

Package ‘gvcR’

July 22, 2025

Type Package

Title Genotypic Variance Components

Version 0.4.0

Maintainer Muhammad Yaseen <myaseen208@gmail.com>

Description Functionalities to compute model based genetic components i.e. genotypic variance, phenotypic variance and heritability for given traits of different genotypes from replicated data using methodology explained by Burton, G. W. & Devane, E. H. (1953) (<[doi:10.2134/agronj1953.00021962004500100005x](https://doi.org/10.2134/agronj1953.00021962004500100005x)>) and Allard, R.W. (2010, ISBN:8126524154).

Depends R (>= 3.2.3)

Imports dplyr, eda4treeR, lme4, magrittr, rlang, stats, supernova, tibble, R6

License GPL-3

URL <https://myaseen208.com/gvcR/> <https://github.com/MYaseen208/gvcR>

BugReports <https://github.com/myaseen208/gvcR/issues>

RoxygenNote 7.3.2

Encoding UTF-8

Suggests testthat

NeedsCompilation no

Author Muhammad Yaseen [aut, cre],
Sami Ullah [aut, ctb]

Repository CRAN

Date/Publication 2024-10-01 06:50:02 UTC

Contents

gvc	2
gvcR	4
Index	5

gvc

*Genotypic Variance***Description**

The 'gvc' class calculates genotypic variance, phenotypic variance, and broad-sense heritability from replicated data.

Details

The 'gvc' class uses methods explained by Burton, G. W. & Devane, E. H. (1953) and Allard, R.W. (2010). It includes methods for calculating genetic variance, phenotypic variance, and heritability.

Public fields

data A 'tibble' containing the data for analysis.

y The name of the response variable.

x The name of the covariate (optional).

rep The name of the replicate factor.

gen The name of the genotype factor.

env The name of the environmental factor.

Methods**Public methods:**

- [gvc\\$new\(\)](#)
- [gvc\\$calculate_gvar\(\)](#)
- [gvc\\$calculate_pvar\(\)](#)
- [gvc\\$calculate_herit\(\)](#)
- [gvc\\$clone\(\)](#)

Method `new()`: Initialize the 'gvc' class with the data and variable names.

Usage:

```
gvc$new(.data, .y, .x = NULL, .rep, .gen, .env)
```

Arguments:

.data A 'data.frame' containing the data for analysis.

.y The response variable.

.x The covariate (optional).

.rep The replicate factor.

.gen The genotype factor.

.env The environmental factor.

Returns: An instance of the 'gvc' class.

Method `calculate_gvar()`: Calculate genetic variance.

Usage:

```
gvc$calculate_gvar()
```

Returns: A list with the genetic variance ('gvar').

Method `calculate_pvar()`: Calculate phenotypic variance.

Usage:

```
gvc$calculate_pvar()
```

Returns: A list with the phenotypic variance ('pvar').

Method `calculate_herit()`: Calculate broad-sense heritability.

Usage:

```
gvc$calculate_herit()
```

Returns: A list with the heritability ('h2').

Method `clone()`: The objects of this class are cloneable with this method.

Usage:

```
gvc$clone(deep = FALSE)
```

Arguments:

`deep` Whether to make a deep clone.

Examples

```
df1 <- data.frame(
  Response = c(
    rnorm(48, mean = 15000, sd = 500)
    , rnorm(48, mean = 5000, sd = 500)
    , rnorm(48, mean = 1000, sd = 500)
  )
, Rep      = as.factor(rep(1:3, each = 48))
, Variety  = gl(n = 4, k = 4, length = 144, labels = letters[1:4])
, Env      = gl(n = 3, k = 16, length = 144, labels = letters[1:3])
)

# Create an instance of the class
gvc1 <- gvc$new(
  .data = df1
  , .y   = Response
  , .rep = Rep
  , .gen = Variety
  , .env = Env
)

# Calculate genetic variance (gvar)
gvc1$calculate_gvar()

# Calculate phenotypic variance (pvar)
gvc1$calculate_pvar()
```

```
# Calculate heritability (h2)
gvc1$calculate_herit()
```

gvcR

Genotypic Variance Components

Description

Functionalities to compute model based genetic components i.e genotypic, phenotypic variances and heritability for given traits of different genotypes from replicated data using methodology explained by Burton, G. W. & Devane, E. H. (1953) (<doi:10.2134/agronj1953.00021962004500100005x>) and Allard, R.W. (2010, ISBN:8126524154).

Author(s)

1. Sami Ullah (<samiullahuos@gmail.com>)
2. Muhammad Yaseen (<myaseen208@gmail.com>)

References

1. R.K. Singh and B.D.Chaudhary *Biometrical Methods in Quantitative Genetic Analysis*. Kalyani Publishers, New Delhi
1. Williams, E.R., Matheson, A.C. and Harwood, C.E. (2002). *Experimental Design and Analysis for Tree Improvement*. CSIRO Publishing.

Index

gvc, [2](#)
gvcR, [4](#)