# Package 'ExpRep'

October 12, 2022

Type Package

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**Title** Experiment Repetitions

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ExpRep-package	Experiment Repetitions

## **Description**

The package ExpRep, which basically responds to educational purposes, allows to calculate the probabilities of occurrences of an event in a great number of repetitions of Bernoulli experiment, through the application of the local and the integral theorem of De Moivre Laplace, and the theorem of Poisson. It gives the possibility to show the results graphically and analytically, and to compare the results obtained by the application of the above theorems with those calculated by the direct application of the Binomial formula.

#### **Details**

#### The DESCRIPTION file:

Package: ExpRep Type: Package

Title: Experiment Repetitions

Version: 1.0

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Author: Larisa Zamora-Matamoros and Jorge Diaz-Silvera Maintainer: Larisa Zamora-Matamoros <larisa@uo.edu.cu>

Description: Allows to calculate the probabilities of occurrences of an event in a great number of repetitions of Bernoulli ex

License: Unlimited

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DeMoivre-Laplace.

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S\_Integral\_Theorem Simulations of the Integral Theorem of

DeMoivre-Laplace.

S\_Local\_Limit\_Theorem Simulations of Local Theorem of

DeMoivre-Laplace

S\_Poisson\_Theorem Simulations of Poisson Theorem

#### Author(s)

Larisa Zamora-Matamoros and Jorge Diaz-Silvera

Maintainer: Larisa Zamora-Matamoros <larisa@uo.edu.cu>

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### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

## **Examples**

```
ProbL<-Local_Theorem(n=100,m=50,p=0.02)
ProbL

ProbI<-Integral_Theorem(n=100,p=0.5,linf=0,lsup=50)
ProbI

ProbP<-Poisson_Theorem(n=100,m=50,p=0.002)
ProbP

beta<-ApplicIntegralTheo(Applic="beta",n=369,p=0.4,alpha=0.05)
beta

alpha<-ApplicIntegralTheo(Applic="alpha",n=369,p=0.4,beta=0.95)
alpha

n<-ApplicIntegralTheo(Applic="n",p=0.4,alpha=0.05,beta=0.95)
n

S_Local_Limit_Theorem(n = 170, p = 0.5, Compare = TRUE, Table = TRUE, Graph = TRUE, GraphE = TRUE)

S_Poisson_Theorem(n = 169, p = 0.002, Compare = TRUE, Table = TRUE, Graph = TRUE, GraphE = TRUE)

S_Integral_Theorem(n=100, p=0.5, linf = 0, lsup = 50, Compare = TRUE, Table = TRUE, Graph = TRUE, GraphE = TRUE)</pre>
```

ApplicIntegralTheo

Applications of the Integral Theorem of DeMoivre-Laplace.

## **Description**

This function shows three applications of the integral theorem of DeMoivre-Laplace: 1. To estimate the probability (beta) that the frequency of occurrence of the successful event will deviate from the probability that this event will happen in any single Bernoulli experiment (p) in a quantity not bigger than alpha. 2. To calculate the least number of experiments that must be carried out (n). 3. To determine the boundary of possible variations between the frequency of occurrence of the successful event and the probability p (alpha).

# Usage

```
ApplicIntegralTheo(Applic = "alpha", n = 10000, p = 0.5, alpha = 0.01, beta = 0.9)
```

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## **Arguments**

Applic	It indicates the calculation to be carried out: if "n", the function calculates the number of repetitions, if "alpha", the function calculates the boundary of possible variations of abs(frequency-p) if "beta", the function calculates the probability that the frequency of occurrence of the successful event will deviate from the probability p by no more than alpha.
n	Number of repetitions of the Bernoulli trial.
р	Probability that a successful event happens in any single Bernoulli trial (called the probability of success).
alpha	The boundary of possible variations of abs(frequency-p).
beta	Probability that the frequency of occurrence of the successful event will deviate from the probability p by no more than alpha.

## Value

Numeric value representing the values of n, alpha or beta according to the value that the parameter "Applic" takes ("n", "alpha" or "beta").

### Note

Department of Mathematics. University of Oriente. Cuba.

## Author(s)

Larisa Zamora and Jorge Diaz

## References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

# See Also

Integral\_Theorem.

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```
Alpha <- function(n, p, beta) {
      a <- (beta + 1)/2
      alpha <- ((p * (1 - p)/n)^0.5) * qnorm(a)
      return(alpha)
  Beta <- function(n, p, alpha) {</pre>
      b \leftarrow alpha * (n/(p * (1 - p)))^0.5
      beta <- 2 * pnorm(b) - 1
      return(beta)
  }
  Repetitions <- function(p, alpha, beta) {</pre>
      a <- (beta + 1)/2
      n \leftarrow (p * (1 - p) * ((qnorm(a)/alpha)^2))%/%1 + 1
      return(n)
  options(digits = 17)
  value <- switch(Applic, alpha = Alpha(n, p, beta), beta = Beta(n,</pre>
      p, alpha), n = Repetitions(p, alpha, beta))
  return(value)
}
```

Buffon

Buffon

# **Description**

Simulations of the experiment of Buffon.

## Usage

```
Buffon(p = 0.5, width = 0.2, r = c(100, 500, 1000, 1500))
```

# Arguments

p Probability of occurrence of some event.

width Width of the band where the probabilities are represented.

r Array of four values, representing the numbers of repetitions of the experiment that will be carried out.

### Value

Four graphics, each one is the simulation of the experiment of Buffon for the number of repetitions contained in the array r.

## Note

Department of Mathematics. University of Oriente. Cuba.

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### Author(s)

Larisa Zamora and Jorge Diaz

#### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

```
Buffon(p = 0.5, width = 0.2, r = c(100, 500, 1000, 1500))
## The function is currently defined as
function (p = 0.5, width = 0.2, r = c(100, 500, 1000, 1500))
    Position <- function(k, colum) {
        PE <- k%/%colum
        Resto <- k%%colum
        if (Resto == 0) {
            fila <- PE
            columna <- colum
        }
        else {
            fila <- PE + 1
            columna <- Resto
        Position <- list(fila, columna)
        return(Position)
    }
    nf \leftarrow layout(matrix(c(1, 2, 3, 4), 2, 2, byrow = TRUE), TRUE)
    k <- 0
    la <- p - width
    lb <- p + width</pre>
    if (la < 0)
        la <- 0
    if (lb > 1)
        lb <- 1
    for (j in 1:4) {
        k < -k + 1
        Probcara \leftarrow array(0, dim = r[j])
        for (i in 1:r[j]) {
            binomial <- rbinom(i, 1, p)</pre>
            cara <- length(binomial[binomial == 1])</pre>
            Probcara[i] <- cara/i
        }
        P <- Position(k, 2)
        fila <- P[[1]]
        colum <- P[[2]]</pre>
        mfg <- c(fila, colum, 2, 2)
        a <- as.character(r[j])</pre>
        plot(Probcara, type = "p", main = paste0("n=", a), xlab = "Repetitions",
            ylab = "Probability", font.main = 3, col = "blue",
            ylim = c(la, lb))
```

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```
abline(h = p, col = "red", lty = 1, lwd = 2)
}
```

Integral\_Theorem

Integral Theorem of DeMoivre-Laplace

# Description

Given n Bernoulli experiments, with success probability p, this function calculates the probability that a successful event occurs between linf and lsup times.

### Usage

```
Integral_Theorem(n = 100, p = 0.5, linf = 0, lsup = 50)
```

# Arguments

n	An integer value representing the number of repetitions of the Bernoulli experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success).
linf	An integer value representing the minimum number of times that the successful event should happen.
lsup	An integer value representing the maximum number of times that the successful event should happen.

## **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. The value of n must be high and the value of p must be small. It is necessary that linf < lsup.

## Value

A real value representing the approximate probability that a successful event occurs between linf and lsup times, in n repetitions of a Bernoulli experiment.

## Note

Department of Mathematics. University of Oriente. Cuba.

# Author(s)

Larisa Zamora and Jorge Diaz

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### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

#### See Also

Poisson\_Theorem, Local\_Theorem.

## **Examples**

```
Prob<-Integral_Theorem(n=100,p=0.5,linf=0,lsup=50)
Prob

## The function is currently defined as
function (n = 100, p = 0.5, linf = 0, lsup = 50)
{
          A <- (linf - n * p)/sqrt(n * p * (1 - p))
          B <- (lsup - n * p)/sqrt(n * p * (1 - p))
          P <- pnorm(B) - pnorm(A)
          return(P)
     }</pre>
```

Local\_Theorem

Local Theorem of DeMoivre-Laplace

## **Description**

Given n Bernoulli experiments, with success probability p, this function calculates the approximate probability that a successful event occurs exactly m times.

## Usage

```
Local_Theorem(n, m, p)
```

# Arguments

n	An integer value representing the number of repetitions of the Bernoulli experiment.
m	An integer value representing the number of times that a successful event occurs in the n repetitions of the Bernoulli experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success).

## **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. The value of n must be high and the value of p must be small.

## Value

A real value representing the approximate probability that a successful event occurs exactly m times in n repetitions of a Bernoulli experiment.

## Note

Department of Mathematics. University of Oriente. Cuba.

### Author(s)

Larisa Zamora and Jorge Diaz

#### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

#### See Also

Integral\_Theorem, Poisson\_Theorem.

# **Examples**

```
Prob<-Local_Theorem(n=100,m=50,p=0.02)
Prob

## The function is currently defined as function (n, m, p)
{
    a <- n * p
    b <- sqrt(a * (1 - p))
    x <- (m - a)/b
    P <- dnorm(x, 0, 1)/b
    return(P)
}</pre>
```

Poisson\_Theorem

Poisson Theorem.

## **Description**

Given n Bernoulli experiments, with success probability p (p small), this function calculates the approximate probability that a successful event occurs exactly m times.

# Usage

```
Poisson_Theorem(n, m, p)
```

### **Arguments**

n	An integer value representing the number of repetitions of the Bernoulli experiment.
m	An integer value representing the number of times that a successful event occurs in the n repetitions of the Bernoulli experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success).

## **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. The value of n must be high and the value of p must be very small.

### Value

A numerical value representing the approximate probability that a successful event occurs exactly m times.

### Note

Department of Mathematics. University of Oriente. Cuba.

#### Author(s)

Larisa Zamora and Jorge Diaz

### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

#### See Also

Integral\_Theorem, Local\_Theorem.

```
Prob<-Poisson_Theorem(n=100,m=50,p=0.002)
Prob

## The function is currently defined as function (n, m, p)
{
    landa <- n * p
    P <- dpois(m, landa)
    return(P)
}</pre>
```

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S_Integral_Theorem Simulations of the Integral Theorem of DeMoivre-Laplace.
---

## **Description**

Given n Bernoulli experiments with success probability p, this function calculates and plots the exact probability and the approximate probability that a successful event occurs between linf+i (0<=i<=lsup-linf-1) and lsup times. It also calculates the difference between these probabilities and shows all the computations in a table.

# Usage

```
S_{Integral_Theorem}(n = 200, p = 0.5, linf = 0, lsup = 100, Compare = TRUE, Table = TRUE, Graph = TRUE, GraphE = TRUE)
```

# **Arguments**

n	An integer vector with the numbers of repetitions of the Bernoulli experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success).
linf	An integer vector, of the same length than n, with the minimum numbers of times that the successful event should happen.
lsup	An integer vector, of the same length than n, with the maximum numbers of times that the successful event should happen.
Compare	A logical value, if TRUE the function calculates both the exact probability and the approximate probability that a successful event occurs and compares these probabilities.
Table	A logical value, if TRUE the function shows a table with the carried out computations.
Graph	A logical value, if TRUE the function plots both the exact probability and the approximate probability that a successful event occurs.
GraphE	A logical value, if TRUE the function shows the graphic of the errors in the approximation.

## **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. It is necessary that linf < lsup.

#### Value

A graph and/or a table.

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### Note

Department of Mathematics. University of Oriente. Cuba.

#### Author(s)

Larisa Zamora and Jorge Diaz

#### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

#### See Also

Integral\_Theorem, Local\_Theorem.

```
S_{Integral_{D}} = 100, p = 0.5, linf = 0, lsup = 100, lsup = 
          Graph = TRUE, GraphE = TRUE)
## The function is currently defined as
function (n = 200, p = 0.5, linf = 0, lsup = 100, Compare = TRUE,
          Table = TRUE, Graph = TRUE, GraphE = TRUE)
    { Integral_Theorem <- function(n = 100, p = 0.5, linf = 0,
                     lsup = 50) {
                     A \leftarrow (linf - n * p)/sqrt(n * p * (1 - p))
                    B \leftarrow (lsup - n * p)/sqrt(n * p * (1 - p))
                     P <- pnorm(B) - pnorm(A)
                     return(P)
         layout(matrix(1))
         PNormal <- numeric()
         Dif <- numeric()</pre>
         PBin <- numeric()
          k \leftarrow lsup - linf - 1
         PNormal[1] <- Integral_Theorem(n, p, linf, lsup)</pre>
         PBin[1] <- 0
          for (j in linf:lsup) PBin[1] <- PBin[1] + dbinom(j, n, p)</pre>
         Dif[1] <- abs(PBin[1] - PNormal[1])</pre>
          for (i in 1:k) {
                     linf_i <- linf + i</pre>
                     PNormal[i + 1] <- Integral_Theorem(n, p, linf_i, lsup)</pre>
                     if (Compare == TRUE) {
                                PBin[i + 1] <- 0
                                for (j in linf_i:lsup) PBin[i + 1] \leftarrow PBin[i + 1] +
                                           dbinom(j, n, p)
                                Dif[i + 1] \leftarrow abs(PBin[i + 1] - PNormal[i + 1])
                     }
          if (Graph == TRUE & GraphE == TRUE) {
                     layout(matrix(c(1, 1, 2, 2), 2, 2, byrow = TRUE))
          }
```

```
if (Graph == TRUE) {
      ymini \leftarrow min(PNormal[k + 1], PBin[k + 1]) - 0.05
      ymaxi <- max(PNormal[1], PBin[1]) + 0.05</pre>
      mfg <- c(1, 1, 2, 2)
   plot(PNormal, ylim = c(ymini, ymaxi), type = "1", main = "The Integral Limit Theorem",
          xlab = "k (linf<=k<=lsup)", ylab = "Probability",</pre>
          col = "red")
      mtext("Integral Theorem", line = -1, side = 1, adj = 1,
          col = "red")
      if (Compare == TRUE) {
          points(PBin, type = "p", col = "blue")
          mtext("Binomial Probability", line = -2, side = 1,
               adj = 1, col = "blue")
      }
  if (GraphE == TRUE) {
      mfg <- c(2, 1, 2, 2)
      dmini <- min(Dif) - 0.01</pre>
      dmaxi \leftarrow max(Dif) + 0.01
      plot(Dif, ylim = c(dmini, dmaxi), type = "b", main = "Errors",
          xlab = "m", ylab = "Errors", col = "green")
      abline(a = 0, b = 0, col = "red")
  }
  if (Table == TRUE) {
      Ak <- array(1:(k + 1))
      if (Compare == TRUE)
          TablaR <- data.frame(k = Ak, PBinomial = PBin, T_Integral = PNormal,
              Difference = Dif)
      else TablaR <- data.frame(K = Ak, T_Integral = PNormal)</pre>
      TablaR
 }
}
```

S\_Local\_Limit\_Theorem Simulations of Local Theorem of DeMoivre-Laplace

#### **Description**

Given n Bernoulli experiments, with success probability p, this function calculates and plots the exact probability and the approximate probability that a successful event occurs exactly m times (0<=m<=n). It also calculates the difference between these probabilities and shows all the computations in a table.

# Usage

### **Arguments**

n	An integer value representing the number of repetitions of the experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success).
Compare	A logical value, if TRUE the function calculates both the exact probability and the approximate probability that a successful event occurs exactly m times and compares these probabilities.
Table	A logical value, if TRUE the function shows a table with the carried out computations.
Graph	A logical value, if TRUE the function plots both the exact probability and the approximate probability that a successful event occurs exactly m times.
GraphE	A logical value, if TRUE the function shows the graphic of the errors in the approximation.

### **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. The value of n must be high and the value of p must be small.

### Value

A graph and/or a table.

#### Note

Department of Mathematics. University of Oriente. Cuba.

# Author(s)

Larisa Zamora and Jorge Diaz

## References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

### See Also

Integral\_Theorem, Local\_Theorem.

```
m <- array(0:n)</pre>
  x <- numeric()</pre>
 PNormal <- numeric()
  a <- n * p
 b <- sqrt(a * (1 - p))
  for (mi in 1:(n + 1)) {
      x[mi] <- (mi - 1 - a)/b
      PNormal[mi] <- dnorm(x[mi], 0, 1)/b
  if (Compare == TRUE) {
      PBin <- numeric()</pre>
      for (mi in 1:(n + 1)) PBin[mi] \leftarrow dbinom(mi - 1, n, p)
      Dif <- abs(PBin - PNormal)</pre>
  if (Graph == TRUE & GraphE == TRUE) {
      layout(matrix(c(1, 1, 2, 2), 2, 2, byrow = TRUE))
  if (Graph == TRUE) {
      mfg <- c(1, 1, 2, 2)
      plot(PNormal, type = "p", main = "The Local Limit Theorem",
          xlab = "m", ylab = "Probability", col = "red")
      mtext("Local Theorem", line = -1, side = 3, adj = 1,
          col = "red")
      if (Compare == TRUE) {
          points(PBin, type = "p", col = "blue")
          mtext("Binomial Probability", line = -2, side = 3,
              adj = 1, col = "blue")
      }
  if (GraphE == TRUE) {
      mfg <- c(2, 1, 2, 2)
      dmini <- min(Dif) - 0.01
      dmaxi \leftarrow max(Dif) + 0.01
      plot(Dif, ylim = c(dmini, dmaxi), type = "b", main = "Errors",
          xlab = "m", ylab = "Errors", col = "green")
      abline(a = 0, b = 0, col = "red")
  if (Table == TRUE) {
      if (Compare == TRUE)
          TablaR \leftarrow data.frame(m = m, x = x, PBinomial = PBin,
              TLocal = PNormal, Difference = Dif)
      else TablaR <- data.frame(m = m, x = x, TLocal = PNormal)
      TablaR
 }
}
```

### **Description**

Given n Bernoulli experiments, with success probability p, this function calculates and plots the exact probability and the approximate probability that a successful event occurs exactly m times (0<=m<=n). It also calculates the difference between theses probabilities and shows all the computations in a table.

# Usage

## **Arguments**

n	An integer value representing the number of repetitions of the Bernoulli experiment.
p	A real value with the probability that a successful event will happen in any single Bernoulli experiment (called the probability of success)
Compare	A logical value, if TRUE the function calculates both the exact probability and the approximate probability that a successful event occurs exactly m times and compares these probabilities.
Table	A logical value, if TRUE the function shows a table with the carried out computations.
Graph	A logical value, if TRUE the function plots both the exact probability and the approximate probability that a successful event occurs exactly m times.
GraphE	A logical value, if TRUE, the function shows the graphics corresponding to the differences between the probabilities using the Binomial approach and Poisson theorem, and the probabilities using the Binomial approach and Local theorem.

### **Details**

Bernoulli experiments are sequences of events, in which successive experiments are independent and at each experiment the probability of appearance of a "successful" event (p) remains constant. The value of n must be high and the value of p must be very small.

#### Value

A graph and/or a table.

#### Note

Department of Mathematics. University of Oriente. Cuba.

## Author(s)

Larisa Zamora and Jorge Diaz

### References

Gnedenko, B. V. (1978). The Theory of Probability. Mir Publishers. Moscow.

#### See Also

Integral\_Theorem, Local\_Theorem.

```
S_Poisson_Theorem(n = 169, p = 0.002, Compare = TRUE, Table = TRUE, Graph = TRUE,
    GraphE = TRUE
## The function is currently defined as
function (n = 2000, p = 0.002, Compare = TRUE, Table = TRUE,
    Graph = TRUE, GraphE = FALSE)
  { layout(matrix(1))
    m \leftarrow array(0:n)
    PPoisson <- numeric()</pre>
    a <- n * p
    for (mi in 1:(n + 1)) PPoisson[mi] <- dpois(mi - 1, a)
    if (Compare == TRUE) {
        PBin <- numeric()</pre>
        x <- numeric()</pre>
        PNormal <- numeric()
        Dif1 <- numeric()</pre>
        Dif2 <- numeric()</pre>
        b <- sqrt(a * (1 - p))
        for (mi in 1:(n + 1)) {
            x[mi] <- (mi - 1 - a)/b
            PBin[mi] <- dbinom(mi - 1, n, p)</pre>
            PNormal[mi] <- dnorm(x[mi], 0, 1)/b
        Dif1 <- abs(PBin - PPoisson)</pre>
        Dif2 <- abs(PBin - PNormal)</pre>
    if (Graph == TRUE & GraphE == TRUE) {
        layout(matrix(c(1, 1, 2, 2), 2, 2, byrow = TRUE))
    if (Graph == TRUE) {
        mfg <- c(1, 1, 2, 2)
        11 <- length(which(Dif1 > 5e-07))
        plot(PPoisson[1:11], type = "b", main = "The Poisson Theorem",
            xlab = "m", ylab = "Probability", col = "red")
        mtext("Poisson Theorem", line = -1, side = 3, adj = 1,
            col = "red")
        if (Compare == TRUE) {
            points(PBin[1:11], type = "b", col = "green")
            points(PNormal[1:11], type = "b", col = "blue")
            mtext("Local Theorem", line = -2, side = 3, adj = 1,
                 col = "blue")
            mtext("Binomial Probability", line = -3, side = 3,
                 adj = 1, col = "green")
        }
```

```
if (GraphE == TRUE) {
     mfg <- c(2, 1, 2, 2)
     11 <- length(which(Dif1 > 5e-07))
     plot(Dif2[1:11], type = "b", main = "Errors", xlab = "m",
          ylab = "Differences", col = "red")
      mtext("Binomial-Poisson", line = -1, side = 3, adj = 1,
          col = "red")
     points(Dif1[1:11], type = "b", col = "green")
     mtext("Binomial-Local Theorem", line = -2, side = 3,
          adj = 1, col = "green")
  if (Table == TRUE) {
      if (Compare == TRUE)
          TablaR <- data.frame(m = m, x = x, PBinomial = PBin,
              TPoisson = PPoisson, Difference1 = Dif1, TLocal = PNormal,
              Difference2 = Dif2)
      else TablaR <- data.frame(m = m, TPoisson = PPoisson)</pre>
      TablaR
 }
}
```

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