

# Package ‘distrom’

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**Title** Distributed Multinomial Regression

**Version** 1.0.1

**Depends** R (>= 2.15), Matrix, gamlr, parallel, methods, stats

**Suggests** MASS, textir

**Description** Fast distributed/parallel estimation for multinomial logistic regression via Poisson factorization and the 'gamlr' package. For details see: Taddy (2015, AoAS), Distributed Multinomial Regression, <[arXiv:1311.6139](https://arxiv.org/abs/1311.6139)>.

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**License** GPL-3

**URL** <https://github.com/TaddyLab/distrom>

**NeedsCompilation** no

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collapse	<i>Data checking and binning</i>
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## Description

Collapses counts along equal levels of binned covariates.

**Usage**

```
collapse(v, counts, mu=NULL, bins=NULL)
```

**Arguments**

v	Either matrix or Matrix of covariates (matches covars in dmr).
counts	Either matrix or Matrix of multinomial counts, or a factor (matches counts in dmr).
mu	Possible pre-specified fixed effects for dmr; otherwise they are calculated here.
bins	The number of quantile bins into which we collapse v. bins=NULL does no collapsing.

**Details**

For each column of v, aggregates the observations into bins defined by their average value. Both v and counts are then collapsed according to levels of the interaction across implied bin-factors, and the number of observations in each bin is recorded as n. Look at the code of the dmr function to see collapse used in practice.

**Value**

A list containing collapsed and formatted v, counts, and nbin, along with  $\mu = \log(\text{rowSums}(\text{counts}))$ , the plug-in fixed effect estimates for dmr.

**Author(s)**

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**See Also**

we8there

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 dmr

*Distributed Multinomial Regression*


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**Description**

Gamma-lasso path estimation for a multinomial logistic regression factorized into independent Poisson log regressions.

**Usage**

```
dmr(cl, covars, counts, mu=NULL, bins=NULL, verb=0, cv=FALSE, ...)
## S3 method for class 'dmr'
coef(object, ...)
## S3 method for class 'dmr'
predict(object, newdata,
type=c("link", "response", "class"), ...)
```

**Arguments**

<code>cl</code>	A parallel library socket cluster. If <code>is.null(cl)</code> , everything is done in serial. See <code>help(parallel)</code> , <code>help(makeCluster)</code> , and our examples here for details.
<code>covars</code>	A dense matrix or sparse <code>Matrix</code> of covariates. This should not include the intercept.
<code>counts</code>	A dense matrix or sparse <code>Matrix</code> of response counts.
<code>mu</code>	Pre-specified fixed effects for each observation in the Poisson regression linear equation. If <code>mu=NULL</code> , then we use <code>log(rowSums(x))</code> . Note that if <code>bins</code> is non-null then this argument is ignored and <code>mu</code> is recalculated on the collapsed data.
<code>bins</code>	Number of bins into which we will attempt to collapse each column of <code>covars</code> . Since sums of multinomials with equal probabilities are also multinomial, the model is then fit to these collapsed ‘observations’. <code>bins=NULL</code> does no collapsing.
<code>verb</code>	Whether to print some info. <code>max(0, verb-1)</code> is passed on to <code>gam1r</code> and will print if you created an outfile when specifying <code>cl</code> .
<code>cv</code>	A flag for whether to use <code>cv.gam1r</code> instead of <code>gam1r</code> for each Poisson regression.
<code>type</code>	For <code>predict.dmr</code> , this is the scale upon which you want prediction. Under "link", just the linear map <code>newdata</code> times object, under "response" the fitted multinomial probabilities, under "class" the max-probability class label. For sufficient reductions see the <code>srproj</code> function of the <code>textir</code> library.
<code>newdata</code>	A <code>Matrix</code> with the same number of columns as <code>covars</code> .
<code>...</code>	Additional arguments to <code>gam1r</code> , <code>cv.gam1r</code> , and their associated methods.
<code>object</code>	A <code>dmr</code> list of fitted <code>gam1r</code> models for each response category.

**Details**

`dmr` fits multinomial logistic regression by assuming that, unconditionally on the ‘size’ (total count across categories) each individual category count has been generated as a Poisson

$$x_{ij} \sim Po(\exp[\mu_i + \alpha_j + \beta v_i]).$$

We [default] plug-in estimate  $\hat{\mu}_i = \log(m_i)$ , where  $m_i = \sum_j x_{ij}$  and  $p$  is the dimension of  $x_i$ . Then each individual is outsourced to Poisson regression in the `gam1r` package via the `parLapply` function of the `parallel` library. The output from `dmr` is a list of `gam1r` fitted models.

`coef.dmr` builds a matrix of multinomial logistic regression coefficients from the `length(object)` list of `gam1r` fits. Default selection under `cv=FALSE` uses an information criteria via `AICc` on Poisson deviance for each individual response dimension (see `gam1r`). Combined coefficients across all dimensions are then returned as a `dmrcoef` `s4`-class object.

`predict.dmr` takes either a `dmr` or `dmrcoef` object and returns predicted values for `newdata` on the scale defined by the `type` argument.

**Value**

`dmr` returns the `dmr` `s3` object: an `ncol(counts)`-length list of fitted `gam1r` objects, with the added attributes `nlambda`, `mu`, and `nobs`.

**Author(s)**

Matt Taddy <mataddy@gmail.com>

**References**

Taddy (2015 AoAS) Distributed Multinomial Regression

Taddy (2017 JCGS) One-step Estimator Paths for Concave Regularization, the Journal of Computational and Graphical Statistics

Taddy (2013 JASA) Multinomial Inverse Regression for Text Analysis

**See Also**

dmrcoef-class, cv.dmr, AICc, and the gamlr and textir packages.

**Examples**

```
library(MASS)
data(fgl)

## make your cluster
## FORK is faster but memory heavy, and doesn't work on windows.
cl <- makeCluster(2,type=ifelse(.Platform$OS.type=="unix","FORK","PSOCK"))
print(cl)

## fit in parallel
fits <- dmr(cl, fgl[,1:9], fgl$type, verb=1)

## its good practice stop the cluster once you're done
stopCluster(cl)

## Individual Poisson model fits and AICc selection
par(mfrow=c(3,2))
for(j in 1:6){
plot(fits[[j]])
mtext(names(fits)[j],font=2,line=2) }

## AICc model selection
B <- coef(fits)

## Fitted probability by true response
par(mfrow=c(1,1))
P <- predict(B, fgl[,1:9], type="response")
boxplot(P[cbind(1:214,fgl$type)]~fgl$type,
ylab="fitted prob of true class")
```

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dmrcoef-class	Class "dmrcoef"
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### Description

The extended `dgCMatrx` class for output from `coef.dmr`.

### Details

This is the class for a covariate matrix from `dmr` regression; it inherits the `dgCMatrx` class as defined in the `Matrix` library. In particular, this is the `ncol(covars)` by `ncol(counts)` matrix of logistic regression coefficients chosen in `coef.dmr` from the regularization paths for each category.

### Objects from the Class

Objects can be created only by a call to the `coef.dmr` function.

### Slots

**i:** From `dgCMatrx`: the row indices.  
**p:** From `dgCMatrx`: the column pointers.  
**Dim:** From `dgCMatrx`: the dimensions.  
**Dimnames:** From `dgCMatrx`: the list of labels.  
**x:** From `dgCMatrx`: the nonzero entries.  
**factors:** From `dgCMatrx`.

### Extends

Class `dgCMatrx`, directly.

### Methods

**predict** signature(object = "dmrcoef"): Prediction for a given `dmrcoef` matrix. Takes the same arguments as `predict.dmr`, but will be faster (since `coef.dmr` is called inside `predict.dmr`).

### Author(s)

Matt Taddy <mataddy@gmail.com>

### See Also

`dmr`, `coef.dmr`, `predict.dmr`

### Examples

```
showClass("dmrcoef")
```

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