Package: pk4adi (via r-universe)

September 9, 2024

```
Type Package
Title PK for Anesthetic Depth Indicators
Version 0.1.3
Maintainer Feng Jiang <silencejiang@zju.edu.cn>
Description Calculate and compare the Anesthetic Depth Indicators PK
     values in R language The prediction probability (PK) is a
     widely used tool for the anesthetic depth indicators, which was
     first proposed by Dr. Warren D. Smith in the paper Warren D.
     Smith; Robert C. Dutton; Ty N. Smith (1996)
     <doi:10.1097/00000542-199601000-00005> and Warren D. Smith;
     Robert C. Dutton; Ty N. Smith (1996)
     <doi:10.1002/(SICI)1097-0258(19960615)15:11<1199::AID-SIM218>3.0.CO;2-Y>.
     They provide the Micro xls files to calculate and compare the
     PK values. This package provide the easy-to-use API to
     calculate and compare the PK values using the R language. The
     package's name, pk4adi, is short for ``PK for Anesthetic Depth
     Indicators".
License MIT + file LICENSE
Encoding UTF-8
LazyData true
URL https://www.r-project.org, https://github.com/xfz329/rpk4adi
BugReports https://github.com/xfz329/rpk4adi/issues
RoxygenNote 7.2.3
Imports data.table (>= 1.10), stats
Repository https://xfz329.r-universe.dev
RemoteUrl https://github.com/xfz329/rpk4adi
RemoteRef HEAD
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RemoteSha 7b64517281f1acdd3e7433a70be5f151b5b55a36

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Description

Compute the PK value to Measure the Performance of Anesthetic Depth Indicators.

Usage

```
calculate_pk(x_in, y_in)
```

Arguments

x_in a vector, the indicator.y_in a vector, the state.

Value

a list containing all the matrices and variables during the calculation. The value list\$type is "PK", which indicated the list is the return-value of the function calculate_pk(). The type of list\$basic is also a list, which contains the most important results of the function. The type of list\$matrices is also a list, which contains all the matrices during the calculation. The type of list\$details is also a list, which contains all the intermediate variables during the calculation.

References

Warren D. Smith, Robert C. Dutton, Ty N. Smith; Measuring the Performance of Anesthetic Depth Indicators. Anesthesiology 1996; 84:38–51 doi: https://doi.org/10.1097/00000542-199601000-00005.

Warren D. Smith, Robert C. Dutton, Ty N. Smith; A measure of association for assessing prediction accuracy that is a generalization of nonparametric ROC area. Statistics in Medicine 1996; 15: 1119-1215 doi: https://doi.org/10.1002/(SICI)1097-0258(19960615)15:11<1199::AID-SIM218>3.0.CO;2-Y.

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Examples

```
x1 <- c(0, 0, 0, 0, 0, 0)
y1 <- c(1, 1, 1, 1, 1, 2)
ans1 <- calculate_pk(x1, y1)

## show the most important results.
print(ans1$basic)

x2 <- c(1, 1, 2, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 4, 5, 5, 5, 5, 6, 6, 6, 6, 6)
y2 <- c(1, 1, 1, 1, 1, 2, 1, 1, 3, 3, 2, 2, 2, 2, 1, 3, 3, 3, 3, 3, 3, 3)
ans2 <- calculate_pk(x2, y2)

## show the full results.
print(ans2)</pre>
```

compare_pks

Compare two answers of the PK values.

Description

Both of the two input have to be the output of the function calculate pk().

Usage

```
compare_pks(pk1, pk2)
```

Arguments

```
pk1 a list, the output of the function calculate_pk().
pk2 a list, the output of the function calculate_pk().
```

Value

a list containing all the variables during the calculation. The value list\$type is "PKC", which indicated the list is the return-value of the function compare_pk(). The type of list\$group is also a list, which contains the normal distribution test results for the group variables. The type of list\$pair is also a list, which contains the t distribution test results for the pair variables. The type of list\$details is also a list, which contains all the intermediate variables during the calculation.

References

Warren D. Smith, Robert C. Dutton, Ty N. Smith; Measuring the Performance of Anesthetic Depth Indicators. Anesthesiology 1996; 84:38–51 doi: https://doi.org/10.1097/00000542-199601000-00005.

Warren D. Smith, Robert C. Dutton, Ty N. Smith; A measure of association for assessing prediction accuracy that is a generalization of nonparametric ROC area. Statistics in Medicine 1996; 15: 1119-1215 doi: https://doi.org/10.1002/(SICI)1097-0258(19960615)15:11<1199::AID-SIM218>3.0.CO;2-Y.

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Examples

```
x1 <- c(1, 1, 2, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 4, 5, 5, 5, 5, 6, 6, 6, 6, 6)
y1 <- c(1, 1, 1, 1, 1, 1, 2, 1, 1, 3, 3, 2, 2, 2, 2, 2, 1, 3, 3, 3, 3, 3, 3, 3, 3)

pk1 <- calculate_pk(x_in = x1, y_in = y1)
print(pk1$basic)

x2 <- c(1, 1, 2, 2, 2, 2, 3, 3, 3, 3, 4, 4, 4, 4, 4, 5, 5, 5, 5, 5, 6, 6, 6, 6, 6)
y2 <- c(1, 1, 2, 1, 1, 2, 1, 2, 3, 3, 2, 2, 1, 2, 2, 2, 3, 3, 3, 3, 2, 3, 3, 2)

pk2 <- calculate_pk(x_in = x2, y_in = y2)
print(pk2$basic)

ans <- compare_pks(pk1, pk2)
print(ans$group)
print(ans$pair)</pre>
```

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