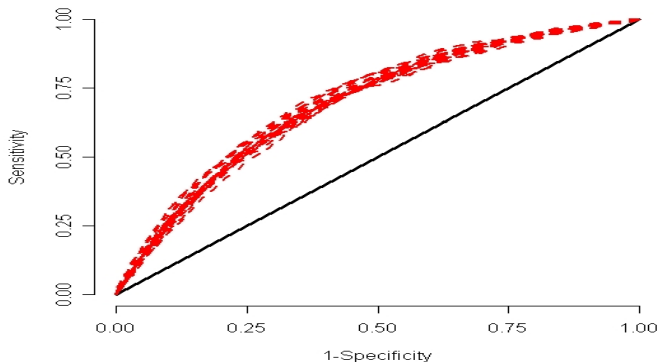
- Have a method for estimating ROC, AUC from data
- Bootstrap subjects nonparametrically (say 10,000 times)
- Repeat the estimation procedure for each data set
- Report the bootstrap distribution of ROCs and AUCs

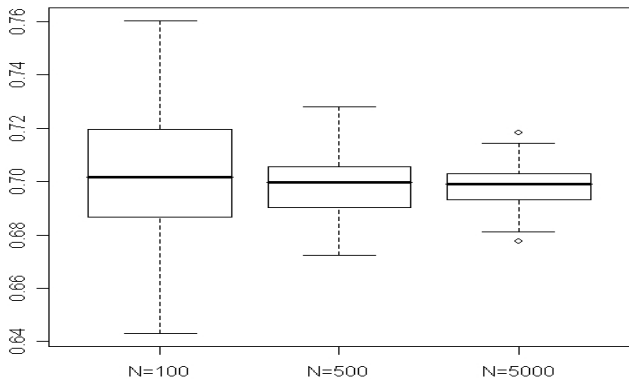
```
for (i in 1:10000)
  {boot<-sample(n,replace=TRUE)}
```

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- 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)”