

Package ‘fairml’

April 15, 2021

Type Package

Title Fair Models in Machine Learning

Version 0.4

Date 2021-04-15

Depends R (>= 3.5.0)

Imports methods, optiSolve, glmnet

Suggests lattice

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Description Fair machine learning regression models which take sensitive attributes into account in model estimation. Currently implementing Komiyama et al. (2018) <<http://proceedings.mlr.press/v80/komiyama18a/komiyama18a.pdf>> and an improvement over the former that uses ridge regression to enforce fairness.

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LazyData yes

NeedsCompilation no

Repository CRAN

Date/Publication 2021-04-15 18:10:02 UTC

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fairml-package	<i>Fair models in machine learning</i>
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Description

Fair machine learning models: estimation, tuning and prediction.

Details

fairml implements key algorithms for learning machine learning models while enforcing fairness with respect to a set of observed sensitive (or protected) attributes.

Currently **fairml** implements the following algorithms (references below):

- `nclm()`: the non-convex formulation of fair linear regression model from Komiyama et al. (2018).
- `frfm()`: the fair ridge regression model.

Author(s)

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References

Komiyama J, Takeda A, Honda J, Shimao H (2018). "Nonconvex Optimization for Regression with Fairness Constraints". Proceedings of the 35th International Conference on Machine Learning (ICML), PMLR **80**:2737–2746.

communities.and.crime *iCommunities and Crime Data Set*

Description

Combined socio-economic data from the 1990 Census, law enforcement data from the 1990 LEMAS survey, and crime data from the 1995 FBI UCR for various communities in the United States.

Usage

```
data(communities.and.crime)
```

Format

The data contains 1969 observations and 104 variables. See the UCI Machine Learning Repository for details.

Note

The data set has been pre-processed as in Komiyama et al. (2018), with the following exceptions:

- the variable `community` has been dropped, as it is non-predictive and contains a sizeable number of missing values;
- the variables `LemasSwornFT`, `LemasSwFTPerPop`, `LemasSwFTFieldOps`, `LemasSwFTFieldPerPop`, `LemasTotalReq`, `LemasTotReqPerPop`, `PolicReqPerOffic`, `PolicPerPop`, `RacialMatchCommPol`, `PctPolicWhite`, `PctPolicBlack`, `PctPolicHisp`, `PctPolicAsian`, `PctPolicMinor`, `OfficAssgnDrugUnits`, `NumKindsDrugsSeiz`, `PolicAveOTWorked`, `PolicCars`, `PolicOperBudg`, `LemasPctPolicOnPatr`, `LemasGangUnitDeploy` and `PolicBudgPerPop` have been dropped because they have more than 80% missing values.

In that paper, `ViolentCrimesPerPop` is the response variable, `racepctblack` and `PctForeignBorn` are the sensitive attributes and the remaining variables are used as predictors.

References

UCI Machine Learning Repository:
<http://archive.ics.uci.edu/ml/datasets/communities+and+crime>

Examples

```
data(communities.and.crime)

# short-hand variable names.
cc = communities.and.crime[complete.cases(communities.and.crime), ]
r = cc[, "ViolentCrimesPerPop"]
s = cc[, c("racepctblack", "PctForeignBorn")]
p = cc[, setdiff(names(cc), c("ViolentCrimesPerPop", names(s)))]
```

```
m = nclm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)
```

```
m = frrm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)
```

compas

Criminal Offenders Screened in Florida

Description

A collection of criminal offenders screened in Florida (US) during 2013-14.

Usage

```
data(compas)
```

Format

The data contains 5855 observations and the following variables:

- age, a continuous variable containing the age (in years) of the person;
- juv_fel_count, a continuous variable containing the number of juvenile felonies;
- decile_score, a continuous variable, the decile of the COMPAS score;
- juv_misd_count, a continuous variable containing the number of juvenile misdemeanors;
- juv_other_count, a continuous variable containing the number of prior juvenile convictions that are not considered either felonies or misdemeanors;
- v_decile_score, a continuous variable containing the predicted decile of the COMPAS score;
- priors_count, a continuous variable containing the number of prior crimes committed;
- sex, a factor with levels "Female" and "Male";
- two_year_recid, a factor with two levels "Yes" and "No" (if the person has recidivated within two years);
- race, a factor encoding the race of the person;
- c_jail_in, a numeric variable containing the date in which the person entered jail (normalized between 0 and 1);
- c_jail_out, a numeric variable containing the date in which the person was released from jail (normalized between 0 and 1);
- c_offense_date, a numeric variable containing the date the offense was committed;
- screening_date, a numeric variable containing the date in which the person was screened (normalized between 0 and 1);
- in_custody, a numeric variable containing the date in which the person was placed in custody (normalized between 0 and 1);
- out_custody, a numeric variable containing the date in which the person was released from custody (normalized between 0 and 1);

Note

The data set has been pre-processed as in Komiyama et al. (2018), with the following exceptions:

- the race variable has not been reduced to a binary factor with levels "African-American" and "not African-American";
- the variables `type_of_assessment`, `v_type_of_assessment` have been dropped from the analysis because they take the same value for all observations;
- variables like `c_jail_in` and `c_jail_out` that encode dates have been jointly rescaled to preserve the temporal ordering of events.

In that paper, `two_year_recid` is the response variable, `sex` and `race` are the sensitive attributes and the remaining variables are used as predictors.

References

Angwin J, Larson J, Mattu S, Kirchner L (2016). "Machine Bias: Theres Software Used Around the Country to Predict Future Criminals." <https://www.propublica.org>.

Examples

```
data(compas)

# convert the response back to a numeric variable.
compas$two_year_recid = as.numeric(compas$two_year_recid) - 1

# short-hand variable names.
r = compas[, "two_year_recid"]
s = compas[, c("sex", "race")]
p = compas[, setdiff(names(compas), c("two_year_recid", "sex", "race"))]

m = nclm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)

m = frrm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)
```

Description

Cross-validation for the models in the **fairml** package.

Usage

```
fairml.cv(response, predictors, sensitive, method = "k-fold", ...,
          unfairness, model, model.args = list())
```

```
cv.loss(x)
cv.unfairness(x)
cv.folds(x)
```

Arguments

response	a numeric vector, the response variable.
predictors	a numeric matrix or a data frame containing numeric and factor columns; the predictors.
sensitive	a numeric matrix or a data frame containing numeric and factor columns; the sensitive attributes.
method	a character string, either k-fold, custom-folds or hold-out. See below for details.
...	additional arguments for the cross-validation method.
unfairness	a positive number in [0, 1], the proportion of the explained variance that can be attributed to the sensitive attributes.
model	a character string, the label of the model. Currently only "nclm" is available.
model.args	additional arguments passed to the model.
x	an object of class fair.kcv or fair.kcv.list.

Details

The following cross-validation methods are implemented:

- *k-fold*: the data are split in k subsets of equal size. For each subset in turn, model is fitted on the other $k - 1$ subsets and the loss function is then computed using that subset. Loss estimates for each of the k subsets are then combined to give an overall loss for data.
- *custom-folds*: the data are manually partitioned by the user into subsets, which are then used as in k -fold cross-validation. Subsets are not constrained to have the same size, and every observation must be assigned to one subset.
- *hold-out*: k subsamples of size m are sampled independently without replacement from the data. For each subsample, model is fitted on the remaining $m - \text{length}(\text{response})$ samples and the loss function is computed on the m observations in the subsample. The overall loss estimate is the average of the k loss estimates from the subsamples.

Cross-validation methods accept the following optional arguments:

- k : a positive integer number, the number of groups into which the data will be split (in k -fold cross-validation) or the number of times the data will be split in training and test samples (in hold-out cross-validation).
- m : a positive integer number, the size of the test set in hold-out cross-validation.

- runs: a positive integer number, the number of times k-fold or hold-out cross-validation will be run.
- folds: a list in which element corresponds to one fold and contains the indices for the observations that are included to that fold; or a list with an element for each run, in which each element is itself a list of the folds to be used for that run.

If cross-validation is used with multiple runs, the overall loss is the average of the loss estimates from the different runs.

The predictive performance of the models is measured using the mean square error as the loss function.

Value

`fairml.cv()` returns an object of class `fair.kcv.list` if `runs` is at least 2, an object of class `fair.kcv` if `runs` is equal to 1.

`cv.loss()` returns a numeric vectors containing the values of the loss function computed for each run of cross-validation.

`cv.unfairness()` returns a numeric vectors containing the values of the unfairness criterion computed on the validation folds for each run of cross-validation.

`cv.folds()` returns a list containing the indexes of the observations in each of the cross-validation folds. In the case of k-fold cross-validation, if `runs` is larger than 1, each element of the list is itself a list with the indexes for the observations in each fold in each run.

Author(s)

Marco Scutari

Examples

```
kcv = fairml.cv(response = vur.test$y, predictors = vur.test$X,
               sensitive = vur.test$S, unfairness = 0.10, model = "nclm",
               method = "k-fold", k = 10, runs = 10)
kcv
cv.loss(kcv)
cv.unfairness(kcv)

# run a second cross-validation with the same folds.
fairml.cv(response = vur.test$y, predictors = vur.test$X,
          sensitive = vur.test$S, unfairness = 0.10, model = "nclm",
          method = "custom-folds", folds = cv.folds(kcv))
```

fairness.profile.plot *Profile Fair Models with Respect to Tuning Parameters*

Description

Visually explore various aspect of a model over the range of possible values of the tuning parameters that control its fairness.

Usage

```
fairness.profile.plot(response, predictors, sensitive, unfairness,
  legend = FALSE, type = "coefficients", model = "nclm", model.args = list())
```

Arguments

response	a numeric vector, the response variable.
predictors	a numeric matrix or a data frame containing numeric and factor columns; the predictors.
sensitive	a numeric matrix or a data frame containing numeric and factor columns; the sensitive attributes.
unfairness	a vector of positive numbers in [0, 1], the proportion of the explained variance that can be attributed to the sensitive attributes. The default value is seq(from = 0.00, to = 1, by = 0.02).
legend	a logical value, whether to add a legend to the plot.
type	a character string, either "coefficients" (the default) or "variance".
model	a character string, the label of the model. Currently only "nclm" is available.
model.args	additional arguments passed to the model.

Details

fairness.profile.plot() fits the model for all the values of the argument unfairness, and produces a profile plot of the regression coefficients or the proportion of explained variance.

If type = "coefficients", the coefficients of the model are plotted against the values of unfairness.

If type = "variance", the following quantities are plotted against the values of unfairness:

1. the proportion of variance explained by the sensitive attributes (with respect to the response;
2. the proportion of variance explained by the predictors (with respect to the response;
3. the proportion of variance explained by the sensitive attributes (with respect to the combined sensitive attributes and predictors).

Value

A trellis object containing a **lattice** plot.

Author(s)

Marco Scutari

Examples

```
data(vur.test)
fairness.profile.plot(response = vur.test$y, predictors = vur.test$X,
  sensitive = vur.test$S, type = "coefficients", legend = TRUE)
fairness.profile.plot(response = vur.test$y, predictors = vur.test$X,
  sensitive = vur.test$S, type = "variance", legend = TRUE)
```

frmm

Fair Ridge Regression Model

Description

A regression model enforcing fairness with a ridge penalty.

Usage

```
frmm(response, predictors, sensitive, unfairness)
```

Arguments

response	a numeric vector, the response variable.
predictors	a numeric matrix or a data frame containing numeric and factor columns; the predictors.
sensitive	a numeric matrix or a data frame containing numeric and factor columns; the sensitive attributes.
unfairness	a positive number in $[0, 1]$, the proportion of the explained variance that can be attributed to the sensitive attributes.

Details

frmm() defines fairness as statistical parity.

The algorithm works like this:

1. regresses the predictors against the sensitive attributes;
2. constructs a new set of predictors that are decorrelated from the sensitive attributes using the residuals of this regression;
3. regresses the response against the decorrelated predictors and the sensitive attributes, while
4. using a ridge penalty to control the proportion of variance the sensitive attributes can explain with respect to the overall explained variance of the model.

Both sensitive and predictors are standardized internally before estimating the regression coefficients, which are then rescaled back to match the original scales of the variables.

Value

`frrm()` returns an object of class `c("frrm", "fair.model")`.

Author(s)

Marco Scutari

See Also

[nclm](#)

german.credit

German Credit Data

Description

A credit scoring data set that can be used to predict defaults on consumer loans in the German market.

Usage

```
data(german.credit)
```

Format

The data contains 1000 observations (700 good loans, 300 bad loans) and the following variables:

- `Account_status`: a factor with four levels representing the amount of money in the account or "no chcking account".
- `Duration`: a continuous variable, the duration in months.
- `Credit_history`: a factor with five levels representing possible credit history backgrounds.
- `Purpose`: a factor with ten levels representing possible reasons for taking out a loan.
- `Credit_amount`: a continuous variable.
- `Savings_bonds`: a factor with five levels representing amount of money available in savings and bonds or "unknown / no savings account".
- `Present_employment_since`: a factor with five levels representing the length of tenure in the current employment or "unemployed".
- `Installment_rate`: a continuous variable, the installment rate in percentage of disposable income.
- `Other_debtors_guarantors`: a factor with levels "none", "co-applicant" and "guarantor".
- `Resident_since`: a continuous variable, number of years in the current residence.
- `Property`: a factor with four levels describing the type of property to be bought or "unknown / no property".
- `Age`: a continuous variable, the age in years.

- Other_installment_plans: a factor with levels "bank", "none" and "stores".
- Housing: a factor with levels "rent", "own" and "for free".
- Existing_credits: a continuous variable, the number of existing credit lines at this bank.
- Job: a factor with four levels for different job descriptions.
- People_maintenance_for: a continuous variable, the number of people being liable to provide maintenance for.
- Telephone: a factor with levels "none" and "yes".
- Foreign_worker: a factor with levels "no" and "yes".
- Credit_risk: a factor with levels "BAD" and "GOOD".
- Gender: a factor with levels "Male" and "Female".

Note

The variable "Personal status and sex" in the original data has been transformed into Gender by dropping the personal status information.

References

UCI Machine Learning Repository:
[https://archive.ics.uci.edu/ml/datasets/statlog+\(german+credit+data\)](https://archive.ics.uci.edu/ml/datasets/statlog+(german+credit+data))

law.school.admissions *Law School Admission Council data*

Description

Survey among students attending law school in the U.S. in 1991.

Usage

```
data(law.school.admissions)
```

Format

The data contains 20800 observations and the following variables:

- age, a continuous variable containing the student's age in years;
- decile1, a continuous variable containing the student's decile in the school given his grades in Year 1;
- decile3, a continuous variable containing the student's decile in the school given his grades in Year 3;
- fam_inc, a continuous variable containing student's family income bracket (from 1 to 5);
- lsat, a continuous variable containing the student's LSAT score;
- ugpa, a continuous variable containing the student's undergraduate GPA;

- gender, a factor with levels "female" and "male";
- race1, a factor with levels "asian", "black", "hisp", "other" and "white";
- cluster, a factor with levels "1", "2", "3", "4", "5" and "6" encoding the tiers of law school prestige;
- fulltime, a factor with levels "FALSE" and "TRUE", whether the student will work full-time or part-time;
- bar, a factor with levels "FALSE" and "TRUE", whether the student passed the bar exam on the first try.

Note

The data set has been pre-processed as in Komiyama et al. (2018), with the following exceptions:

- DOB_yr, the year of birth, has been dropped because it is (nearly) perfectly collinear with age, and thus it is redundant;
- decile1b has been dropped because it is (nearly) perfectly collinear with decile1, and thus it is redundant.

In that paper, ugpa is the response variable, age and race1 are the sensitive attributes and the remaining variables are used as predictors.

References

Sander RH (2004). "A Systemic Analysis of Affirmative Action in American Law Schools". Stanford Law Review, 57:367–483.

Examples

```
data(law.school.admissions)

# short-hand variable names.
ll = law.school.admissions
r = ll[, "ugpa"]
s = ll[, c("age", "race1")]
p = ll[, setdiff(names(ll), c("ugpa", "age", "race1"))]

m = nclm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)

m = frrm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)
```

methods for fair.model objects

Extract information from fair.model objects

Description

Extract various quantities of interest from an object of class `fair.model`.

Usage

```
# methods for all fair.model objects.
## S3 method for class 'fair.model'
coef(object, ...)
## S3 method for class 'fair.model'
residuals(object, ...)
## S3 method for class 'fair.model'
fitted(object, ...)
## S3 method for class 'fair.model'
sigma(object, ...)
## S3 method for class 'fair.model'
nobs(object, ...)
## S3 method for class 'fair.model'
print(x, digits, ...)
## S3 method for class 'fair.model'
summary(object, ...)
## S3 method for class 'fair.model'
all.equal(target, current, ...)
## S3 method for class 'fair.model'
plot(x, diagonal = FALSE, regression = FALSE, ...)

# methods for nclm objects.
## S3 method for class 'nclm'
predict(object, new.predictors, new.sensitive, ...)
## S3 method for class 'nclm'
summary(object, ...)
## S3 method for class 'nclm'
deviance(object, ...)
```

Arguments

`object, x, target, current` an object of class `fair.model` or `nclm`.

`digits` a non-negative integer, the number of significant digits.

`new.predictors` a numeric matrix or a data frame containing numeric and factor columns; the predictors for the new observations.

<code>new.sensitive</code>	a numeric matrix or a data frame containing numeric and factor columns; the sensitive attributes for the new observations.
<code>diagonal</code>	a logical value, whether to draw the diagonal of the first quadrant in <code>plot()</code> .
<code>regression</code>	a logical value, whether to draw the regression line of the observed values on the fitted values from the model in <code>plot()</code> .
<code>...</code>	additional arguments, currently ignored.

`national.longitudinal.survey`

Income and Labour Market Activities

Description

Survey results from the U.S. Bureau of Labor Statistics to gather information on the labour market activities and other life events of several groups.

Usage

```
data(national.longitudinal.survey)
```

Format

The data contains 4908 observations and the following variables:

- `age`, a numeric variable containing the interviewee's age in years;
- `race`, a factor with 20 levels denoting various racial/ethnic origins;
- `gender`, a factor with levels "Male" and "Female".
- `grade90`, a factor containing the highest completed school grade from "3RD GRADE" to "8TH YR COL OR MORE", with 18 levels;
- `income06`, a numeric variable, income in 2006 in 10000-USD units;
- `income96`, a numeric variable, income in 1996 in 10000-USD units;
- `income90`, a numeric variable, income in 1990 in 10000-USD units;
- `partner`, a factor encoding whether the interviewee has a partner, with levels "No" and "Yes";
- `height`, a numeric variable, the height of the interviewee;
- `weight`, a numeric variable, the weight of the interviewee;
- `famsize`, a numeric variable, the number of family members;
- `genhealth`, a factor with levels "Excellent", "Very Good", "Good", "Fair", "Poor" encoding the general health status of the interviewee;
- `illegalact`, a numeric variable containing the number of illegal acts committed by the interviewee;
- `charged`, a numeric variable containing the number of illegal acts for which the interviewee has been charged;

- jobsnum90, a numeric value, the number of different jobs ever reported;
- afqt89, a numeric value, the percentile score of the "Profiles, Armed Forces Qualification Test" (AFQT);
- typejob90, a factor with 13 levels encoding different job types;
- jobtrain90, a factor with levels "No" and "Yes" encoding whether the job was classified as training.

Note

The data set has been pre-processed differently from Komiyama et al. (2018). In particular:

- the variables income96 and income06 have been retained as alternative responses;
- the variables height, weight, race, partner and famsize have been retained;
- the variables grade90 and genhealth are coded as ordered factors because they do not make sense on a numeric scale.

In that paper, income90 is the response variable, gender and age are the sensitive attributes.

References

U.S. Bureau of Labor Statistics: <https://www.bls.gov/nls/>

Examples

```
data(national.longitudinal.survey)

# short-hand variable names.
nn = national.longitudinal.survey
# remove alternative response variables.
nn = nn[, setdiff(names(nn), c("income96", "income06"))]
# short-hand variable names.
r = nn[, "income90"]
s = nn[, c("gender", "age")]
p = nn[, setdiff(names(nn), c("income90", "gender", "age"))]

m = nclm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)

m = frrm(response = r, sensitive = s, predictors = p, unfairness = 0.05)
summary(m)
```

Description

Fair regression model based on nonconvex optimization from Komiyama et al. (2018).

Usage

```
nclm(response, predictors, sensitive, unfairness, covfun, lambda)
```

Arguments

response	a numeric vector, the response variable.
predictors	a numeric matrix or a data frame containing numeric and factor columns; the predictors.
sensitive	a numeric matrix or a data frame containing numeric and factor columns; the sensitive attributes.
unfairness	a positive number in $[0, 1]$, the proportion of the explained variance that can be attributed to the sensitive attributes.
covfun	a function computing covariance matrices. It defaults to the <code>cov()</code> function from the stats package.
lambda	a non-negative number, a ridge-regression penalty coefficient. It defaults to zero.

Details

`nclm()` defines fairness as statistical parity.

The algorithm proposed by Komiyama et al. (2018) works like this:

1. regresses the predictors against the sensitive attributes;
2. constructs a new set of predictors that are decorrelated from the sensitive attributes using the residuals of this regression;
3. regresses the response against the decorrelated predictors and the sensitive attributes, while
4. bounding the proportion of variance the sensitive attributes can explain with respect to the overall explained variance of the model.

Both `sensitive` and `predictors` are standardized internally before estimating the regression coefficients, which are then rescaled back to match the original scales of the variables. `response` is only standardized if it has a variance smaller than 1, as that seems to improve the stability of the solutions provided by the optimizer (as far as the data included in **fairml** are concerned).

The `covfun` argument makes it possible to specify a custom function to compute the covariance matrices used in the constrained optimization. Some examples are the kernel estimators described in Komiyama et al. (2018) and the shrinkage estimators in the **corpcor** package.

Value

`nclm()` returns an object of class `c("nclm", "fair.model")`.

Author(s)

Marco Scutari

References

Komiyama J, Takeda A, Honda J, Shimao H (2018). "Nonconvex Optimization for Regression with Fairness Constraints". *Proceedings of the 35th International Conference on Machine Learning (ICML)*, PMLR **80**:2737–2746.

See Also

[frm](#)

vur.test

Synthetic data set to test fair regression models

Description

This is a synthetic data set used as a test case in the **fairml** package.

Usage

```
data(vur.test)
```

Format

The `vur.test` data set is a list with following three elements:

- `y`, the response variable;
- `X`, a numeric matrix containing 3 predictors called `X1`, `X2` and `X3`;
- `S`, a numeric matrix containing 3 sensitive attributes called `S1`, `S2` and `S3`.

Note

This data set is called `vur.test` because it is generated from a very *unfair* regression model in which sensitive attributes explain the lion's share of the overall explained variance. The code used to generate the data is as follows.

```
library(mvtnorm)
sigma = matrix(0.3, nrow = 6, ncol = 6)
diag(sigma) = 1
n = 1000
X = rmvnorm(n, mean = rep(0, 6), sigma = sigma)
```

```
y = 2 + 2 * X[, 1] + 3 * X[, 2] + 4 * X[, 3] + 5 * X[, 4] +  
      6 * X[, 5] + 7 * X[, 6] + rnorm(n, sd = 10)  
S = X[, 4:6]  
X = X[, 1:3]  
colnames(X) = c("X1", "X2", "X3")  
colnames(S) = c("S1", "S2", "S3")  
vur.test = list(y = y, X = X, S = S)
```

Examples

```
data(vur.test)  
sensitive.attributes.model = lm(y ~ S, data = vur.test)  
summary(sensitive.attributes.model)$r.squared  
overall.model = lm(y ~ X + S, data = vur.test)  
summary(overall.model)$r.squared
```

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