

Package ‘gamm4’

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Title Generalized additive mixed models using mgcv and lme4

Description Fit generalized additive mixed models via a version of mgcv’s gamm function, using lme4 for estimation via Fabian Scheipl’s trick.

Depends R (>= 2.9.0), methods, Matrix, lme4 (>= 0.999375-31), mgcv (>= 1.7-7)

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gamm4 *Generalized Additive Mixed Models using lme4 and mgcv*

Description

Fits the specified generalized additive mixed model (GAMM) to data, by a call to `lmer` in the normal errors identity link case, or by a call to `glmer` otherwise (see [lmer](#)). Smoothness selection is by REML in the gaussian additive case and ML otherwise.

`gamm4` is based on [gamm](#) from package `mgcv`, but uses `lme4` rather than `nlme` as the underlying fitting engine via a trick due to Fabian Scheipl. `gamm4` is more robust numerically than [gamm](#), and by avoiding PQL gives better performance for binary and low mean count data. Its main disadvantage is that it can not handle most multi-penalty smooths (i.e. not `te` type tensor products or adaptive

smooths) and there is no facility for nlme style correlation structures. Tensor product smoothing is available via `t2` terms.

For fitting generalized additive models without random effects, `gamm4` is much slower than `gam` and has slightly worse MSE performance than `gam` with REML smoothness selection.

To use this function effectively it helps to be quite familiar with the use of `gam` and `lmer`.

Usage

```
gamm4(formula, random=NULL, family=gaussian(), data=list(), weights=NULL,
       subset=NULL, na.action, knots=NULL, ...)
```

Arguments

<code>formula</code>	A GAM formula (see also <code>formula.gam</code> and <code>gam.models</code>). This is like the formula for a <code>glm</code> except that smooth terms (<code>s</code> and <code>t2</code> but not <code>te</code>) can be added to the right hand side of the formula. Note that <code>ids</code> for smooths and fixed smoothing parameters are not supported.
<code>random</code>	An optional formula specifying the random effects structure in <code>lmer</code> style. See example below.
<code>family</code>	A family as used in a call to <code>glm</code> or <code>gam</code> .
<code>data</code>	A data frame or list containing the model response variable and covariates required by the formula. By default the variables are taken from <code>environment(formula)</code> , typically the environment from which <code>gamm4</code> is called.
<code>weights</code>	a vector of prior weights on the observations. <code>NULL</code> is equivalent to a vector of 1s. Used, in particular, to supply the number-of-trials for binomial data, when the response is proportion of successes.
<code>subset</code>	an optional vector specifying a subset of observations to be used in the fitting process.
<code>na.action</code>	a function which indicates what should happen when the data contain 'NA's. The default is set by the 'na.action' setting of 'options', and is 'na.fail' if that is unset. The "factory-fresh" default is 'na.omit'.
<code>knots</code>	this is an optional list containing user specified knot values to be used for basis construction. Different terms can use different numbers of knots, unless they share a covariate.
<code>...</code>	further arguments for passing on e.g. to <code>lmer</code>

Details

A generalized additive mixed model is a generalized linear mixed model in which the linear predictor depends linearly on unknown smooth functions of some of the covariates ('smooths' for short). `gamm4` follows the approach taken by package `mgcv` and represents the smooths using penalized regression spline type smoothers, of moderate rank. For estimation purposes the penalized component of each smooth is treated as a random effect term, while the unpenalized component is treated as fixed. The wiggleness penalty matrix for the smooth is in effect the precision matrix when the smooth is treated as a random effect. Estimating the degree of smoothness of the term amounts to estimating the variance parameter for the term.

`gamm4` uses the same reparameterization trick employed by `gamm` to allow any single quadratic penalty smoother to be used (see Wood, 2004, or 2006 for details). Given the reparameterization then Fabian Scheipl's trick for getting `lmer` to fit a GAMM can be employed (see package `amer`). Estimation is by Maximum Likelihood in the generalized case, and REML in the gaussian additive model case. `gamm4` allows the random effects specifiable with `lmer` to be combined with any number of any of the (single penalty) smooth terms available in `gam` from package `mgcv` as well as `t2` tensor product smooths. Note that the model comparison on the basis of the (Laplace approximate) log likelihood is possible with GAMMs fitted by `gamm4`.

As in `gamm` the smooth estimates are assumed to be of interest, and a covariance matrix is returned which enables Bayesian credible intervals for the smooths to be constructed, which treat all the terms in random as random.

For details on how to condition smooths on factors, set up varying coefficient models, do signal regression or set up terms involving linear functionals of smooths, see `gam.models`, but note that the type tensor product and adaptive smooths are not available with `gamm4`.

Value

Returns a list with two items:

<code>gam</code>	an object of class <code>gam</code> . At present this contains enough information to use <code>predict</code> , <code>plot</code> , <code>summary</code> and <code>print</code> methods and <code>vis.gam</code> , from package <code>mgcv</code> but not to use e.g. the <code>anova</code> method function to compare models.
<code>mer</code>	the fitted model object returned by <code>lmer</code> or <code>glmer</code> . Extra random and fixed effect terms will appear relating to the estimation of the smooth terms. Note that unlike <code>lme</code> objects returned by <code>gamm</code> , everything in this object always relates to the fitted model itself, and never to a PQL working approximation: hence the usual methods of model comparison are entirely legitimate.

WARNINGS

If you don't need random effects in addition to the smooths, then `gam` is substantially faster, gives fewer convergence warnings, and slightly better MSE performance (based on simulations).

Models must contain at least one random effect: either a smooth with non-zero smoothing parameter, or a random effect specified in argument `random`.

Note that the `gam` object part of the returned object is not complete in the sense of having all the elements defined in `gamObject` and does not inherit from `glm`: hence e.g. multi-model `anova` calls will not work.

Linked smoothing parameters, adaptive smoothing and `te` terms are not supported.

This routine is obviously less well tested than `gamm`.

Author(s)

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References

Bates D. and M. Maechler (2009). lme4: Linear mixed-effects models using S4 classes. <http://CRAN.R-project.org/package=lme4>

Scheipl, F. (2009) amer: Additive mixed models with lme4. <http://CRAN.R-project.org/package=amer>

Wood, S.N. (2004) Stable and efficient multiple smoothing parameter estimation for generalized additive models. Journal of the American Statistical Association. 99:673-686

Wood S.N. (2006) Generalized Additive Models: An Introduction with R. Chapman and Hall/CRC Press.

For more GAMM references see [gamm](#)

<http://www.maths.bath.ac.uk/~sw283/>

See Also

[gam](#), [gamm](#), [gam.models](#), [lmer](#), [predict.gam](#), [plot.gam](#), [summary.gam](#), [s](#), [vis.gam](#)

Examples

```
## NOTE: some 'gamm' calls are flagged as do not run simply to
## save time in package checking.

#####
## A simple additive model...
#####
library(gamm4)

set.seed(0)
dat <- gamSim(1,n=400,scale=2) ## simulate data

## Not run:
b <- gamm(y~s(x0)+s(x1)+s(x2)+s(x3),data=dat)
plot(b$gam,pages=1)
summary(b$gam) ## gam style summary of fitted model

## End(Not run)

br <- gamm4(y~s(x0)+s(x1)+s(x2)+s(x3),data=dat)
plot(br$gam,pages=1)

summary(br$gam) ## same from 'gamm4'
summary(br$mer) ## underlying mixed model
anova(br$gam)
rm(dat)

#####
## Add a factor to the linear predictor, to be modelled as random
## and make response Poisson. Again compare 'gamm' and 'gamm4'
#####
```

```

dat <- gamSim(6,n=400,scale=.2,dist="poisson")

## Not run:
b2<-gamm(y~s(x0)+s(x1)+s(x2)+s(x3),family=poisson,
         data=dat,random=list(fac=~1))
plot(b2$gam,pages=1)

## End(Not run)
b2r<-gamm4(y~s(x0)+s(x1)+s(x2)+s(x3),family=poisson,
          data=dat,random = ~ (1|fac))

plot(b2r$gam,pages=1)

rm(dat)
vis.gam(b2r$gam)

#####
# Multivariate varying coefficient
# With crossed and nested random
# effects.
#####

## Start by simulating data...

f0 <- function(x, z, sx = 0.3, sz = 0.4) {
  (pi^sx * sz) * (1.2 * exp(-(x - 0.2)^2/sx^2 - (z -
    0.3)^2/sz^2) + 0.8 * exp(-(x - 0.7)^2/sx^2 -
    (z - 0.8)^2/sz^2))
}
f1 <- function(x2) 2 * sin(pi * x2)
f2 <- function(x2) exp(2 * x2) - 3.75887
f3 <- function(x2) 0.2 * x2^11 * (10 * (1 - x2))^6 + 10 * (10 * x2)^3 *
  (1 - x2)^10

n <- 1000

## first set up a continuous-within-group effect...

g <- factor(sample(1:50,n,replace=TRUE)) ## grouping factor
x <- runif(n) ## continuous covariate
X <- model.matrix(~g-1)
mu <- X%*%rnorm(50)*.5 + (x*X)%*%rnorm(50)

## now add nested factors...
a <- factor(rep(1:20,rep(50,20)))
b <- factor(rep(rep(1:25,rep(2,25)),rep(20,50)))
Xa <- model.matrix(~a-1)
Xb <- model.matrix(~a/b-a-1)
mu <- mu + Xa%*%rnorm(20) + Xb%*%rnorm(500)*.5

## finally simulate the smooth terms

```

```

v <- runif(n);w <- runif(n);z <- runif(n)
r <- runif(n)
mu <- mu + f0(v,w)*z*10 + f3(r)

y <- mu + rnorm(n)*2 ## response data

## First compare gamm and gamm4 on a reduced model

br <- gamm4(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),random = ~ (1|a/b))

## Not run:
ba <- gamm(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),random = list(a=~1,b=~1),method="REML")

## End(Not run)

par(mfrow=c(2,2))
plot(br$gam)
## Not run:
plot(ba$gam)

## End(Not run)
## now fit the full model

br <- gamm4(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),random = ~ (x+0|g) + (1|g) + (1|a/b))

br$mer
br$gam
plot(br$gam)

## try a Poisson example, based on the same linear predictor...

lp <- mu/5
y <- rpois(exp(lp),exp(lp)) ## simulated response

## again compare gamm and gamm4 on reduced model

br <- gamm4(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),family=poisson,random = ~ (1|a/b))

## Not run:
ba <- gamm(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),family=poisson,random = list(a=~1,b=~1))

## End(Not run)

par(mfrow=c(2,2))
plot(br$gam);

## Not run:
plot(ba$gam)

## End(Not run)
## and now fit full version (very slow)...
## Not run:
br <- gamm4(y ~ s(v,w,by=z) + s(r,k=20,bs="cr"),family=poisson,random = ~ (x|g) + (1|a/b))

```

```

br$mer
br$gam
plot(br$gam)

## End(Not run)

#####
# Different smooths of x2 depending
# on factor 'fac'...
#####
dat <- gamSim(4)

br <- gamm4(y ~ fac+s(x2,by=fac)+s(x0),data=dat)
plot(br$gam,pages=1)
summary(br$gam)

#####
# Timing comparison with 'gam'... #
#####

dat <- gamSim(1,n=600,dist="binary",scale=.33)

system.time(lr.fit0 <- gam(y~s(x0)+s(x1)+s(x2)+s(x3),
                          family=binomial,data=dat,method="ML"))

system.time(lr.fit <- gamm4(y~s(x0)+s(x1)+s(x2)+s(x3),
                           family=binomial,data=dat))

lr.fit0;lr.fit$gam
cor(fitted(lr.fit0),fitted(lr.fit$gam))

## plot model components with truth overlaid in red
op <- par(mfrow=c(2,2))
fn <- c("f0","f1","f2","f3");xn <- c("x0","x1","x2","x3")
for (k in 1:4) {
  plot(lr.fit$gam,select=k)
  ff <- dat[[fn[k]];xx <- dat[[xn[k]]]
  ind <- sort.int(xx,index.return=TRUE)$ix
  lines(xx[ind],(ff-mean(ff))[ind]*.33,col=2)
}
par(op)

#####
# Simple random effects comparison
# with 'gam'
#####

dat <- gamSim(1,n=400,scale=2) ## simulate 4 term additive truth
## Now add some random effects to the simulation. Response is
## grouped into one of 20 groups by 'fac' and each groups has a
## random effect added...
fac <- as.factor(sample(1:20,400,replace=TRUE))

```

```

X <- model.matrix(~fac-1)
b <- rnorm(20)*.5
dat$y <- dat$y + X%*%b
dat$fac <- fac

## now fit appropriate random effect model using 'gam'...

rm <- gam(y~ s(fac,bs="re")+s(x0)+s(x1)+s(x2)+s(x3),data=dat,
          method="REML")
plot(rm,pages=1)
## Get estimated random effects standard deviation...
gam.vcomp(rm)

## Now do same thing with gamm4
rmr <- gamm4(y~ s(x0)+s(x1)+s(x2)+s(x3),data=dat,
             random = ~ (1|fac))
plot(rmr$gam,pages=1)
rmr$mer

#####
## A "signal" regression example, in
## which a univariate response depends
## on functional predictors.
#####

## simulate data first...

rf <- function(x=seq(0,1,length=100)) {
## generates random functions...
  m <- ceiling(runif(1)*5) ## number of components
  f <- x*0;
  mu <- runif(m,min(x),max(x));sig <- (runif(m)+.5)*(max(x)-min(x))/10
  for (i in 1:m) f <- f+ dnorm(x,mu[i],sig[i])
  f
}

x <- seq(0,1,length=100) ## evaluation points

## example functional predictors...
par(mfrow=c(3,3));for (i in 1:9) plot(x,rf(x),type="l",xlab="x")

## simulate 200 functions and store in rows of L...
L <- matrix(NA,200,100)
for (i in 1:200) L[i,] <- rf() ## simulate the functional predictors

f2 <- function(x) { ## the coefficient function
  (0.2*x^11*(10*(1-x))^6+10*(10*x)^3*(1-x)^10)/10
}

f <- f2(x) ## the true coefficient function

y <- L%*%f + rnorm(200)*20 ## simulated response data

```

```
## Now fit the model  $E(y) = L * f(x)$  where  $f$  is a smooth function.
## The summation convention is used to evaluate smooth at each value
## in matrix  $X$  to get matrix  $F$ , say. Then  $\text{rowSum}(L * F)$  gives  $E(y)$ .

## create matrix of eval points for each function. Note that
## 'smoothCon' is smart and will recognize the duplication...
X <- matrix(x,200,100,byrow=TRUE)

## compare 'gam' and 'gamm4' this time

b <- gam(y~s(X,by=L,k=20),method="REML")
br <- gamm4(y~s(X,by=L,k=20))
par(mfrow=c(2,1))
plot(b,shade=TRUE);lines(x,f,col=2)
plot(br$gam,shade=TRUE);lines(x,f,col=2)
```

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