

# Package ‘ewhorm’

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**Title** Statistical Considerations for Designing e-WHORM Adaptive Trial

**Version** 0.1

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**Description** Data simulation and analysis for the design of e-WHORM trial

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**Encoding** UTF-8

**Imports** mvtnorm,  
stats,  
multcomp,  
gtools

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.2.1

**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

## R topics documented:

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get_hyp_mat	<i>Function to compute the hypotheses to test (closed test)</i>
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## Description

Function to compute the hypotheses to test (closed test)

**Usage**

```
get_hyp_mat(n_hypothesis = 3, selected_hypothesis = 1)
```

**Arguments**

n\_hypothesis     num elementary hypotheses  
selected\_hypothesis  
                  selected hypothesis for closed test

**Details**

eWHORM simulations

**Value**

maximum value in a row

**Author(s)**

Marta Bofill Roig

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get_max_col_index	<i>Function to get the column index of the maximum value in a row</i>
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**Description**

Function to get the column index of the maximum value in a row

**Usage**

```
get_max_col_index(row)
```

**Arguments**

row                    selected row

**Details**

eWHORM simulations

**Value**

maximum value in a row

**Author(s)**

Marta Bofill Roig

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sim_data	<i>Simulate data from a multi-arm trial with shared control</i>
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**Description**

Function to simulate trial data (1-stage, multiple arms)

**Usage**

```
sim_data(n_arms, N, mu_6m, mu_12m, sigma, rmonth)
```

**Arguments**

n_arms	number of arms (including control)
N	total sample size
mu_6m	6-month mean response per arm (vector of length n_arm)
mu_12m	12-month mean response per arm (vector of length n_arm)
sigma	covariance matrix between 6- and 12-month responses assumed equal across arms (matrix of dim 2x2)
rmonth	recruitment per month (recruitment speed assumed constant over time)

**Details**

eWHORM simulations

**Value**

simulated data consisting of the responses at 6 and 12 months, treatment arm, and recruitment time for each subject.

**Author(s)**

Marta Bofill Roig

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sim_trial	<i>Simulate data from a multi-arm multi-stage trial with shared control and dose selection</i>
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**Description**

Function to simulate trial data (2-stages, with dose selection)

**Usage**

```

sim_trial(
  n_arms = 4,
  N1 = 30 * 4,
  N2 = 30 * 2,
  mu_6m,
  mu_12m,
  sigma,
  rmonth,
  alpha1 = 0.5,
  alpha = 0.05,
  p_safety = c(0.9, 0.8, 0.7),
  safety = T
)

```

**Arguments**

n_arms	number of arms (including control)
N1	sample size stage 1
N2	sample size stage 2
mu_6m	6-month mean response per arm (vector of length n_arm)
mu_12m	12-month mean response per arm (vector of length n_arm)
sigma	covariance matrix between 6- and 12-month responses assumed equal across arms (matrix of dim 2x2)
rmonth	recruitment per month (recruitment speed assumed constant over time)
alpha1	significance level for dose selection
alpha	significance level for selected dose vs control comparison
p_safety	probability of each dose to be safe
safety	indicator - if true, it simulates safety according to p_safety

**Details**

eWHORM simulations

**Value**

Combined p-value, selected dose and safety for each dose (if argument safety=TRUE)

**Author(s)**

Marta Bofill Roig

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