

# Package ‘MIWilson’

August 23, 2021

**Title** Implementing the MI-Wilson Confidence Interval

**Version** 1.0.0

**Description** Implements the Wilson confidence interval for binomial proportions given multiple imputations of missing data (detailed theory provided in “Wilson Confidence Intervals for Binomial Proportions With Multiple Imputation for Missing Data” (A. Lott & J. Reiter, 2018)). Our package also implements a Wald confidence interval and allows for both MIDs object and proportion vector arguments.

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**Encoding** UTF-8

**RoxygenNote** 7.1.1

**URL** <https://github.com/hungf8342/MIWilson>

**BugReports** <https://github.com/hungf8342/MIWilson/issues>

**Imports** dplyr, magrittr, mice, stats

**Suggests** rmarkdown, knitr, testthat (>= 3.0.0)

**Config/testthat.edition** 3

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Repository** CRAN

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**Index****11****Bm***Calculate between-imputation variance of the response mean*

$$\frac{\sum(\hat{Q}_l - \bar{Q})}{m - 1}$$

**Description**

Calculate between-imputation variance of the response mean

$$\frac{\sum(\hat{Q}_l - \bar{Q})}{m - 1}$$

**Usage**

```
Bm(qhats, m)
```

**Arguments**

- |       |   |
|-------|---|
| qhats | vector of Qhats(means of response for each imputed dataset) |
| m     | number of imputed datasets                                  |

**Value**

Bm: the between-dataset variance of the response mean

**Examples**

```
imp = mice::mice(mice::nhanes)
qhats = Qhats(imp, "hyp")
m = imp$m
Bm(qhats, m)
```

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dof	<i>Calculate degrees of freedom used in calculating confidence intervals of t-distributed proportion point estimate <math>\bar{Q}_m</math></i>
-----	--

---

**Description**

Calculate degrees of freedom used in calculating confidence intervals of t-distributed proportion point estimate  $\bar{Q}_m$

**Usage**

```
dof(qhats, m, nrow)
```

**Arguments**

qhats	vector of Qhats(means of response for each imputed dataset)
m	number of imputed datasets
nrow	number of observations in the imputed dataset

**Value**

degrees of freedom

**Examples**

```
imp = mice::mice(mice::nhanes)
qhats = Qhats(imp, "hyp")
m = imp$m
nrow = imp$data %>% nrow()
dof(qhats, m, nrow)
```

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mi_wald	<i>Calculates the specified Wald CI of a binomial proportion variable, given imputed data sets.</i>
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---

**Description**

Calculates the specified Wald CI of a binomial proportion variable, given imputed data sets.

**Usage**

```
mi_wald(mids_obj, response, ci_level = 0.95, summaries = TRUE)
```

**Arguments**

<code>mids_obj</code>	mids object created by mice package
<code>response</code>	string name of response variable (must be 0-1 valued)
<code>ci_level</code>	desired confidence interval level (defaults to 95%)
<code>summaries</code>	boolean: should summary helper values be printed (default TRUE)

**Value**

two-length vector of Wald lower CI and upper CI

**Examples**

```
imp = mice::mice(mice::nhanes %>% dplyr::mutate(hyp = hyp-1))
mi_wald(imp, "hyp", 0.95)
```

**mi\_wald\_phat**

*Calculates the MI-Wald interval if given a vector of observed binomial proportions (one for each imputed data frame)*

**Description**

Calculates the MI-Wald interval if given a vector of observed binomial proportions (one for each imputed data frame)

**Usage**

```
mi_wald_phat(phats, n, ci_level = 0.95, summaries = TRUE)
```

**Arguments**

<code>phats</code>	vector of binomial proportions (one for each imputation)
<code>n</code>	the common number of observations over the imputed dataframes
<code>ci_level</code>	desired confidence interval level (default 95%)
<code>summaries</code>	boolean: should summary helper values be printed (default TRUE)

**Value**

Two-length vector of Wilson lower CI and upper CI

**Related Functions**

- [mi\\_wald](#)
- [mi\\_wilson\\_phat](#)

**Examples**

```
phats = c(0.2, 0.23, 0.25)
mi_wald_phat(phats, 100, 0.99, TRUE)
```

**mi\_wilson**

*Calculates the specified Wilson CI of a binomial proportion variable, given imputed data sets.*

**Description**

Calculates the specified Wilson CI of a binomial proportion variable, given imputed data sets.

**Usage**

```
mi_wilson(mids_obj, response, ci_level = 0.95, summaries = TRUE)
```

**Arguments**

mids_obj	mids object created by mice package
response	string name of response variable (must be 0-1 valued)
ci_level	desired confidence interval level (defaults to 95%)
summaries	boolean: should summary helper values be printed (default TRUE)

**Value**

two-length vector of Wilson lower CI and upper CI

**Examples**

```
imp = mice::mice(mice:::nhanes %>% dplyr::mutate(hyp = hyp-1))
mi_wilson(imp, "hyp", 0.95)
```

**mi\_wilson\_phat**

*Calculates the MI-Wilson interval if given a vector of observed binomial proportions (one for each imputed data frame)*

**Description**

Calculates the MI-Wilson interval if given a vector of observed binomial proportions (one for each imputed data frame)

**Usage**

```
mi_wilson_phat(phats, n, ci_level = 0.95, summaries = TRUE)
```

**Arguments**

<code>phats</code>	vector of binomial proportions (one for each imputation)
<code>n</code>	the common number of observations over the imputed dataframes
<code>ci_level</code>	desired confidence interval level (default 95%)
<code>summaries</code>	boolean: should summary helper values be printed (default TRUE)

**Value**

two-length vector of Wilson lower CI and upper CI

**Examples**

```
phats = c(0.2, 0.23, 0.25)
mi_wilson_phat(phats, 100, 0.99, TRUE)
```

Qbar

*Calculate Qbar (average response over MICE datasets)*

**Description**

Calculate Qbar (average response over MICE datasets)

**Usage**

```
Qbar(qhats)
```

**Arguments**

<code>qhats</code>	vector of Qhats(response means for each imputed dataset)
--------------------	--

**Value**

Qbar: the average response over MICEd datasets.

**Examples**

```
imp = mice::mice(mice::nhanes)
qhats = Qhats(imp, "hyp")
Qbar(qhats)
```

Qhats

*Calculate Qhats (means of response for each imputed dataset)***Description**

Calculate Qhats (means of response for each imputed dataset)

**Usage**

```
Qhats(mids_obj, response)
```

**Arguments**

mids_obj	mids object created by mice package
response	string name of binary response variable

**Value**

Qhats: vector of response means for each imputed dataset

**Examples**

```
imp = mice::mice(mice::nhanes)
Qhats(imp, "hyp")
```

Rm

*Helper function for getting rm, a key component for calculating degrees of freedom and the wilson CI directly***Description**

Helper function for getting rm, a key component for calculating degrees of freedom and the wilson CI directly

**Usage**

```
Rm(qhats, m, nrow)
```

**Arguments**

qhats	vector of Qhats(means of response for each imputed dataset)
m	number of imputed datasets
nrow	number of observations in the imputed dataset

**Value**

```
rm
```

**Examples**

```
imp = mice::mice(mice:::nhanes)
qhats = Qhats(imp, "hyp")
m = imp$m
nrow = imp$data %>% nrow()
Tm(qhats, m, nrow)
```

**Tm***Estimate variance of proportion point estimate  $\bar{Q}_m$* **Description**

Estimate variance of proportion point estimate  $\bar{Q}_m$

**Usage**

```
Tm(qhats, m, nrow)
```

**Arguments**

<code>qhats</code>	vector of <code>Qhats</code> (means of response for each imputed dataset)
<code>m</code>	number of imputed datasets
<code>nrow</code>	number of observations in the imputed dataset

**Value**

variance of proportion point estimate

**Examples**

```
imp = mice::mice(mice:::nhanes)
qhats = Qhats(imp, "hyp")
m = imp$m
nrow = imp$data %>% nrow()
Tm(qhats, m, nrow)
```

---

**Ubar***Calculate Ubar (average response variance over MICE datasets)*

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**Description**

Calculate Ubar (average response variance over MICE datasets)

**Usage**

```
Ubar(qhats, m, nrow)
```

**Arguments**

qhats	vector of Qhats(means of response for each imputed dataset)
m	number of imputed datasets
nrow	number of observations in the imputed dataset

**Value**

Ubar: average response variance over MICE datasets

**Examples**

```
imp = mice::mice(mice::nhanes)
qhats = Qhats(imp, "hyp")
m = imp$m
nrow = imp$data %>% nrow()
Ubar(qhats, m, nrow)
```

---

**Uhats***Calculate Uhats (variance for each imputed dataset)*

---

**Description**

Calculate Uhats (variance for each imputed dataset)

**Usage**

```
Uhats(qhats, nrow)
```

**Arguments**

qhats	vector of Qhats(means of response for each imputed dataset)
nrow	number of observations in the imputed dataset

**Value**

*Uhats*: vector of response variances for each imputed dataset

**Examples**

```
imp = mice::mice(mice::nhanes)
qhats = Qhats(imp, "hyp")
nrow = imp$data %>% nrow()
Uhats(qhats, nrow)
```

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