

# Package ‘magree’

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**Type** Package

**Title** Implements the O’Connell-Dobson-Schouten Estimators of Agreement for Multiple Observers

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**Depends** graphics

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**Description** Implements an interface to the legacy Fortran code from O’Connell and Dobson (1984) <DOI:10.2307/2531148>. Implements Fortran 77 code for the methods developed by Schouten (1982) <DOI:10.1111/j.1467-9574.1982.tb00774.x>. Includes estimates of average agreement for each observer and average agreement for each subject.

**License** GPL-3 | GPL-2

**LazyData** yes

**NeedsCompilation** yes

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landis	<i>Landis and Koch dataset.</i>
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### Description

Canonical dataset for agreement for multiple observers described in Landis and Koch (Biometrics 1977; 33: 363-374).

### Usage

```
data("landis")
```

### Format

The format is: int [1:118, 1:7] 4 1 3 4 3 2 1 3 2 1 ... - attr(\*, "dimnames")=List of 2 ..\$ : chr [1:118] "1" "2" "3" "4" ... ..\$ : chr [1:7] "A" "B" "C" "D" ...

### Source

Landis and Koch (Biometrics 1977; 33: 363-374)

### Examples

```
data(landis)
## maybe str(landis) ; plot(landis) ...
```

---

magree	<i>O'Connell-Dobson-Schouten estimators for multiobserver agreement.</i>
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### Description

Use the O'Connell-Dobson-Schouten estimators of agreement for nominal or ordinal data.

### Usage

```
magree(X, weights=c("unweighted", "linear", "quadratic"), score = NULL)
```

### Arguments

X	A matrix or data-frame with observations/subjects as rows and raters as columns.
weights	<b>"unweighted"</b> For nominal categories - only perfect agreement is counted. <b>"linear"</b> For ordinal categories where disagreement is proportional to the distance between the categories. This is analogous to the agreement weights $w_{i,j} = 1 -  score[i] - score[j]  / (max(score) - min(score))$ .

	<b>"quadratic"</b> For ordinal categories where disagreement is proportional to the square of the distance between the categories. This is analogous to the agreement weights $w_{i,j} = 1 - (score[i] - score[j])^2 / (max(score) - min(score))^2$ .
score	The scores that are to be assigned to the categories. Currently, this defaults to a sorted list of the unique values.

## Details

The Fortran code from Professor Dianne O'Connell was adapted for R.

The output object is very similar to the Fortan code. Not all of the variance terms are currently used in the `print` and `summary` methods.

## Value

<code>oconnell</code>	object from the <code>oconnell</code> function
<code>schouten</code>	object from the <code>schouten</code> function
<code>call</code>	As per <code>sys.call()</code> , to allow for using <code>update</code>

## See Also

[oconnell](#), [schouten](#).

## Examples

```
## Table 1 (O'Connell and Dobson, 1984)
summary(fit <- magree(landis, weights="unweighted"))
update(fit, weights="linear")
update(fit, weights="quadratic")

## Table 5, O'Connell and Dobson (1984)
magree(landis==1)
magree(landis==2)
magree(landis==3)
magree(landis==4)
magree(landis==5)

## Plot of the marginal distributions
plot(fit)

## Plot of the average agreement by observer
plot(fit, type="kappa by observer")
```

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 oconnell

*O'Connell-Dobson estimators for multiobserver agreement.*


---

### Description

Use the O'Connell-Dobson estimator of agreement for nominal or ordinal data. This includes a range of statistics on agreement for assuming either distinct or homogeneous items.

### Usage

```
oconnell(X, weights=c("unweighted", "linear", "quadratic"), i=NULL, score = NULL)
```

### Arguments

- |         |   |
|---------|---|
| X       | A matrix or data-frame with observations/subjects as rows and observers as columns.   |
| weights | <p><b>"unweighted"</b> For nominal categories - only perfect agreement is counted.</p> <p><b>"linear"</b> For ordinal categories where disagreement is proportional to the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 -  score[i] - score[j]  / (max(score) - min(score))</math>.</p> <p><b>"quadratic"</b> For ordinal categories where disagreement is proportional to the square of the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 - (score[i] - score[j])^2 / (max(score) - min(score))^2</math>.</p>   |
| i       | <ol style="list-style-type: none"> <li>1. For nominal categories - only perfect agreement is counted.</li> <li>2. For ordinal categories where disagreement is proportional to the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 -  score[i] - score[j]  / (max(score) - min(score))</math>.</li> <li>3. For ordinal categories where disagreement is proportional to the square of the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 - (score[i] - score[j])^2 / (max(score) - min(score))^2</math>.</li> </ol> <p>This argument takes precedence over weights if it is specified.</p> |
| score   | The scores that are to be assigned to the categories. Currently, this defaults to 1:L, where codeL is the number of categories.   |

### Details

The Fortran code from Professor Dianne O'Connell was adapted for R.

The output object is very similar to the Fortan code. Not all of the variance terms are currently used in the print, summary and plot methods.

**Value**

X	As input
i	As input
nrater	Number of observers
nscore	Number of categories
nsubj	Number of subjects
p1[j, k]	Probability of observer j giving score k when observers are distinct
p2[k]	Probability of score k when observers are homogeneous
w1[j, k]	Weighted average of d[] for observer j, score k
w2[k]	Weighted average of d[] for score k when observers are homogeneous
d[j]	Amount of disagreement for subject j
s1[j]	Chance-corrected agreement statistic for subject j when observers are distinct
s2[j]	Chance-corrected agreement statistic for subject j when observers are homogeneous; $s[j]=1-d[j]/\text{expdel}$ .
delta[j, k]	j<k: amount of disagreement expected by change for observers j and k; j>k amount of disagreement expected by chance for observers j and k when observers are homogeneous
expd1	Amount of disagreement expected by chance in null case when observers are distinct
expd2	Amount of disagreement expected by chance when observers are homogeneous
dbar	Average value of d[] over all subjects
sav1	Chance-corrected agreement statistic over all subjects when observers are distinct
sav2	Chance-corrected agreement statistic over all subjects when observers are homogeneous
var@s1	Null variance of S when observers are distinct
var@s2	Null variance of S when observers are homogeneous
vars1	Unconstrained variance of S when observers are distinct
vars2	Unconstrained variance of S when observers are homogeneous
v@sav1	Null variance of Sav when observers are distinct
v@sav2	Null variance of Sav when observers are homogeneous
vsav1	Unconstrained variance of Sav when observers are distinct
vsav2	Unconstrained variance of Sav when observers are homogeneous
p@sav1	Probability of overall agreement due to chance when observers are distinct
p@sav2	Probability of overall agreement due to chance when observers are homogeneous
resp[i, j]	Response for observer i on subject j; transpose of X (BEWARE)
score(i)	Score associated with i'th category
call	As per <code>sys.call()</code> , to allow for using update

**See Also**

[magree, schouten.](#)

**Examples**

```
## Table 1 (O'Connell and Dobson, 1984)
summary(fit <- oconnell(landis, weights="unweighted"))
update(fit, weights="linear")
update(fit, weights="quadratic")

## Table 3 (O'Connell and Dobson, 1984)
slideTypeGroups <-
  list(c(2, 3, 5, 26, 31, 34, 42, 58, 59, 67, 70, 81, 103, 120),
       c(7, 10:13, 17, 23, 30, 41, 51, 55, 56, 60, 65, 71, 73, 76, 86, 87, 105, 111, 116, 119, 124),
       c(4, 6, 24, 25, 27, 29, 39, 48, 68, 77, 79, 94, 101, 102, 117),
       c(9, 32, 36, 44, 52, 62, 84, 95),
       c(35, 53, 69, 72),
       c(8, 15, 18, 19, 47, 64, 82, 93, 98, 99, 107, 110, 112, 115, 121),
       c(1, 16, 22, 49, 63, 66, 78, 90, 100, 113),
       c(28, 37, 40, 61, 108, 114, 118),
       106,
       43,
       83,
       c(54, 57, 88, 91, 126),
       c(74, 104),
       38,
       46,
       c(89, 122),
       c(80, 92, 96, 123),
       85)
data.frame(SlideType=1:18,
           S1=sapply(slideTypeGroups,
                    function(ids) mean(fit$s1[as.character(ids)])),
           S2=sapply(slideTypeGroups,
                    function(ids) mean(fit$s2[as.character(ids)])))

## Table 5, O'Connell and Dobson (1984)
oconnell(landis==1)
oconnell(landis==2)
oconnell(landis==3)
oconnell(landis==4)
oconnell(landis==5)

## Plot of the marginal distributions
plot(fit)
```

**Description**

plot methods for magree, oconnell and schouten objects

**Usage**

```
## S3 method for class 'magree'
plot(x, type = c("p1", "kappa by observer"),
     xlab = NULL, ylab = NULL, main = NULL, ...)
## S3 method for class 'oconnell'
plot(x, type = c("p1"), xlab = NULL, ylab = NULL, main = NULL, ...)
## S3 method for class 'schouten'
plot(x, type = c("kappa by observer"), xlab = NULL,
     ylab = NULL,
     main = NULL, xdelta = 0.1, axes = TRUE, ...)
```

**Arguments**

x	magree, oconnell or schouten object.
type	Type of plot. For "p1", plot the probabilities by observer. For "kappa by observer", plot the kappas for each observer.
xlab	
ylab	
main	
xdelta	For plot.schouten and "kappa by observer", specifies the width of the brackets for the confidence intervals.
axes	Bool for whether to plot the axes.
...	

**Examples**

```
fit <- schouten(landis)
plot(fit)
fit <- oconnell(landis)
plot(fit,type="p1")
```

---

print.magree

*print methods for magree objects*


---

**Description**

print methods for magree objects

**Usage**

```
## S3 method for class 'magree'  
print(x, ...)  
## S3 method for class 'oconnell'  
print(x, ...)  
## S3 method for class 'schouten'  
print(x, ...)
```

**Arguments**

x	the object to print
...	other arguments

**Examples**

```
print(magree(landis))
```

---

print.summary.magree *print method for summary.magree objects*

---

**Description**

print method for summary.magree objects

**Usage**

```
## S3 method for class 'summary.magree'  
print(x, ...)  
## S3 method for class 'summary.oconnell'  
print(x, ...)  
## S3 method for class 'summary.schouten'  
print(x, ...)
```

**Arguments**

x
...

**Examples**

```
summary(magree(landis))
```



schouten

*Schouten estimators for multiobserver agreement.***Description**

Use the Schouten estimator of agreement for nominal or ordinal data. This includes a range of statistics on agreement.

**Usage**

```
schouten(X, weights=c("unweighted", "linear", "quadratic", "user"), w=NULL,
score=NULL)
```

**Arguments**

X	A matrix or data-frame with subjects as rows and observers as columns.
weights	<p><b>"unweighted"</b> For nominal categories - only perfect agreement is counted.</p> <p><b>"linear"</b> For ordinal categories where disagreement is proportional to the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 -  i - j  / (c - 1)</math>.</p> <p><b>"quadratic"</b> For ordinal categories where disagreement is proportional to the square of the distance between the categories. This is analogous to the agreement weights <math>w_{i,j} = 1 - (i - j)^2 / (c - 1)^2</math>.</p> <p><b>"user"</b> An indicator for a user-defined weight matrix. The weights argument will be defined as "user" if the w argument is specified.</p>
w	A user-defined weights matrix. This argument takes precedence over weights and score if it is specified and the weight argument will be defined as "user".
score	A user-defined set of scores for each category. If this is not specified, it is assumed that score=1:L, where L is the number of categories. This is used with the weights argument to define the w matrix.

**Details**

Fortran code was written by Mark Clements based on the algorithms in Schouten (1982).

The output object is closely related to the Fortan code. Not all of the variance terms are currently used in the print, summary and plot methods.

**Value**

N	Number of subjects
M	Number of observers
L	Number of categories
data	Re-formatted X
w	Weight matrix

<code>kab</code>	Kappas between each pair of observers
<code>ka</code>	Average kappas for each observer
<code>kappa</code>	Average kappa
<code>pab, pa, p, ma, qab, qa, q, oab, eab, oa, ea, o, e, wa, wab</code>	Working fields
<code>varkab</code>	Variances for kab
<code>varka</code>	Variances for ka
<code>vark</code>	Variance for the kappa
<code>covkka</code>	Covariance term between the overall average kappa and the average kappas for each observer
<code>chi</code>	Chi-squared statistics comparing the overall average kappa and the average kappa for each observer (df=1 under the null hypothesis)
<code>pchi</code>	P-values that the overall average kappa equals the average kappa for each observer
<code>var0kab</code>	Variance for kab under the null hypothesis
<code>var0ka</code>	Variance for ka under the null hypothesis
<code>var0k</code>	Variance for the overall average kappa under the null hypothesis
<code>p0</code>	P-value for kappa=0
<code>p0a</code>	P-values that the average kappa for a observer equals zero (i.e. ka=0)
<code>weights</code>	As input
<code>X</code>	As input
<code>call</code>	As per <code>sys.call()</code> , to allow for using update

**See Also**

[magree](#), [oconnell](#).

**Examples**

```
## Weights matrix used by Schouten (1982)
w <- outer(1:5,1:5,function(x,y) ((x<=2 & y<=2) | (x>=3 & y>=3))+0)
fit <- schouten(landis,w=w) # user-defined weights

summary(fit) # Schouten (1982), Tables 2 and 5

## we can fit the same model with oconnell() or magree() using the score argument
magree(landis,score=c(1,1,2,2,2))

## plot of the average kappas by observer
plot(fit, type="kappa by observer")
```

---

summary.magree	<i>summary method for magree objects</i>
----------------	--

---

**Description**

summary method for magree objects

**Usage**

```
## S3 method for class 'magree'  
summary(object, ...)  
## S3 method for class 'oconnell'  
summary(object, ci.transform = c("logit", "identity"), ci.p = 0.95, ...)  
## S3 method for class 'schouten'  
summary(object, ci.transform = c("logit", "identity"), ci.p = 0.95, ...)
```

**Arguments**

object

ci.transform    transformation used to calculate the confidence intervals. Either "logit" for a logit transform or "identity" for no transform.

ci.p            p value for the confidence interval.

...

**Examples**

```
summary(magree(landis))
```

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