

Package ‘deepregression’

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Title Fitting Deep Distributional Regression

Version 0.1

Description Allows for the specification of semi-structured deep distributional regression models which are fitted in a neural network as proposed by Ruegamer et al. (2021) <[arXiv:2104.02705](https://arxiv.org/abs/2104.02705)>. Predictors can be modeled using structured (penalized) linear effects, structured non-linear effects or using an unstructured deep network model.

Config/reticulate list(packages = list(list(package = ``six'', pip = TRUE), list(package = ``tensorflow'', version = ``2.5.0rc0'', pip = TRUE), list(package = ``tensorflow_probability'', version = ``0.12'', pip = TRUE), list(package = ``keras'', version = ``2.5.0rc0'', pip = TRUE)))

Depends R (>= 4.0.0)

Suggests testthat, knitr

Imports tensorflow (>= 2.2.0), tfprobability, keras, mgcv, dplyr, purrr, R6, reticulate (>= 1.14), Matrix, magrittr, Metrics, tfruns, methods, utils

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check_and_install	<i>Function to check python environment and install necessary packages</i>
-------------------	--

Description

Note: The package currently relies on tensorflow version 2.0.0 which is not available for the latest python versions 3.9 and later. If you encounter problems with installing the required python modules please make sure, that a correct python version is configured using ‘py_discover_config’ and change the python version if required. Internally uses keras::install_keras.

Usage

```
check_and_install(force = FALSE)
```

Arguments

force	if TRUE, forces the installations
-------	-----------------------------------

Value

Function that checks if a Python environment is available and contains TensorFlow. If not the recommended version is installed.

create_family	<i>Function to create (custom) family</i>
---------------	---

Description

Function to create (custom) family

Usage

```
create_family(tfd_dist, trafo_list, output_dim = 1L)
```

Arguments

tfd_dist	a tensorflow probability distribution
trafo_list	list of transformations h for each parameter (e.g, exp for a variance parameter)
output_dim	integer defining the size of the response

Value

a function that can be used by `tfp$layers$DistributionLambda` to create a new distributional layer

`cv`*Generic cv function***Description**

Generic cv function

Usage

```
cv(x, ...)
```

Arguments

<code>x</code>	model to do cv on
<code>...</code>	further arguments passed to the class-specific function

`deepregression`*Fitting Semi-Structured Deep Distributional Regression***Description**

Fitting Semi-Structured Deep Distributional Regression

Usage

```
deepregression(
  y,
  list_of_formulas,
  list_of_deep_models = NULL,
  family = "normal",
  data,
  tf_seed = as.integer(1991 - 5 - 4),
  return_prepoc = FALSE,
  subnetwork_builder = subnetwork_init,
  model_builder = keras_dr,
  fitting_function = utils::getFromNamespace("fit.keras.engine.training.Model",
    "keras"),
  additional_processors = list(),
  penalty_options = penalty_control(),
  orthog_options = orthog_control(),
  verbose = FALSE,
  ...
)
```

Arguments

<code>y</code>	response variable
<code>list_of_formulas</code>	a named list of right hand side formulas, one for each parameter of the distribution specified in <code>family</code> ; set to ~ 1 if the parameter should be treated as constant. Use the <code>s()</code> -notation from <code>mgcv</code> for specification of non-linear structured effects and <code>d(...)</code> for deep learning predictors (predictors in brackets are separated by commas), where <code>d</code> can be replaced by an name name of the names in <code>list_of_deep_models</code> , e.g., $\sim 1 + s(x) + \text{my_deep_mod}(a, b, c)$, where <code>my_deep_mod</code> is the name of the neural net specified in <code>list_of_deep_models</code> and <code>a, b, c</code> are features modeled via this network.
<code>list_of_deep_models</code>	a named list of functions specifying a keras model. See the examples for more details.
<code>family</code>	a character specifying the distribution. For information on possible distribution and parameters, see make_tfd_dist . Can also be a custom distribution.
<code>data</code>	data.frame or named list with input features
<code>tf_seed</code>	a seed for TensorFlow (only works with R version $\geq 2.2.0$)
<code>return_prepoc</code>	logical; if TRUE only the pre-processed data and layers are returned (default FALSE).
<code>subnetwork_builder</code>	function to build each subnetwork (network for each distribution parameter; per default <code>subnetwork_init</code>). Can also be a list of the same size as <code>list_of_formulas</code> .
<code>model_builder</code>	function to build the model based on additive predictors (per default <code>keras_dr</code>). In order to work with the methods defined for the class <code>deepregression</code> , the model should behave like a keras model
<code>fitting_function</code>	function to fit the instantiated model when calling <code>fit</code> . Per default the keras <code>fit</code> function.
<code>additional_processors</code>	a named list with additional processors to convert the formula(s). Can have an attribute "controls" to pass additional controls
<code>penalty_options</code>	options for smoothing and penalty terms defined by penalty_control
<code>orthog_options</code>	options for the orthogonalization defined by orthog_control
<code>verbose</code>	logical; whether to print progress of model initialization to console
<code>...</code>	further arguments passed to the <code>model_builder</code> function

References

Ruegamer, D. et al. (2021): deepregression: a Flexible Neural Network Framework for Semi-Structured Deep Distributional Regression. <https://arxiv.org/abs/2104.02705>.

Examples

```

library(deepregression)

n <- 1000
data = data.frame(matrix(rnorm(4*n), nrow=n, ncol=4))
colnames(data) <- c("x1", "x2", "x3", "xa")
formula <- ~ 1 + deep_model(x1,x2,x3) + s(xa) + x1

deep_model <- function(x) x %>%
  layer_dense(units = 32, activation = "relu", use_bias = FALSE) %>%
  layer_dropout(rate = 0.2) %>%
  layer_dense(units = 8, activation = "relu") %>%
  layer_dense(units = 1, activation = "linear")

y <- rnorm(n) + data$xa^2 + data$x1

mod <- deepregression(
  list_of_formulas = list(loc = formula, scale = ~ 1),
  data = data, y = y,
  list_of_deep_models = list(deep_model = deep_model)
)

if(!is.null(mod)){

  # train for more than 10 epochs to get a better model
  mod %>% fit(epochs = 10, early_stopping = TRUE)
  mod %>% fitted() %>% head()
  cvres <- mod %>% cv()
  mod %>% get_partial_effect(name = "s(xa)")
  mod %>% coef()
  mod %>% plot()

}

mod <- deepregression(
  list_of_formulas = list(loc = ~ 1 + s(xa) + x1, scale = ~ 1,
    dummy = ~ -1 + deep_model(x1,x2,x3) %OZ% 1),
  data = data, y = y,
  list_of_deep_models = list(deep_model = deep_model),
  mapping = list(1,2,1:2)
)

```

Description

Function to define output distribution based on dist_fun

Usage

```
distfun_to_dist(dist_fun, preds)
```

Arguments

dist_fun	a distribution function as defined by <code>make_tfd_dist</code>
preds	tensors with predictions

Value

a symbolic tfp distribution

extractval	<i>Extract value in term name</i>
------------	-----------------------------------

Description

Extract value in term name

Usage

```
extractval(term, name, null_for_missing = FALSE)
```

Arguments

term	character representing a formula term
name	character; the value to extract
null_for_missing	logical; if TRUE, returns NULL if argument is missing

Value

the value used for name

Examples

```
extractval("s(a, la = 2)", "la")
```

family_to_tfd *Character-tfd mapping function*

Description

Character-tfd mapping function

Usage

```
family_to_tfd(family)
```

Arguments

family	character defining the distribution
--------	-------------------------------------

Value

a tfp distribution

family_to_trafo *Character-to-transformation mapping function*

Description

Character-to-transformation mapping function

Usage

```
family_to_trafo(family, add_const = 1e-08)
```

Arguments

family	character defining the distribution
add_const	see make_tfd_dist

Value

a list of transformation for each distribution parameter

fit	<i>Generic train function</i>
-----	-------------------------------

Description

Generic train function

Usage

```
fit(object, ...)
```

Arguments

object	object to apply fit on
...	further arguments passed to the class-specific function

from_dist_to_loss	<i>Function to transform a distribution layer output into a loss function</i>
-------------------	---

Description

Function to transform a distribution layer output into a loss function

Usage

```
from_dist_to_loss(  
  family,  
  ind_fun = function(x) tfd_independent(x),  
  weights = NULL  
)
```

Arguments

family	see ?deepregression
ind_fun	function applied to the model output before calculating the log-likelihood. Per default independence is assumed by applying tfd_independent.
weights	sample weights

Value

loss function

from_preds_to_dist *Define Predictor of a Deep Distributional Regression Model*

Description

Define Predictor of a Deep Distributional Regression Model

Usage

```
from_preds_to_dist(
  list_pred_param,
  family = NULL,
  output_dim = 1L,
  mapping = NULL,
  from_distfun_to_dist = distfun_to_dist,
  add_layer_shared_pred = function(x, units) layer_dense(x, units = units, use_bias =
    FALSE)
)
```

Arguments

list_pred_param	list of input-output(-lists) generated from <code>subnetwork_init</code>
family	see <code>?deepregression</code> ; if <code>NULL</code> , concatenated <code>list_pred_param</code> entries are returned (after applying <code>mapping</code> if provided)
output_dim	dimension of the output
mapping	a list of integers. The i-th list item defines which element elements of <code>list_pred_param</code> are used for the i-th parameter. For example, <code>map = list(1, 2, 1:2)</code> means that <code>list_pred_param[[1]]</code> is used for the first distribution parameter, <code>list_pred_param[[2]]</code> for the second distribution parameter and <code>list_pred_param[[3]]</code> for both distribution parameters (and then added once to <code>list_pred_param[[1]]</code> and once to <code>list_pred_param[[2]]</code>)
from_distfun_to_dist	function creating a tfp distribution based on the prediction tensors and <code>dist_fun</code> . See <code>?distfun_to_dist</code>
add_layer_shared_pred	layer to extend shared layers defined in <code>mapping</code>

Value

a list with input tensors and output tensors that can be passed to, e.g., `keras_model`

get_distribution	<i>Function to return the fitted distribution</i>
------------------	---

Description

Function to return the fitted distribution

Usage

```
get_distribution(x, data = NULL, force_float = FALSE)
```

Arguments

x	the fitted deepregression object
data	an optional data set
force_float	forces conversion into float tensors

get_partial_effect	<i>Return partial effect of one smooth term</i>
--------------------	---

Description

Return partial effect of one smooth term

Usage

```
get_partial_effect(  
  object,  
  name,  
  return_matrix = FALSE,  
  which_param = 1,  
  newdata = NULL  
)
```

Arguments

object	deepregression object
name	string; for partial match with smooth term
return_matrix	logical; whether to return the design matrix or
which_param	integer; which distribution parameter the partial effect (FALSE, default)
newdata	data.frame; new data (optional)

`get_type_pfc`*Function to subset parsed formulas***Description**

Function to subset parsed formulas

Usage

```
get_type_pfc(pfc, type = NULL)
```

Arguments

<code>pfc</code>	list of parsed formulas
<code>type</code>	either <code>NULL</code> (all types of coefficients are returned), "linear" for linear coefficients or "smooth" for coefficients of

`get_weight_by_name`*Function to retrieve the weights of a structured layer***Description**

Function to retrieve the weights of a structured layer

Usage

```
get_weight_by_name(mod, name, param_nr = 1)
```

Arguments

<code>mod</code>	fitted deepregression object
<code>name</code>	name of partial effect
<code>param_nr</code>	distribution parameter number

Value

weight matrix

handle_gam_term	<i>Function to define smoothness and call mgcv's smooth constructor</i>
-----------------	---

Description

Function to define smoothness and call mgcv's smooth constructor

Usage

```
handle_gam_term(object, data, controls)
```

Arguments

object	character defining the model term
data	data.frame or list
controls	controls for penalization

Value

constructed smooth term

keras_dr	<i>Compile a Deep Distributional Regression Model</i>
----------	---

Description

Compile a Deep Distributional Regression Model

Usage

```
keras_dr(  
  list_pred_param,  
  weights = NULL,  
  optimizer = tf$keras$optimizers$Adam(),  
  model_fun = keras_model,  
  monitor_metrics = list(),  
  from_preds_to_output = from_preds_to_dist,  
  loss = from_dist_to_loss(family = list(...)$family, weights = weights),  
  additional_penalty = NULL,  
  ...  
)
```

Arguments

<code>list_pred_param</code>	list of input-output(-lists) generated from <code>subnetwork_init</code>
<code>weights</code>	vector of positive values; optional (default = 1 for all observations)
<code>optimizer</code>	optimizer used. Per default Adam
<code>model_fun</code>	which function to use for model building (default <code>keras_model</code>)
<code>monitor_metrics</code>	Further metrics to monitor
<code>from_preds_to_output</code>	function taking the <code>list_pred_param</code> outputs and transforms it into a single network output
<code>loss</code>	the model's loss function; per default evaluated based on the arguments <code>family</code> and <code>weights</code> using <code>from_dist_to_loss</code>
<code>additional_penalty</code>	a penalty that is added to the negative log-likelihood; must be a function of <code>model\$trainable_weights</code> with suitable subsetting
<code>...</code>	arguments passed to <code>from_preds_to_output</code>

Value

a list with input tensors and output tensors that can be passed to, e.g., `keras_model`

`layer_add_identity` *Convenience layer function*

Description

Convenience layer function

Usage

```
layer_add_identity(inputs)
layer_concatenate_identity(inputs)
```

Arguments

<code>inputs</code>	list of tensors
---------------------	-----------------

Details

convenience layers to work with list of inputs where `inputs` can also have length one

Value

`tensor`

log_score	<i>Function to return the log_score</i>
-----------	---

Description

Function to return the log_score

Usage

```
log_score(
  x,
  data = NULL,
  this_y = NULL,
  ind_fun = function(x) tfd_independent(x, 1),
  convert_fun = as.matrix,
  summary_fun = function(x) x
)
```

Arguments

x	the fitted deepregression object
data	an optional data set
this_y	new y for optional data
ind_fun	function indicating the dependency; per default (iid assumption) <code>tfd_independent</code> is used.
convert_fun	function that converts Tensor; per default <code>as.matrix</code>
summary_fun	function summarizing the output; per default the identity

loop_through_pfc_and_call_trafo
<i>Function to loop through parsed formulas and apply data trafo</i>

Description

Function to loop through parsed formulas and apply data trafo

Usage

```
loop_through_pfc_and_call_trafo(pfc, newdata = NULL)
```

Arguments

pfc	list of processor transformed formulas
newdata	list in the same format as the original data

Value

list of matrices or arrays

`makeInputs`

Convenience layer function

Description

Convenience layer function

Usage

```
makeInputs(pp, param_nr)
```

Arguments

<code>pp</code>	processed predictors
<code>param_nr</code>	integer for the parameter

Value

input tensors with appropriate names

`make_folds`

Generate folds for CV out of one hot encoded matrix

Description

Generate folds for CV out of one hot encoded matrix

Usage

```
make_folds(mat, val_train = 0, val_test = 1)
```

Arguments

<code>mat</code>	matrix with columns corresponding to folds and entries corresponding to a one hot encoding
<code>val_train</code>	the value corresponding to train, per default 0
<code>val_test</code>	the value corresponding to test, per default 1

Details

`val_train` and `val_test` can both be a set of value

make_generator *creates a generator for training*

Description

creates a generator for training

Usage

```
make_generator(  
    input_x,  
    input_y = NULL,  
    batch_size,  
    sizes,  
    shuffle = TRUE,  
    seed = 42L  
)
```

Arguments

input_x	list of matrices
input_y	list of matrix
batch_size	integer
sizes	sizes of the image including colour channel
shuffle	logical for shuffling data
seed	seed for shuffling in generators

Value

generator for all x and y

make_generator_from_matrix

Make a DataGenerator from a data.frame or matrix

Description

Creates a Python Class that internally iterates over the data.

Usage

```
make_generator_from_matrix(
  x,
  y = NULL,
  generator = image_data_generator(),
  batch_size = 32L,
  shuffle = TRUE,
  seed = 1L
)
```

Arguments

x	matrix;
y	vector;
generator	generator as e.g. obtained from ‘keras::image_data_generator’. Used for consistent train-test splits.
batch_size	integer
shuffle	logical; Should data be shuffled?
seed	integer; seed for shuffling data.

Description

Families for deepregression

Usage

```
make_tfd_dist(family, add_const = 1e-08, output_dim = 1L, trafo_list = NULL)
```

Arguments

family	character vector
add_const	small positive constant to stabilize calculations
output_dim	number of output dimensions of the response (larger 1 for multivariate case)
trafo_list	list of transformations for each distribution parameter. Per default the transformation listed in details is applied.

Details

To specify a custom distribution, define the a function as follows `function(x) do.call(your_tfd_dist, lapply(1:ncol(your_trafo_list_on_inputs[[i]](x[, i, drop=FALSE])))` and pass it to `deepregression` via the `dist_fun` argument. Currently the following distributions are supported with parameters (and corresponding inverse link function in brackets):

- "normal": normal distribution with location (identity), scale (exp)
- "bernoulli": bernoulli distribution with logits (identity)
- "bernoulli_prob": bernoulli distribution with probabilities (sigmoid)
- "beta": beta with concentration 1 = alpha (exp) and concentration 0 = beta (exp)
- "betar": beta with mean (sigmoid) and scale (sigmoid)
- "cauchy": location (identity), scale (exp)
- "chi2": cauchy with df (exp)
- "chi": cauchy with df (exp)
- "exponential": exponential with lambda (exp)
- "gamma": gamma with concentration (exp) and rate (exp)
- "gamma": gamma with location (exp) and scale (exp)
- "gumbel": gumbel with location (identity), scale (exp)
- "half_cauchy": half cauchy with location (identity), scale (exp)
- "half_normal": half normal with scale (exp)
- "horseshoe": horseshoe with scale (exp)
- "inverse_gamma": inverse gamma with concentration (exp) and rate (exp)
- "inverse_gamma_ls": inverse gamma with location (exp) and variance (1/exp)
- "inverse_gaussian": inverse Gaussian with location (exp) and concentration (exp)
- "laplace": Laplace with location (identity) and scale (exp)
- "log_normal": Log-normal with location (identity) and scale (exp) of underlying normal distribution
- "logistic": logistic with location (identity) and scale (exp)
- "negbinom": neg. binomial with count (exp) and prob (sigmoid)
- "negbinom_ls": neg. binomial with mean (exp) and clutter factor (exp)
- "pareto": Pareto with concentration (exp) and scale (1/exp)
- "pareto_ls": Pareto location scale version with mean (exp) and scale (exp), which corresponds to a Pareto distribution with parameters scale = mean and concentration = 1/sigma, where sigma is the scale in the pareto_ls version.
- "poisson": poisson with rate (exp)
- "poisson_lograte": poisson with lograte (identity))
- "student_t": Student's t with df (exp)
- "student_t_ls": Student's t with df (exp), location (identity) and scale (exp)
- "uniform": uniform with upper and lower (both identity)
- "zinb": Zero-inflated negative binomial with mean (exp), variance (exp) and prob (sigmoid)
- "zip": Zero-inflated poisson distribution with mean (exp) and prob (sigmoid)

names_families	<i>Returns the parameter names for a given family</i>
----------------	---

Description

Returns the parameter names for a given family

Usage

```
names_families(family)
```

Arguments

family	character specifying the family as defined by deepregression
--------	--

Value

vector of parameter names

orthog_control	<i>Options for orthogonalization</i>
----------------	--------------------------------------

Description

Options for orthogonalization

Usage

```
orthog_control(
  split_fun = split_model,
  orthog_type = c("tf", "manual"),
  orthogonalize = options()$orthogonalize,
  identify_intercept = options()$identify_intercept,
  deep_top = NULL
)
```

Arguments

split_fun	a function separating the deep neural network in two parts so that the orthogonalization can be applied to the first part before applying the second network part; per default, the function <code>split_model</code> is used which assumes a dense layer as penultimate layer and separates the network into a first part without this last layer and a second part only consisting of a single dense layer that is fed into the output layer
-----------	--

orthog_type	one of two options; If "manual", the QR decomposition is calculated before model fitting, otherwise ("tf") a QR is calculated in each batch iteration via TF. The first only works well for larger batch sizes or ideally batch_size == NROW(y).
orthogonalize	logical; if set to TRUE, automatic orthogonalization is activated
identify_intercept	whether to orthogonalize the deep network w.r.t. the intercept to make the intercept identifiable
deep_top	function; optional function to put on top of the deep network instead of splitting the function using split_fun

Value

Returns a list with options

penalty_control	<i>Options for penalty setup in the pre-processing</i>
-----------------	--

Description

Options for penalty setup in the pre-processing

Usage

```
penalty_control(
  defaultSmoothing = NULL,
  df = 10,
  null_space_penalty = FALSE,
  absorb_cons = FALSE,
  anisotropic = TRUE,
  zero_constraint_for_smooths = TRUE,
  hat1 = FALSE,
  sp_scale = function(x) 1/NROW(x)
)
```

Arguments

defaultSmoothing	function applied to all s-terms, per default (NULL) the minimum df of all possible terms is used. Must be a function the smooth term from mgcv's smoothCon and an argument df.
df	degrees of freedom for all non-linear structural terms (default = 7); either one common value or a list of the same length as number of parameters; if different df values need to be assigned to different smooth terms, use df as an argument for s(), te() or ti()

<code>null_space_penalty</code>	logical value; if TRUE, the null space will also be penalized for smooth effects. Per default, this is equal to the value give in <code>variational</code> .
<code>absorb_cons</code>	logical; adds identifiability constraint to the basisi. See <code>?mgcv::smoothCon</code> for more details.
<code>anisotropic</code>	whether or not use anisotropic smoothing (default is TRUE)
<code>zero_constraint_for_smooths</code>	logical; the same as <code>absorb_cons</code> , but done explicitly. If true a constraint is put on each smooth to have zero mean. Can be a vector of <code>length(list_of_formulas)</code> for each distribution parameter.
<code>hat1</code>	logical; if TRUE, the smoothing parameter is defined by the trace of the hat matrix <code>sum(diag(H))</code> , else <code>sum(diag(2*H-HH))</code>
<code>sp_scale</code>	function of response; for scaling the penalty (1/n per default)

Value

Returns a list with options

`plot.deepregression` *Generic functions for deepregression models*

Description

- Generic functions for deepregression models
- Predict based on a deepregression object
- Function to extract fitted distribution
- Fit a deepregression model (pendant to fit for keras)
- Extract layer weights / coefficients from model
- Print function for deepregression model
- Cross-validation for deepgression objects
- mean of model fit
- Standard deviation of fit distribution
- Calculate the distribution quantiles

Usage

```
## S3 method for class 'deepregression'
plot(
  x,
  which = NULL,
  which_param = 1,
  only_data = FALSE,
  grid_length = 40,
```

```
    type = "b",
    ...
)

## S3 method for class 'deepregression'
predict(
  object,
  newdata = NULL,
  batch_size = NULL,
  apply_fun = tfd_mean,
  convert_fun = as.matrix,
  ...
)

## S3 method for class 'deepregression'
fitted(object, apply_fun = tfd_mean, ...)

## S3 method for class 'deepregression'
fit(
  object,
  batch_size = 32,
  epochs = 10,
  early_stopping = FALSE,
  early_stopping_metric = "val_loss",
  verbose = TRUE,
  view_metrics = FALSE,
  patience = 20,
  save_weights = FALSE,
  validation_data = NULL,
  validation_split = ifelse(is.null(validation_data), 0.1, 0),
  callbacks = list(),
  convertfun = function(x) tf$constant(x, dtype = "float32"),
  ...
)

## S3 method for class 'deepregression'
coef(object, which_param = 1, type = NULL, ...)

## S3 method for class 'deepregression'
print(x, ...)

## S3 method for class 'deepregression'
cv(
  x,
  verbose = FALSE,
  patience = 20,
  plot = TRUE,
  print_folds = TRUE,
```

```

  cv_folds = 5,
  stop_if_nan = TRUE,
  mylapply = lapply,
  save_weights = FALSE,
  callbacks = list(),
  save_fun = NULL,
  ...
)

## S3 method for class 'deepregression'
mean(x, data = NULL, ...)

## S3 method for class 'deepregression'
stddev(x, data = NULL, ...)

## S3 method for class 'deepregression'
quant(x, data = NULL, probs, ...)

```

Arguments

<code>x</code>	a deepregression object
<code>which</code>	character vector or number(s) identifying the effect to plot; default plots all effects
<code>which_param</code>	integer, indicating for which distribution parameter coefficients should be returned (default is first parameter)
<code>only_data</code>	logical, if TRUE, only the data for plotting is returned
<code>grid_length</code>	the length of an equidistant grid at which a two-dimensional function is evaluated for plotting.
<code>type</code>	either NULL (all types of coefficients are returned), "linear" for linear coefficients or "smooth" for coefficients of smooth terms
<code>...</code>	arguments passed to the <code>predict</code> function
<code>object</code>	a deepregression model
<code>newdata</code>	optional new data, either <code>data.frame</code> or <code>list</code>
<code>batch_size</code>	integer, the batch size used for mini-batch training
<code>apply_fun</code>	function applied to fitted distribution, per default <code>tfd_mean</code>
<code>convert_fun</code>	how should the resulting tensor be converted, per default <code>as.matrix</code>
<code>epochs</code>	integer, the number of epochs to fit the model
<code>early_stopping</code>	logical, whether early stopping should be user.
<code>early_stopping_metric</code>	character, based on which metric should early stopping be triggered (default: "val_loss")
<code>verbose</code>	whether to print training in each fold
<code>view_metrics</code>	logical, whether to trigger the Viewer in RStudio / Browser.
<code>patience</code>	number of patience for early stopping

save_weights	logical, whether to save weights in each epoch.
validation_data	optional specified validation data
validation_split	float in [0,1] defining the amount of data used for validation
callbacks	a list of callbacks used for fitting
convertfun	function to convert R into Tensor object
plot	whether to plot the resulting losses in each fold
print_folds	whether to print the current fold
cv_folds	an integer if list with train and test data sets
stop_if_nan	logical; whether to stop CV if NaN values occur
mylapply	lapply function to be used; defaults to lapply
save_fun	function applied to the model in each fold to be stored in the final result
data	either NULL or a new data set
probs	the quantile value(s)

Value

Returns an object drCV, a list, one list element for each fold containing the model fit and the weighthistory.

plot_cv

Plot CV results from deepregression

Description

Plot CV results from deepregression

Usage

```
plot_cv(x, what = c("loss", "weight"), ...)
```

Arguments

x	drCV object returned by cv.deepregression
what	character indicating what to plot (currently supported 'loss' or 'weights')
...	further arguments passed to matplot

prepare_data

Function to prepare data based on parsed formulas

Description

Function to prepare data based on parsed formulas

Usage

```
prepare_data(pfc)
```

Arguments

pfc	list of processor transformed formulas
-----	--

Value

list of matrices or arrays

prepare_newdata

Function to prepare new data based on parsed formulas

Description

Function to prepare new data based on parsed formulas

Usage

```
prepare_newdata(pfc, newdata)
```

Arguments

pfc	list of processor transformed formulas
-----	--

newdata	list in the same format as the original data
---------	--

Value

list of matrices or arrays

processor

Control function to define the processor for terms in the formula

Description

Control function to define the processor for terms in the formula

Usage

```
processor(  
  form,  
  data,  
  controls,  
  output_dim,  
  param_nr,  
  specials_to_oz = c(),  
  automatic_oz_check = TRUE,  
  identify_intercept = FALSE,  
  ...  
)
```

Arguments

form	the formula to be processed
data	the data for the terms in the formula
controls	controls for gam terms
output_dim	the output dimension of the response
param_nr	integer; identifier for the distribution parameter
specials_to_oz	specials that should be automatically checked for
automatic_oz_check	logical; whether to automatically check for DNNs to be orthogonalized
identify_intercept	logical; whether to make the intercept automatically identifiable
...	further processors

Value

returns a processor function

quant	<i>Generic quantile function</i>
-------	----------------------------------

Description

Generic quantile function

Usage

```
quant(x, ...)
```

Arguments

x	object
...	further arguments passed to the class-specific function

separate_define_relation	
--------------------------	--

Function to define orthogonalization connections in the formula

Description

Function to define orthogonalization connections in the formula

Usage

```
separate_define_relation(
  form,
  specials,
  specials_to_oz,
  automatic_oz_check = TRUE,
  identify_intercept = FALSE
)
```

Arguments

form	a formula for one distribution parameter
specials	specials in formula to handle separately
specials_to_oz	parts of the formula to orthogonalize
automatic_oz_check	logical; automatically check if terms must be orthogonalized
identify_intercept	logical; whether to make the intercept identifiable

Value

Returns a list of formula components with ids and assignments for orthogonalization

stddev*Generic sd function*

Description

Generic sd function

Usage

```
stddev(x, ...)
```

Arguments

x	object
...	further arguments passed to the class-specific function

stop_iter_cv_result *Function to get the stoppting iteration from CV*

Description

Function to get the stoppting iteration from CV

Usage

```
stop_iter_cv_result(  
  res,  
  thisFUN = mean,  
  loss = "validloss",  
  whichFUN = which.min  
)
```

Arguments

res	result of cv call
thisFUN	aggregating function applied over folds
loss	which loss to use for decision
whichFUN	which function to use for decision

<code>subnetwork_init</code>	<i>Initializes a Subnetwork based on the Processed Additive Predictor</i>
------------------------------	---

Description

Initializes a Subnetwork based on the Processed Additive Predictor

Usage

```
subnetwork_init(
  pp,
  deep_top = NULL,
  orthog_fun = orthog_tf,
  split_fun = split_model,
  param_nr = 1
)
```

Arguments

<code>pp</code>	processed predictor list from processor
<code>deep_top</code>	keras layer if the top part of the deep network after orthogonalization is different to the one extracted from the provided network
<code>orthog_fun</code>	function used for orthogonalization
<code>split_fun</code>	function to split the network to extract head
<code>param_nr</code>	integer number for the distribution parameter

Value

returns a list of input and output for this additive predictor

<code>tfd_zinb</code>	<i>Implementation of a zero-inflated negbinom distribution for TFP</i>
-----------------------	--

Description

Implementation of a zero-inflated negbinom distribution for TFP

Usage

```
tfd_zinb(mu, r, probs)
```

Arguments

<code>mu, r</code>	parameter of the negbin_ls distribution
<code>probs</code>	vector of probabilités of length 2 (probability for poisson and probability for 0s)

`tfd_zip`*Implementation of a zero-inflated poisson distribution for TFP*

Description

Implementation of a zero-inflated poisson distribution for TFP

Usage

```
tfd_zip(lambda, probs)
```

Arguments

lambda	scalar value for rate of poisson distribution
probs	vector of probabilities of length 2 (probability for poisson and probability for 0s)

`tf_stride_cols`*Function to index tensors columns*

Description

Function to index tensors columns

Usage

```
tf_stride_cols(A, start, end = NULL)
```

Arguments

A	tensor
start	first index
end	last index (equals start index if NULL)

Value

sliced tensor

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