# Package 'micer'

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Type Package
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Title Map Image Classification Efficacy

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**Description** Map image classification efficacy (MICE) adjusts the accuracy rate relative to a random classification base-

line (Shao et al. (2021)<doi:10.1109/ACCESS.2021.3116526> and Tang et al. (2024)<doi:10.1109/TGRS.2024.3446950>) portions from the reference labels are considered, as opposed to the proportions from the reference and predictions, as is the case for the Kappa statistic. This package offers means to calculate MICE and adjusted versions of class-level user's accuracy (i.e., precision) and producer's accuracy (i.e., recall) and F1-scores. Class-level metrics are aggregated using macroaveraging. Functions are also made available to estimate confidence intervals using bootstrapping and statistically compare two classification results.

**Depends** R (>= 2.10)

**Imports** dplyr (>= 1.1.3),

License GPL (>= 3)

URL https://github.com/maxwell-geospatial/micer

BugReports https://github.com/maxwell-geospatial/micer/issues

NeedsCompilation no

**Encoding UTF-8** 

RoxygenNote 7.3.2

Suggests knitr, rmarkdown

VignetteBuilder knitr

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Repository CRAN

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# **Description**

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Example binary classification dataset. "Mine" is the positive case and "Not Mine" is the background class. There are 178 samples from the "Mine" class and 4,822 samples from the "Not Mine" class. Counts are relative to reference labels. Class proportions are based on landscape proportions. There are a total of 5,000 samples.

#### **Format**

ref reference label pred predicted label

# References

Maxwell, A.E., Bester, M.S., Guillen, L.A., Ramezan, C.A., Carpinello, D.J., Fan, Y., Hartley, F.M., Maynard, S.M. and Pyron, J.L., 2020. Semantic segmentation deep learning for extracting surface mine extents from historic topographic maps. Remote Sensing, 12(24), p.4145.

compareData

Data for multiclass classification comparison

# **Description**

Example multiclass classification dataset with the following wetland-related classes: "PFO", "PEM", "RLP", and "Not". PFO = Palustrine Forested; PEM = Palustrine Emergent; RLP = River, Lake, Pond; Not = Not Wetland. There are 600 examples from each class relative to the reference labels.

#### **Format**

ref correct label rfRred random forest prediction dfRred single decision tree prediction mcData 3

# References

These data are unpublished

mcData

Example multiclass classification dataset

# **Description**

Example multiclass classification dataset with the following classes (counts relative to reference labels): "Barren" (n=163), "Forest" (n=20,807), "Impervious" (n=426), "Low Vegetation" (n=3,182), "Mixed Dev" (n=520), and "Water" (n=200). There are a total of 25,298 samples.

#### **Format**

```
ref reference labelpred predicted label
```

#### References

Maxwell, A.E., Strager, M.P., Warner, T.A., Ramezan, C.A., Morgan, A.N. and Pauley, C.E., 2019. Large-area, high spatial resolution land cover mapping using random forests, GEOBIA, and NAIP orthophotography: Findings and recommendations. Remote Sensing, 11(12), p.1409.

mice mice

# Description

Calculate map image classification efficacy (MICE) and other metrics using columns/vectors of reference and predicted classes

# Usage

```
mice(
  reference,
  prediction,
  mappings = levels(as.factor(reference)),
  multiclass = TRUE,
  positiveIndex = 1
)
```

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# Arguments

reference column/vector of reference labels as factor data type.

prediction column/vector of predicted labels as factor data type.

mappings names of classes (if not provided, factor levels are used).

multiclass TRUE or FALSE. If TRUE, treats classification as multiclass. If FALSE, treats

classification as binary. Default is TRUE.

positiveIndex index for positive case for binary classification. Ignored for multiclass classifi-

cation. Default is 1 or first factor level.

#### **Details**

For multiclass classification, returns a list object with the following items: \$Mappings = class names; \$confusionMatrix = confusion matrix where columns represent the reference data and rows represent the classification result; \$referenceCounts = count of samples in each reference class; \$predictionCounts = count of predictions in each class; \$overallAccuracy = overall accuracy; \$MICE = map image classification efficacy; \$usersAccuracies = class-level user's accuracies (1 - commission error); \$CTBICEs = classification-total-based image classification efficacies (adjusted user's accuracies); \$producersAccuracies = class-level producer's accuracies (1 - omission error); \$RTBICEs = reference-total-based image classification efficacies (adjusted producer's accuracies); \$F1Scores = class-level harmonic mean of user's and producer's accuracies; \$F1Efficacies = F1-score efficacies; \$macroPA = class-aggregated, macro-averaged producer's accuracy; \$macroRTBICE = class-aggregated, macro-averaged reference-total-based image classification efficacy; \$macroCTBICE = class-aggregated, macro-averaged user's accuracy; \$macroCTBICE = class-aggregated, macro-averaged classification-total-based image classification efficacy; \$macroF1 = class-aggregated, macro-averaged F1-score; \$macroF1Efficacy = class-aggregated, macro-averaged F1 efficacy;

For binary classification, returns a list object with the following items: \$Mappings = class names; \$confusionMatrix = confusion matrix where columns represent the reference data and rows represent the classification result; \$referenceCounts = count of samples in each reference class; \$predictionCounts = count of predictions in each class; \$postiveCase = name or mapping for the positive case; \$overallAccuracy = overall accuracy; \$MICE = map image classification efficacy; \$Precision = precision (1 - commission error relative to positive case); \$precisionEfficacy = precision efficacy; \$NPV = negative predictive value (1 - commission error relative to negative case); \$npvEfficacy = negative predictive value efficacy; \$Recall = recall (1 - omission error relative to positive case); \$recallEfficacy = recall efficacy; \$specificity = specificity (1 - omission error relative to negative case); \$specificityEfficacy = specificity efficacy; \$f1Score = harmonic mean of precision and recall; \$f1Efficacy = F1-score efficacy;

# Value

multiclass or binary assessment metrics in a list object. See details for description of generated metrics.

# **Examples**

#Multiclass example
data(mcData)

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```
mice(mcData$ref,
mcData$pred,
mappings=c("Barren", "Forest", "Impervious", "Low
Vegetation", "Mixed Dev", "Water"),
multiclass=TRUE)

#Binary example
data(biData)
mice(biData$ref,
biData$pred,
mappings = c("Mined", "Not Mined"),
multiclass=FALSE,
positiveIndex=1)
```

miceCI

miceCI

# **Description**

Calculate confidence intervals (CIs) for MICE and associated metrics using bootstrap sampling and the percentile method.

# Usage

```
miceCI(
  reps = 200,
  frac = 0.7,
  lowPercentile,
  highPercentile,
  reference,
  prediction,
  mappings = levels(as.factor(reference)),
  multiclass = TRUE,
  positiveIndex = 1
)
```

# **Arguments**

reps number of bootstrap replicates to use. Default is 200.

frac proportion of samples to include in each bootstrap sample. Default is 0.7.

lowPercentile lower percentile for confidence interval. Default is 0.025 for a 95% CI.

highPercentile upper percentile for confidence interval. Default is 0.975 for a 95% CI.

reference column of reference labels as factor data type.

prediction column of predicted labels as factor data type.

mappings names of classes (if not provided, factor levels are used).

multiclass TRUE or FALSE. If TRUE, treats classification as multiclass. If FALSE, treats

classification as binary. Default is TRUE.

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positiveIndex index for positive case for binary classification. Ignored for multiclass classification. Default is 1 or first factor level

# **Details**

Confidence intervals are estimated for overall accuracy, MICE, and all class-aggregated, macro-averaged metrics produced by mice() or miceCM(). Returns metric name, mean metric value, median metric value, lower confidence interval bounds (low.ci), and upper confidence interval bounds (upper.ci) as a dataframe object.

#### Value

dataframe object of metric name and estimated mean value, median value, and lower and upper CIs.

# **Examples**

```
#Multiclass example
data(mcData)
ciResultsMC <- miceCI(rep=100,</pre>
frac=.7,
mcData$ref,
mcData$pred,
lowPercentile=0.025,
highPercentile=0.975,
mappings=c("Barren", "Forest", "Impervious", "Low Vegetation", "Mixed Dev", "Water"),
multiclass=TRUE)
print(ciResultsMC)
#Binary example
data(biData)
ciResultsBi <- miceCI(rep=100,</pre>
frac=.7,
biData$ref,
biData$pred,
lowPercentile=0.025,
highPercentile=0.975,
mappings = c("Mined", "Not Mined"),
multiclass=FALSE,
positiveIndex=1)
print(ciResultsBi)
```

miceCM

miceCM

# Description

Calculate map image classification efficacy (MICE) and other metrics using confusion matrix

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# Usage

```
miceCM(
   cm,
   mappings = levels(as.factor(row.names(cm))),
   multiclass = TRUE,
   positiveIndex = 1
)
```

# **Arguments**

cm confusion matrix as table object where rows define predictions and columns

define reference labels.

mappings names of classes (if not provided, factor levels are used).

multiclass TRUE or FALSE. If TRUE, treats classification as multiclass. If FALSE, treats

classification as binary. Default is TRUE.

positiveIndex index for positive case for binary classification. Ignored for multiclass classifi-

cation. Default is 1 or first factor level.

#### **Details**

For multiclass classification, returns a list object with the following items: \$Mappings = class names; \$confusionMatrix = confusion matrix where columns represent the reference data and rows represent the classification result; \$referenceCounts = count of samples in each reference class; \$predictionCounts = count of predictions in each class; \$overallAccuracy = overall accuracy; \$MICE = map image classification efficacy; \$usersAccuracies = class-level user's accuracies (1 - commission error); \$CTBICEs = classification-total-based image classification efficacies (adjusted user's accuracies); \$producersAccuracies = class-level producer's accuracies (1 - omission error); \$RTBICEs = reference-total-based image classification efficacies (adjusted producer's accuracies); \$F1Scores = class-level harmonic mean of user's and producer's accuracies; \$F1Efficacies = F1-score efficacies; \$macroPA = class-aggregated, macro-averaged producer's accuracy; \$macroRTBICE = class-aggregated, macro-averaged reference-total-based image classification efficacy; \$macroCTBICE = class-aggregated, macro-averaged classification-total-based image classification efficacy; \$macroF1 = class-aggregated, macro-averaged F1-score; \$macroF1Efficacy = class-aggregated, macro-averaged F1 efficacy;

For binary classification, returns a list object with the following items: \$Mappings = class names; \$confusionMatrix = confusion matrix where columns represent the reference data and rows represent the classification result; \$referenceCounts = count of samples in each reference class; \$predictionCounts = count of predictions in each class; \$postiveCase = name or mapping for the positive case; \$overallAccuracy = overall accuracy; \$MICE = map image classification efficacy; \$Precision = precision (1 - commission error relative to positive case); \$precisionEfficacy = precision efficacy; \$NPV = negative predictive value (1 - commission error relative to negative case); \$npvEfficacy = negative predictive value efficacy; \$Recall = recall (1 - omission error relative to positive case); \$recallEfficacy = recall efficacy; \$specificity = specificity (1 - omission error relative to negative case); \$specificityEfficacy = specificity efficacy; \$f1Score = harmonic mean of precision and recall; \$f1Efficacy = F1-score efficacy;

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# Value

multiclass or binary assessment metrics in a list object. See details for description of generated metrics.

# **Examples**

```
#Multiclass example
data(mcData)
cmMC <- table(mcData$pred, mcData$ref)
miceCM(cmMC,
mappings=c("Barren", "Forest", "Impervious", "Low Vegetation", "Mixed Dev", "Water"),
multiclass=TRUE)

#Binary example
data(biData)
cmB <- table(biData$pred, biData$ref)
miceMCResult <- miceCM(cmB,
mappings=c("Mined", "Not Mined"),
multiclass=FALSE,
positiveIndex=1)
print(miceMCResult)</pre>
```

miceCompare

miceCompare

# **Description**

Statistically compare two models using a paired t-test and bootstrap samples of the assessment results

# Usage

```
miceCompare(ref, result1, result2, reps, frac)
```

# Arguments

ref column of reference labels as factor data type.

result1 column of predicted labels as factor data type (first result to compare).

column of predicted labels as factor data type (second result to compare).

reps number of bootstrap replicates to use. Default is 200.

frac proportion of samples to include in each bootstrap sample. Default is 0.7.

#### Value

paired t-test results including t-statistic, degrees of freedom, p-value, 95% confidence interval, and mean difference

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# Examples

```
data(compareData)
compareResult <- miceCompare(ref=compareData$ref,
result1=compareData$rfPred,
result2=compareData$dtPred,
reps=100,
frac=.7)
print(compareResult)</pre>
```

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