

# CondProb

*Dr. B*

*Monday, November 03, 2014*

```
## First let's load a library
if (!require("xtable")){
  install.packages("xtable")
}
```

```
## Loading required package: xtable
```

```
if (!require("datasets")){
  install.packages("datasets")
}

if (!require("knitr")){
  install.packages("knitr")
}
```

```
## Loading required package: knitr
```

```
if (!require("UsingR")){
  install.packages("UsingR")
}
```

```
## Loading required package: UsingR
## Loading required package: MASS
## Loading required package: HistData
## Loading required package: Hmisc
## Loading required package: grid
## Loading required package: lattice
## Loading required package: survival
## Loading required package: splines
## Loading required package: Formula
##
## Attaching package: 'Hmisc'
##
## The following objects are masked from 'package:xtable':
##
##   label, label<-
##
## The following objects are masked from 'package:base':
##
##   format.pval, round.POSIXt, trunc.POSIXt, units
##
## Loading required package: quantreg
## Loading required package: SparseM
##
## Attaching package: 'SparseM'
```

```
##
## The following object is masked from 'package:base':
##
##      backsolve
##
##
## Attaching package: 'quantreg'
##
## The following object is masked from 'package:Hmisc':
##
##      latex
##
## The following object is masked from 'package:survival':
##
##      untangle.specials
##
##
## Attaching package: 'UsingR'
##
## The following object is masked from 'package:survival':
##
##      cancer

## set global options
opts_chunk$set(echo=FALSE, fig.height=4, fig.width=4,warning=FALSE)
```

In line text computations. The current time is Fri Nov 11/07/14 10:23:26 PM 2014 A random number is -0.446595

## Conditional Probabilty

Let + or - be the results of a diagnostic test, positive or negative, for a specific disease. Let D or D' indicate if an individual has or does not have the disease.

The SENSITIVITY is the probability that the test is positive given that the individual actually has the disease,  $P(+|D)$

The SPECIFICITY is the probability that the test is negative given that the indiviudal does not actually have the disease,  $P(-|D')$ .

The positive predictive value is the probability that the individual has the disease given that the test is positive,  $P(D|+)$ .

The negative predictive value is the probabiltly that the subject does not have the disease given that the test is negative,  $P(D'|-)$ .

The prevalence of the disease is the marginal probability of disease,  $P(D)$ .

The diagnostic likelihood ratio of a positive test, DLR+, is  $P(+|D)/P(+|D')$  which is:

$$\text{sensitivity}/(1-\text{specificity})$$

The diagnostic likelihood ratio of a negative test, DLR-, is  $P(-|D)/P(-|D')$  which is:

$(1 - \text{sensitivity}) / \text{specificity}$

A test for a certain disease has a sensitivity of 99.7% and a specificity of 98.5%

Suppose that an individual from a population with a .1% prevalence of the disease receives a positive test result. What is the positive predictive value?

First calculate the DLR+

```
## [1] 66.46667
```

The results of a positive test is that the odds of disease is now 66 times the prevalence of the disease.

```
## [1] 6.646667
```

In other words, there is a 6.65% probability that the individual has the disease given the positive test result.

```
## [1] 68.51605 68.85209
```

```
## [1] 0.4627099 0.6572901
```

```
## [1] 0.4571875 0.6591640
```

```
## attr(,"conf.level")
```

```
## [1] 0.95
```

```
## [1] 0.007 0.099
```

```
## [1] 0.01721254 0.12371005
```

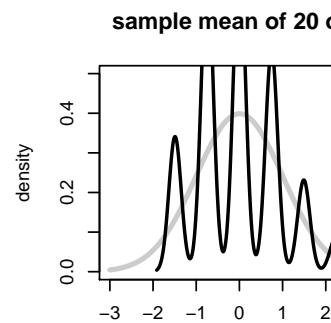
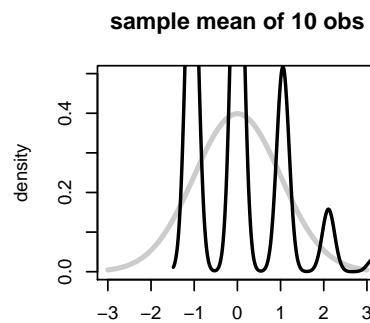
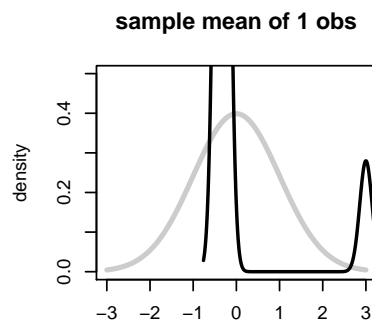
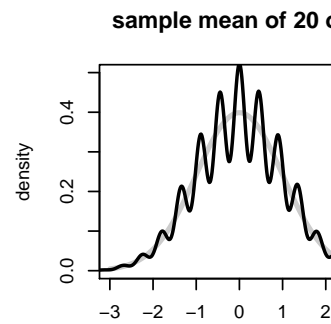
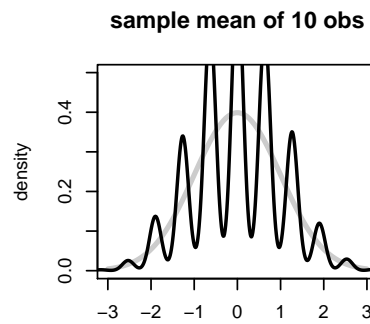
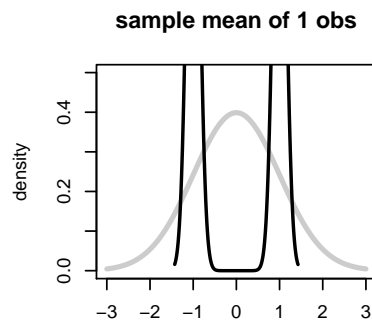
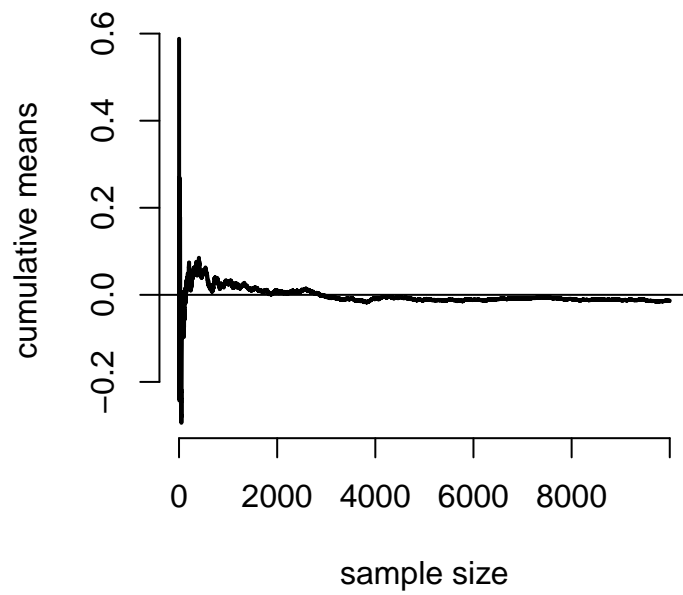
```
## attr(,"conf.level")
```

```
## [1] 0.95
```

```
## Waiting for profiling to be done...
```

```
##      2.5 %      97.5 %
```

```
## 0.01900677 0.11393446
```



Sample flip of biased coin

Estimate

Std. Error

t value
Pr(> t )
(Intercept)
-64.3421
23.0547
-2.79
0.0062
Wind
-3.3336
0.6544
-5.09
0.0000
Temp
1.6521
0.2535
6.52
0.0000
Solar.R
0.0598
0.0232
2.58
0.0112