

# Package ‘biglmm’

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**Type** Package

**Title** Bounded Memory Linear and Generalized Linear Models

**Version** 0.9-1

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**Description** Regression for data too large to fit in memory. This package functions exactly like the 'biglm' package, but works with later versions of R.

**License** GPL

**Suggests** RSQLite, RODB

**Depends** DBI, methods

**Enhances** leaps

**NeedsCompilation** yes

**Repository** CRAN

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**RoxygenNote** 6.0.1

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bigglm

*Bounded memory linear regression***Description**

bigglm creates a generalized linear model object that uses only  $p^2$  memory for  $p$  variables.

**Usage**

```
bigglm(formula, data, family=gaussian(),...)
## S3 method for class 'data.frame'
bigglm(formula, data,...,chunksize=5000)
## S3 method for class 'function'
bigglm(formula, data, family=gaussian(),
        weights=NULL, sandwich=FALSE, maxit=8, tolerance=1e-7,
        start=NULL,quiet=FALSE,...)
## S3 method for class 'RODBC'
bigglm(formula, data, family=gaussian(),
        tablename, ..., chunksize=5000)
## S4 method for signature 'ANY,DBIConnection'
bigglm(formula, data, family=gaussian(),
        tablename, ..., chunksize=5000)
## S3 method for class 'bigglm'
vcov(object,dispersion=NULL, ...)
## S3 method for class 'bigglm'
deviance(object,...)
## S3 method for class 'bigglm'
family(object,...)
## S3 method for class 'bigglm'
AIC(object,...,k=2)
```

**Arguments**

formula	A model formula
data	See Details below. Method dispatch is on this argument
family	A glm family object
chunksize	Size of chunks for processng the data frame
weights	A one-sided, single term formula specifying weights
sandwich	TRUE to compute the Huber/White sandwich covariance matrix (uses $p^4$ memory rather than $p^2$ )
maxit	Maximum number of Fisher scoring iterations
tolerance	Tolerance for change in coefficient (as multiple of standard error)
start	Optional starting values for coefficients. If NULL, maxit should be at least 2 as some quantities will not be computed on the first iteration

object	A bigglm object
dispersion	Dispersion parameter, or NULL to estimate
tablename	For the SQLiteConnection method, the name of a SQL table, or a string specifying a join or nested select
k	penalty per parameter for AIC
quiet	When FALSE, warn if the fit did not converge
...	Additional arguments

### Details

The data argument may be a function, a data frame, or a SQLiteConnection or RODBConnection object.

When it is a function the function must take a single argument `reset`. When this argument is FALSE it returns a data frame with the next chunk of data or NULL if no more data are available. When `reset=TRUE` it indicates that the data should be reread from the beginning by subsequent calls. The chunks need not be the same size or in the same order when the data are reread, but the same data must be provided in total. The `bigglm.data.frame` method gives an example of how such a function might be written, another is in the Examples below.

The model formula must not contain any data-dependent terms, as these will not be consistent when updated. Factors are permitted, but the levels of the factor must be the same across all data chunks (empty factor levels are ok). Offsets are allowed (since version 0.8).

The SQLiteConnection and RODBConnection methods load only the variables needed for the model, not the whole table. The code in the SQLiteConnection method should work for other DBI connections, but I do not have any of these to check it with.

### Value

An object of class `bigglm`

### References

Algorithm AS274 Applied Statistics (1992) Vol.41, No. 2

### See Also

[biglm](#), [glm](#)

### Examples

```
data(trees)
ff<-log(Volume)~log(Girth)+log(Height)
a <- bigglm(ff,data=trees, chunksize=10, sandwich=TRUE)
summary(a)

gg<-log(Volume)~log(Girth)+log(Height)+offset(2*log(Girth)+log(Height))
b <- bigglm(gg,data=trees, chunksize=10, sandwich=TRUE)
summary(b)
```

```

## requires internet access
make.data<-function(urlname, chunksize,...){
  conn<-NULL
  function(reset=FALSE){
    if(reset){
      if(!is.null(conn)) close(conn)
      conn<<-url(urlname,open="r")
    } else{
      rval<-read.table(conn, nrows=chunksize,...)
      if (nrow(rval)==0) {
        close(conn)
        conn<<-NULL
        rval<-NULL
      }
      return(rval)
    }
  }
}

airpoll<-make.data("http://faculty.washington.edu/tlumley/NO2.dat",
  chunksize=150,
  col.names=c("logno2", "logcars", "temp", "windsp",
    "tempgrad", "winddir", "hour", "day"))

b<-bigglm(exp(logno2)~logcars+temp+windsp,
  data=airpoll, family=Gamma(log),
  start=c(2,0,0,0),maxit=10)
summary(b)

```

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biglm

*Bounded memory linear regression*


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### Description

biglm creates a linear model object that uses only  $p^2$  memory for  $p$  variables. It can be updated with more data using update. This allows linear regression on data sets larger than memory.

### Usage

```

biglm(formula, data, weights=NULL, sandwich=FALSE)
## S3 method for class 'biglm'
update(object, moredata,...)
## S3 method for class 'biglm'
vcov(object,...)
## S3 method for class 'biglm'
coef(object,...)
## S3 method for class 'biglm'

```

```
summary(object,...)
## S3 method for class 'biglm'
AIC(object,...,k=2)
## S3 method for class 'biglm'
deviance(object,...)
```

### Arguments

formula	A model formula
weights	A one-sided, single term formula specifying weights
sandwich	TRUE to compute the Huber/White sandwich covariance matrix (uses $p^4$ memory rather than $p^2$ )
object	A biglm object
data	Data frame that must contain all variables in formula and weights
moredata	Additional data to add to the model
...	Additional arguments for future expansion
k	penalty per parameter for AIC

### Details

The model formula must not contain any data-dependent terms, as these will not be consistent when updated. Factors are permitted, but the levels of the factor must be the same across all data chunks (empty factor levels are ok). Offsets are allowed (since version 0.8).

### Value

An object of class biglm

### References

Algorithm AS274 Applied Statistics (1992) Vol.41, No. 2

### See Also

lm

### Examples

```
data(trees)
ff<-log(Volume)~log(Girth)+log(Height)

chunk1<-trees[1:10,]
chunk2<-trees[11:20,]
chunk3<-trees[21:31,]

a <- biglm(ff,chunk1)
a <- update(a,chunk2)
a <- update(a,chunk3)
```

```
summary(a)
deviance(a)
AIC(a)
```

---

predict.bigglm      *Predictions from a biglm/bigglm*

---

### Description

Computes fitted means and standard errors at new data values after fitting a model with biglm or bigglm.

### Usage

```
## S3 method for class 'bigglm'
predict(object, newdata, type = c("link", "response"),
        se.fit = FALSE, make.function = FALSE, ...)
## S3 method for class 'biglm'
predict(object, newdata=NULL, se.fit = FALSE, make.function = FALSE, ...)
```

### Arguments

object	fitted model
newdata	data frame with variables for new values
type	link is on the linear predictor scale, response is the response
se.fit	Compute standard errors?
make.function	If TRUE return a prediction function, see Details below
...	not used

### Details

When `make.function` is TRUE, the return value is either a single function that computes the fitted values or a list of two functions that compute the fitted values and standard errors. The input to these functions is the design matrix, without the intercept column. This allows the relatively time-consuming calls to `model.frame()` and `model.matrix()` to be avoided.

### Value

Either a vector of predicted values or a data frame with predicted values and standard errors.

### Author(s)

based on code by Christophe Dutang

### References

~put references to the literature/web site here ~

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### **See Also**

[predict.glm](#), [biglm](#), [bigglm](#)

### **Examples**

```
example(biglm)
predict(a,newdata=trees)
f<-predict(a,make.function=TRUE)
X<- with(trees, cbind(log(Girth),log(Height)))
f(X)
```

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