

# Package ‘BonEV’

October 12, 2022

**Type** Package

**Title** An Improved Multiple Testing Procedure for Controlling False  
Discovery Rates

**Version** 1.0

**Date** 2016-02-10

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**Depends** R (>= 3.2.0), qvalue

**Description** An improved multiple testing procedure for controlling false discovery rates which is developed based on the Bonferroni procedure with integrated estimates from the Benjamini-Hochberg procedure and the Storey's q-value procedure. It controls false discovery rates through controlling the expected number of false discoveries.

**License** GPL (>= 2)

**NeedsCompilation** no

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## Description

BonEV is an improved multiple testing procedure for controlling false discovery rates which is developed based on the Bonferroni procedure with integrated estimates from the Benjamini-Hochberg procedure and the Storey's q-value procedure. It controls false discovery rates through controlling the expected number of false discoveries.

## Details

Package: BonEV  
Type: Package  
Version: 1.0.0  
Date: 2015-02-10  
Depends: R (>= 3.2.0), qvalue  
License: GPL (>= 2)

## Author(s)

Dongmei Li Maintainer: Dongmei Li <dongmei\_li@urmc.rochester.edu>

## See Also

The [Bon\\_EV](#) function defined in this package. The [qvalue](#) package.

## Examples

```
library(qvalue)
data(hedenfalk)
summary(hedenfalk)
pvalues <- hedenfalk$p
adjp <- Bon_EV(pvalues, 0.05)
summary(adjp)
results <- cbind(adjp$raw_P_value, adjp$BH_adjp, adjp$Storey_adjp, adjp$Bon_EV_adjp)
results

##Compare with Benjamini-Hochberg and Storey's q-value procedures
sum(adjp$raw_P_value <= 0.05)
sum(adjp$BH_adjp <= 0.05)
sum(adjp$Storey_adjp <= 0.05)
sum(adjp$Bon_EV_adjp <= 0.05)
```

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Bon\_EV

*Bon\_EV: A R Function of Improved Multiple Testing Procedure for Controlling False Discovery Rates*

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## Description

Bon\_EV is an improved multiple testing procedure for controlling false discovery rates which is developed based on the Bonferroni procedure with integrated estimates from the Benjamini-Hochberg procedure and the Storey's q-value procedure. It controls false discovery rates through controlling the expected number of false discoveries.

## Usage

```
Bon_EV(pvalue, alpha)
```

## Arguments

pvalue	The input data is a vector of P-values ranged from 0 to 1
alpha	The alpha is the level of false discovery rates (FDR) to control for

## Details

Bon\_EV is a function for getting adjusted P-values with FDR controlled at level alpha.

## Value

Bon\_EV produces a named list with the following components:

raw_P_value	Vector of raw P-values
BH_adjp	Adjusted P-values from the Benjamini-Hochberg procedure
Storey_adjp	Adjusted P-values from the Storey's q-value procedure
Bon_EV_adjp	Adjusted P-values from the Bon-EV multiple testing procedure

## Author(s)

Dongmei Li

## See Also

The qvalue package.

**Examples**

```
library(qvalue)
data(hedenfalk)
summary(hedenfalk)
pvalues <- hedenfalk$p
adjp <- Bon_EV(pvalues, 0.05)
summary(adjp)
sum(adjp$Bon_EV_adjp <= 0.05)
```

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