

# Package ‘corrsieve’

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**Title** Software for Summarising and Evaluating STRUCTURE Output

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**Description** Statistical summary of STRUCTURE output. STRUCTURE is a K-means clustering method for inferring population structure and assigning individuals to populations using genetic data. Pritchard JK, Stephens M, Donnelly PJ (2000) <DOI:10.1093/genetics/155.2.945>. <<https://web.stanford.edu/group/pritchardlab/structure.html>>.

**License** GPL (>= 3)

**Depends** methods, stats

**URL** <https://github.com/campanam/rCorrSieve>

**BugReports** <https://github.com/campanam/rCorrSieve>

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calc.delta	<i>Calc.delta</i>
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**Description**

Calculates delta Fst or delta K from the output of `summarise.Fst` or `summarise.lnPD`.

**Usage**

```
calc.delta(input, Fst = FALSE)
```

**Arguments**

input	a table containing Fst or lnPD data generated by <code>summarise.Fst</code> or <code>summarise.lnPD</code> .
Fst	when FALSE, data is lnPD data and calculates delta K. When true, data is Fst data and calculates delta Fst

**Value**

Returns a table listing K values and delta F or delta K statistics

**Author(s)**

Michael G. Campana <[mcampana63@gmail.com](mailto:mcampana63@gmail.com)>

**See Also**

[summarise.Fst](#) [summarise.lnPD](#)

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corr.Qmatrix	<i>Corr.Qmatrix</i>
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**Description**

Calculates Q matrix correlations from structure files in the folder specified in the `filepath` option

**Usage**

```
corr.Qmatrix(filepath = "./", instruct = FALSE, rowncol = TRUE,  
             avmax = TRUE, pvalue = FALSE, raw = TRUE, r = 0.99, p = 0.05)
```

**Arguments**

filepath	a character string listing the folder's path from the current directory
instruct	when TRUE, data is in INSTRUCT format, else data is in STRUCTURE format
rowncol	when TRUE, calculates and returns filtered Q matrix correlations using the rows-and-columns criterion
avmax	when TRUE, calculates and returns filtered Q matrix correlations using the average maximum correlation criterion
pvalue	when TRUE, calculates and returns Q matrix correlations using permutation tests
raw	when TRUE, returns the raw unfiltered Q matrix correlations
r	the minimum r value to classify a correlation as significant
p	the maximum p value to classify a correlation as significant. Ignored unless pvalue = TRUE

**Value**

Returns a S4 object of class QmatrixFilt listing Q matrix correlation results for all STRUCTURE results files in the designated folder

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

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matrixCorr

*MatrixCorr*

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**Description**

The S4 class matrixCorr lists raw, unfiltered Q matrices between Structure runs

**Objects from the Class**

Objects can be created by calls of the form `new("matrixCorr", ...)`.

**Slots**

**K** A numeric listing the K value of the runs correlated

**Run1** A numeric identifying the first of the runs correlated

**Run2** A numeric identifying the second of the runs correlated

**CorrMatrix** A matrix listing raw Q matrix correlations

**Pvalues** A matrix listing raw Q matrix correlation significances

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

**See Also**[matrixCorr](#)

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matrixCorr-method	<i>MatrixCorr</i> constructor
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**Description**

Constructor for [matrixCorr](#) objects

**Usage**

```
matrixCorr(K, Run1, Run2, CorrMatrix, Pvalues = matrix(NA))
```

**Arguments**

K	A numeric corresponding to the @K slot listing the K value of the runs correlated
Run1	A numeric corresponding to the @Run1 slot identifying the first of the runs correlated
Run2	A numeric corresponding to the @Run2 slot identifying the second of the runs correlated
CorrMatrix	A matrix corresponding to the @CorrMatrix slot listing raw Q matrix correlations
Pvalues	A matrix corresponding to the @Pvalues slot listing raw Q matrix correlation significances

**Value**

Returns a S4 object of class `matrixCorr` listing raw Q matrix correlation results

**Author(s)**

Michael G. Campana <[mcampana63@gmail.com](mailto:mcampana63@gmail.com)>

**See Also**[matrixCorr](#)**Examples**

```
test <- matrixCorr(K = 1, Run1 = 2, Run2 = 3, CorrMatrix = matrix(NA))
```

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QmatrixFilt

*QmatrixFilt*


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### Description

The S4 class QmatrixFilt lists for Q matrix correlation output

### Objects from the Class

Objects can be created by calls of the form `new("QmatrixFilt", ...)`.

### Slots

**rowncol** A list listing filtered Q matrix correlations by the rows-and-columns method

**avmaxcorr** A table listing filtered Q matrix correlations by the rows-and-columns method

**rawcorr** A list listing raw Q matrix correlations

### Author(s)

Michael G. Campana <[mcampana63@gmail.com](mailto:mcampana63@gmail.com)>

### See Also

[QmatrixFilt](#)

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QmatrixFilt-method

*QmatrixFilt constructor*


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### Description

Constructor for [QmatrixFilt](#) objects

### Usage

```
QmatrixFilt(rowncol = list(""), avmaxcorr = as.table(matrix(NA)), rawcorr = list(""))
```

### Arguments

**rowncol** A list corresponding to the @rowncol slot listing filtered Q matrix correlations by the rows-and-columns method

**avmaxcorr** A table corresponding to the @avmaxcorr slot listing filtered Q matrix correlations by the rows-and-columns method

**rawcorr** A list corresponding to the @rawcorr slot listing raw Q matrix correlations

**Value**

Returns a S4 object of class `QmatrixFilt` listing Q matrix correlation results

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

**See Also**

[QmatrixFilt](#)

**Examples**

```
test <- QmatrixFilt(rowncol = list(c("a", "b", "c")))
test@rowncol
```

---

read.struct

*Read.struct*

---

**Description**

Reads the K values, Fsts, lnPDs from structure files in the folder specified in the filepath option

**Usage**

```
read.struct(filepath = "./", instruct = FALSE)
```

**Arguments**

filepath            a character string listing the folder's path from the current directory  
instruct            when TRUE, data is in INSTRUCT format, else data is in STRUCTURE format

**Value**

Returns a table listing K values, lnPDs and Fsts for all STRUCTURE results files in the designated folder

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

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rowncolMatrix	<i>RowncolMatrix</i>
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**Description**

The S4 class RowncolMatrix lists filtered Q matrix output by the row-and-column method

**Objects from the Class**

Objects can be created by calls of the form `new("rowncolMatrix", ...)`.

**Slots**

**K** A numeric listing the K value of the runs correlated

**filterMatrix** A table listing filtered Q matrix correlations by the row-and-column method

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

**See Also**

[rowncolMatrix](#)

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rowncolMatrix-method	<i>RowncolMatrix constructor</i>
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**Description**

Constructor for [rowncolMatrix](#) objects

**Usage**

```
rowncolMatrix(K, filtermatrix)
```

**Arguments**

**K** A numeric corresponding to the @K slot listing the K value of the runs correlated

**filtermatrix** A table corresponding to the @filtermatrix slot listing filtered Q matrix correlations

**Value**

Returns a S4 object of class rowncolMatrix listing raw Q matrix correlation results

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

**See Also**

[rowncolMatrix](#)

**Examples**

```
## Make a table of correlation determinations
filtmat <- table(matrix(c("Y","Y","Y",NA,"Y","Y",NA,NA,"Y"),ncol = 3, byrow = TRUE))
## Make a rowncolMatrix
test <- rowncolMatrix(K = 3, filtermatrix = filtmat)
```

---

summarise.Fst

*Summarise.Fst*

---

**Description**

Summarises Fst from structure output read by read.struct.

**Usage**

```
summarise.Fst(input, stdevopt = 1)
```

**Arguments**

input	a table containing InPD Fst generated by read.struct
stdevopt	Chooses the optimisation procedure for the Fst summaries. 1: no optimisation, 2: order the clusters by value, 3: order the clusters by correlation coefficients

**Value**

Returns a table listing K values and summarised Fst statistics

**Author(s)**

Michael G. Campana <mcampana63@gmail.com>

**See Also**

[read.struct](#) [calc.delta](#)

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`summarise.lnPD`

*Summarise.lnPD*

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**Description**

Summarises lnP(D) from structure output read by `read.struct`.

**Usage**

`summarise.lnPD(input)`

**Arguments**

`input` a table containing lnPD data generated by `read.struct`

**Value**

Returns a table listing K values and summarised lnPD statistics

**Author(s)**

Michael G. Campana <[mcampana63@gmail.com](mailto:mcampana63@gmail.com)>

**See Also**

[read.struct](#) [calc.delta](#)

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