Package: fairness (via r-universe)

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Title Algorithmic Fairness Metrics

Version 1.2.2

Maintainer Nikita Kozodoi <n.kozodoi@icloud.com>

Description Offers calculation, visualization and comparison of algorithmic fairness metrics. Fair machine learning is an emerging topic with the overarching aim to critically assess whether ML algorithms reinforce existing social biases. Unfair algorithms can propagate such biases and produce predictions with a disparate impact on various sensitive groups of individuals (defined by sex, gender, ethnicity, religion, income, socioeconomic status, physical or mental disabilities). Fair algorithms possess the underlying foundation that these groups should be treated similarly or have similar prediction outcomes. The fairness R package offers the calculation and comparisons of commonly and less commonly used fairness metrics in population subgroups. These methods are described by Calders and Verwer (2010) <doi:10.1007/s10618-010-0190-x>, Chouldechova (2017) <doi:10.1089/big.2016.0047>, Feldman et al. (2015) <doi:10.1145/2783258.2783311>, Friedler et al. (2018) <doi:10.1145/3287560.3287589> and Zafar et al. (2017) <doi:10.1145/3038912.3052660>. The package also offers convenient visualizations to help understand fairness metrics.

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LazyData true

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URL https://kozodoi.me/r/fairness/packages/2020/05/01/fairness-tutorial.html

BugReports https://github.com/kozodoi/fairness/issues

Depends R (>= 3.5.0)

Imports caret, devtools, e1071, ggplot2, pROC

Suggests testthat, knitr, rmarkdown

VignetteBuilder knitr Repository https://kozodoi.r-universe.dev RemoteUrl https://github.com/kozodoi/fairness RemoteRef HEAD RemoteSha 62c545802010af3b89d2eb8aaaf9dfba0bcfc3b8

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acc_parity Accuracy parity

Description

This function computes the Accuracy parity metric Formula: (TP + TN) / (TP + FP + TN + FN)

Usage

```
acc_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

acc_parity

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5 .
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number $>= 2$ giving the number of intervals into which group feature is to be cut.

Details

This function computes the Accuracy parity metric as described by Friedler et al., 2018. Accuracy metrics are calculated by the division of correctly predicted observations (the sum of all true positives and true negatives) with the number of all predictions. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their accuracies are lower or higher compared to the reference group. Lower accuracies will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw accuracy metrics for all groups and metrics standardized for the base group (accuracy parity metric). Lower values compared to the reference group mean lower accuracies in the selected subgroups
Metric_plot	Bar plot of Accuracy parity metric
 Probability_p	lot
	Density plot of predicted probabilities per subgroup. Only plotted if probabili-

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
acc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
acc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

ties are defined

compas

Description

compas is a landmark dataset to study algorithmic (un)fairness. This data was used to predict recidivism (whether a criminal will reoffend or not) in the USA. The tool was meant to overcome human biases and offer an algorithmic, fair solution to predict recidivism in a diverse population. However, the algorithm ended up propagating existing social biases and thus, offered an unfair algorithmic solution to the problem. In this dataset, a model to predict recidivism has already been fit and predicted probabilities and predicted status (yes/no) for recidivism have been concatenated to the original data.

Usage

compas

Format

A data frame with 6172 rows and 9 variables:

- Two_yr_Recidivism factor, yes/no for recidivism or no recidivism. This is the outcome or target in this dataset
- **Number_of_Priors** numeric, number of priors, normalized to mean = 0 and standard deviation = 1

Age_Above_FourtyFive factor, yes/no for age above 45 years or not

Age_Below_TwentyFive factor, yes/no for age below 25 years or not

Female factor, female/male for gender

Misdemeanor factor, yes/no for having recorded misdemeanor(s) or not

ethnicity factor, Caucasian, African American, Asian, Hispanic, Native American or Other

probability numeric, predicted probabilities for recidivism, ranges from 0 to 1

predicted numeric, predicted values for recidivism, 0/1 for no/yes

Source

The dataset is downloaded from Kaggle https://www.kaggle.com/danofer/compass and has undergone modifications (e.g. ethnicity was originally encoded using one-hot encoding, number or priors have been normalized, variables have been renamed, prediction model was fit and predicted probabilities and predicted status were concatenated to the original dataset).

dem_parity

Description

This function computes the Demographic parity metric

```
Formula: (TP + FP)
```

Usage

```
dem_parity(
    data,
    outcome,
    group,
    probs = NULL,
    preds = NULL,
    outcome_base = NULL,
    cutoff = 0.5,
    base = NULL,
    group_breaks = NULL
)
```

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Demographic parity metric (also known as Statistical Parity, Equal Parity, Equal Acceptance Rate or Independence) as described by Calders and Verwer 2010. Demographic parity is calculated based on the comparison of the absolute number of all positively classified individuals in all subgroups of the data. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their proportion of positively predicted observations are lower or higher compared to the reference group. Lower proportions will be reflected in numbers lower than 1 in the returned named vector.

Value

Metric	Absolute number of positive classifications for all groups and metrics standard-
	ized for the base group (demographic parity metric). Lower values compared to
	the reference group mean lower number of positively predicted observations in
	the selected subgroups
Metric_plot	Bar plot of Demographic parity metric
Probability_p	lot
	Density plot of predicted probabilities per subgroup. Only plotted if probabili-
	ties are defined

Examples

```
data(compas)
```

```
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
dem_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
dem_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

equal_odds

Equalized Odds

Description

This function computes the Equalized Odds metric Formula: TP / (TP + FN)

Usage

```
equal_odds(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
```

equal_odds

```
cutoff = 0.5,
base = NULL,
group_breaks = NULL
)
```

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5 .
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Equalized Odds metric (also known as Equal Opportunity, Positive Rate Parity or Separation). Equalized Odds are calculated by the division of true positives with all positives (irrespective of predicted values). This metrics equals to what is traditionally known as sensitivity. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their sensitivities are lower or higher compared to the reference group. Lower sensitivities will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw sensitivities for all groups and metrics standardized for the base group (equalized odds parity metric). Lower values compared to the reference group mean lower sensitivities in the selected subgroups	
Metric_plot	Bar plot of Equalized Odds metric	
Probability_plot		
	Density plot of predicted probabilities per subgroup. Only plotted if probabilities are defined	

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
equal_odds(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
equal_odds(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

fairness

fairness: Algorithmic Fairness Metrics

Description

The **fairness** package offers calculation, visualization and comparison of algorithmic fairness metrics. Fair machine learning is an emerging topic with the overarching aim to critically assess whether ML algorithms reinforce existing social biases. Unfair algorithms can propagate such biases and produce predictions with a disparate impact on various sensitive groups of individuals (defined by sex, gender, ethnicity, religion, income, socioeconomic status, physical or mental disabilities). Fair algorithms possess the underlying foundation that these groups should be treated similarly or have similar prediction outcomes. The **fairness** R package offers the calculation and comparisons of commonly and less commonly used fairness metrics in population subgroups. The package also offers convenient visualizations to help understand fairness metrics.

Details

Package:	fairness
Depends:	R (>= 3.5.0)
Type:	Package
Version:	1.2.2
Date:	2021-04-14
License:	MIT
LazyLoad:	Yes

Author(s)

- Nikita Kozodoi <n.kozodoi@icloud.com>
- Tibor V. Varga <tirgit@hotmail.com>

See Also

https://github.com/kozodoi/fairnesshttps://kozodoi.me/r/fairness/packages/2020/ 05/01/fairness-tutorial.html

fnr_parity

Description

This function computes the False Negative Rate (FNR) parity metric

```
Formula: FN / (TP + FN)
```

Usage

```
fnr_parity(
    data,
    outcome,
    group,
    probs = NULL,
    preds = NULL,
    outcome_base = NULL,
    cutoff = 0.5,
    base = NULL,
    group_breaks = NULL
)
```

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the False Negative Rate (FNR) parity metric as described by Chouldechova 2017. False negative rates are calculated by the division of false negatives with all positives (irrespective of predicted values). In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their false negative rates are lower or higher compared to the reference group. Lower false negative error rates will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean BETTER prediction for the subgroup.

Value

Metric	Raw false negative rates for all groups and metrics standardized for the base group (false negative rate parity metric). Lower values compared to the refer- ence group mean lower false negative error rates in the selected subgroups
Metric_plot	Bar plot of False Negative Rate parity metric
Probability_p	lot
	Density plot of predicted probabilities per subgroup. Only plotted if probabili- ties are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
fnr_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
fnr_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

fpr_parity

False Positive Rate parity

Description

This function computes the False Positive Rate (FPR) parity metric Formula: FP / (TN + FP)

Usage

```
fpr_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
```

fpr_parity

```
base = NULL,
group_breaks = NULL
)
```

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the False Positive Rate (FPR) parity metric as described by Chouldechova 2017. False positive rates are calculated by the division of false positives with all negatives (irrespective of predicted values). In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their false positive rates are lower or higher compared to the reference group. Lower false positives error rates will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean BETTER prediction for the subgroup.

Value

Metric_plot Bar plot of False Positives Rate metric Probability_plot Density plot of predicted probabilities per subgroup. Only plotted if probabi ties are defined	Metric	Raw false positive rates for all groups and metrics standardized for the base group (false positive rate parity metric). Lower values compared to the reference group mean lower false positive error rates in the selected subgroups
Probability_plot Density plot of predicted probabilities per subgroup. Only plotted if probabi ties are defined	Metric_plot	Bar plot of False Positives Rate metric
Density plot of predicted probabilities per subgroup. Only plotted if probabi ties are defined	Probability_plo	ot
		Density plot of predicted probabilities per subgroup. Only plotted if probabili- ties are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
fpr_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',</pre>
```

```
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
fpr_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')
```

germancredit

Modified german credit dataset

Description

germancredit is a credit scoring data set that can be used to study algorithmic (un)fairness. This data was used to predict defaults on consumer loans in the German market. In this dataset, a model to predict default has already been fit and predicted probabilities and predicted status (yes/no) for default have been concatenated to the original data.

Usage

germancredit

Format

A data frame with 1000 rows and 23 variables:

Account_status factor, status of existing checking account

Duration numeric, loan duration in month

Credit_history factor, previous credit history

Purpose factor, loan purpose

Amount numeric, credit amount

Savings factor, savings account/bonds

Employment factor, present employment since

Installment_rate numeric, installment rate in percentage of disposable income

Guarantors factor, other debtors / guarantors

Resident_since factor, present residence since

Property factor, property

Age numeric, age in years

Other_plans factor, other installment plans

Housing factor, housing

Num_credits numeric, Number of existing credits at this bank

Job factor, job

People_maintenance numeric, number of people being liable to provide maintenance for

Phone factor, telephone

Foreign factor, foreign worker

mcc_parity

BAD factor, GOOD/BAD for whether a customer has defaulted on a loan. This is the outcome or target in this dataset

Female factor, female/male for gender

probability numeric, predicted probabilities for default, ranges from 0 to 1

predicted numeric, predicted values for default, 0/1 for no/yes

Source

The dataset has undergone modifications (e.g. categorical variables were encoded, prediction model was fit and predicted probabilities and predicted status were concatenated to the original dataset).

mcc_parity Matthews Correlation Coefficient parity

Description

This function computes the Matthews Correlation Coefficient (MCC) parity metric Formula: $(TP \times TN - FP \times FN) / sqrt((TP + FP) \times (TP + FN) \times (TN + FP) \times (TN + FN))$

Usage

```
mcc_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.

cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number $>= 2$ giving the number of intervals into which group feature is to be cut.

Details

This function computes the Matthews Correlation Coefficient (MCC) parity metric. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their Matthews Correlation Coefficients are lower or higher compared to the reference group. Lower Matthews Correlation Coefficients rates will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw Matthews Correlation Coefficient metrics for all groups and metrics stan- dardized for the base group (parity metric). Lower values compared to the ref- erence group mean Matthews Correlation Coefficients in the selected subgroups	
Metric_plot	Bar plot of Matthews Correlation Coefficient metric	
Probability_plot		
	Density plot of predicted probabilities per subgroup. Only plotted if probabili- ties are defined	

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
mcc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
mcc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

npv_parity

Negative Predictive Value parity

Description

This function computes the Negative Predictive Value (NPV) parity metric Formula: TN / (TN + FN)

npv_parity

Usage

```
npv_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Negative Predictive Value (NPV) parity metric as described by the Aequitas bias toolkit. Negative Predictive Values are calculated by the division of true negatives with all predicted negatives. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their negative predictive values are lower or higher compared to the reference group. Lower negative predictive values will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric Raw negative predictive values for all groups and metrics standardized for the base group (negative predictive value parity metric). Lower values compared to the reference group mean lower negative predictive values in the selected subgroups

Metric_plot Bar plot of Negative Predictive Value metric Probability_plot Density plot of predicted probabilities per subgroup. Only plotted if probabilities are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
npv_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
npv_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

pred_rate_parity Predictive Rate Parity

Description

This function computes the Predictive Rate Parity metric. Formula: TP / (TP + FP)

Usage

```
pred_rate_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.

prop_parity

outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5 .
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Predictive Rate Parity metric (also known as Sufficiency) as described by Zafar et al., 2017. Predictive rate parity is calculated by the division of true positives with all observations predicted positives. This metrics equals to what is traditionally known as precision or positive predictive value. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their precisions are lower or higher compared to the reference group. Lower precisions will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw precision metrics for all groups and metrics standardized for the base group
	(predictive rate parity metric). Lower values compared to the reference group
	mean lower precisions in the selected subgroups
Metric_plot	Bar plot of Predictive Rate Parity metric
Probability_plot	
	Density plot of predicted probabilities per subgroup. Only plotted if probabili

Density plot of predicted probabilities per subgroup. Only plotted if probabilities are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
pred_rate_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
pred_rate_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

prop_parity

Proportional parity

Description

This function computes the Proportional parity metric Formula: (TP + FP) / (TP + FP + TN + FN)

Usage

```
prop_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Proportional parity metric (also known as Impact Parity or Minimizing Disparate Impact) as described by Calders and Verwer 2010. Proportional parity is calculated based on the comparison of the proportion of all positively classified individuals in all subgroups of the data. In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their proportion of positively predicted observations are lower or higher compared to the reference group. Lower proportions will be reflected in numbers lower than 1 in the returned named vector.

Value

Metric

Raw proportions for all groups and metrics standardized for the base group (proportional parity metric). Lower values compared to the reference group mean lower proportion of positively predicted observations in the selected subgroups

roc_parity

 Metric_plot
 Bar plot of Proportional parity metric

 Probability_plot
 Density plot of predicted probabilities per subgroup. Only plotted if probabilities are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
prop_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
prop_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

roc_parity ROC AUC parity

Description

This function computes the ROC AUC parity metric

Usage

```
roc_parity(data, outcome, group, probs, base = NULL, group_breaks = NULL)
```

Arguments

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).
probs	Column name or vector with the predicted probabilities (numeric between 0 - 1).
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number $>= 2$ giving the number of intervals into which group feature is to be cut.

Details

This function computes the ROC AUC values for each subgroup. In the returned table, the reference group will be assigned 1, while all other groups will be assigned values according to whether their ROC AUC values are lower or higher compared to the reference group. Lower ROC AUC will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw ROC AUC metrics for all groups and metrics standardized for the base	
	group (parity metric). Lower values compared to the reference group mean	
	lower ROC AUC values in the selected subgroups	
Metric_plot	Bar plot of ROC AUC metric	
Probability_plot		
	Density plot of predicted probabilities per subgroup	
ROCAUC_plot	ROC plots for all subgroups	

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
roc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', base = 'Caucasian')
roc_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', base = 'African_American')</pre>
```

spec_parity

Specificity parity

Description

This function computes the Specificity parity metric Formula: TN / (TN + FP)

Usage

```
spec_parity(
   data,
   outcome,
   group,
   probs = NULL,
   preds = NULL,
   outcome_base = NULL,
   cutoff = 0.5,
   base = NULL,
   group_breaks = NULL
)
```

data	Data.frame that contains the necessary columns.
outcome	Column name indicating the binary outcome variable (character).
group	Column name indicating the sensitive group (character).

probs	Column name or vector with the predicted probabilities (numeric between 0 - 1). Either probs or preds need to be supplied.
preds	Column name or vector with the predicted binary outcome (0 or 1). Either probs or preds need to be supplied.
outcome_base	Base level of the outcome variable (i.e., negative class). Default is the first level of the outcome variable.
cutoff	Cutoff to generate predicted outcomes from predicted probabilities. Default set to 0.5.
base	Base level of the sensitive group (character).
group_breaks	If group is continuous (e.g., age): either a numeric vector of two or more unique cut points or a single number ≥ 2 giving the number of intervals into which group feature is to be cut.

Details

This function computes the Specificity parity metric. Specificities are calculated by the division of true negatives with all negatives (irrespective of predicted values). In the returned named vector, the reference group will be assigned 1, while all other groups will be assigned values according to whether their specificities are lower or higher compared to the reference group. Lower specificities will be reflected in numbers lower than 1 in the returned named vector, thus numbers lower than 1 mean WORSE prediction for the subgroup.

Value

Metric	Raw specificity metrics for all groups and metrics standardized for the base group (specificity parity metric). Lower values compared to the reference group mean lower specificities in the selected subgroups
Metric_plot	Bar plot of Specificity parity metric
Probability_p	lot
	Density plot of predicted probabilities per subgroup. Only plotted if probabili- ties are defined

Examples

```
data(compas)
compas$Two_yr_Recidivism_01 <- ifelse(compas$Two_yr_Recidivism == 'yes', 1, 0)
spec_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
probs = 'probability', cutoff = 0.4, base = 'Caucasian')
spec_parity(data = compas, outcome = 'Two_yr_Recidivism_01', group = 'ethnicity',
preds = 'predicted', cutoff = 0.5, base = 'Hispanic')</pre>
```

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