

mgcv

tips sheet

Getting Help

What is mgcv ?

mgcv is an R package for fitting generalized additive models (GAMs). That means we can fit models where the predictors are smooth functions of the covariates. Often these smooth functions are splines, but that's not all they can be.

The main functions in mgcv

gam
For fitting GAMs

gamm
For fitting generalized additive mixed models. Can include correlation structures and performance can be better for random effects. You can specify random effects using `lme` syntax.

bam
For fitting big additive models. Includes some special tricks for fitting to large datasets.

ginla
For fitting using a general integrated nested Laplace approximation (like R-INLA).

Useful references

Wood. Generalized Additive Models. An Introduction with R. 2nd ed. CRC Press, 2017

Pedersen, Miller, Simpson and Ross. Hierarchical Generalized Additive Models in Ecology: An Introduction with mgcv . PeerJ (2019). <https://doi.org/10.7717/peerj.6876>

Basic operation of gam

formula=

We can write a model formula in mgcv just as we can when we use `lm` or `glm`, with some additions.

`s()` is the general setup for a smooth.

`te()` allows us to construct an interaction using a tensor product.

Response distribution family=

Binomial	binomial
Normal	gaussian
Gamma	Gamma
Inverse normal	inverse.gaussian
Poisson	poisson
Quasi	quasi
Quasi-binomial	quasibinomial
Quasi-Poisson	quasipoisson
Tweedie	tw/Tweedie
Negative binomial	nb/negbin
Beta	betar
Censored normal	cnorm
Ordered categorical	ocat
Scaled t	scat
Zero inflated Poisson	ziP
Zero inflated Poisson	ziplss
location-scale Cox proportional hazards	cox.ph
Generalized extreme value location-scale Normal	gevlss
location-scale model	gaulss
Multivariate normal	mvn
Gamma location-scale	gammals
Gumbel location-scale	gumbels
Multinomial	multinom
Tweedie location-scale	twlss
General family	gfam

Smoothers

Using the `bs=` argument in `s()`, `te()`, etc. Further details can be found in `?smooth.construct.*.smooth.spec`

Univariate only smoothers

Cubic regression splines `cr`

Cubic regression splines with shrinkage `cs`

Cyclic cubic splines `cc`

B-splines `bs`

P-splines `ps`

Special smoothers

Adaptive smoothers `ad`

Factor-smooth interactions `sz`

Random factor-smooth interactions `fs`

Smoothers in ≥ 1 dimension

Thin plate regression splines `tp`

Thin plate regression splines within shrinkage `ts`

Duchon splines `ds`

Random effects `re`

Markov random fields `mrf`

Gaussian process smooths `gp`

Smoothers in 2 dimensions

Splines on the sphere `sos`

Soap film smoothing `so` (`sw` and `sf`)

Model checking

`gam.check`

Knots and basis complexity

Something

Fitting criterion `method=`

"GCV.Cp"	Generalized cross validation, default
"REML"	REstricted Maximum Likelihood, preferred
"ML"	Maximum Likelihood
"NCV"	Neighbourhood Cross-Validation

Extras

<code>gam.mh</code>	Metropolis-Hastings sampling of the posterior
<code>concurvity</code>	Assess concurvity between terms
<code>gam.vcomp</code>	Random effects style output
<code>gamSim</code>	Simulate GAM-type data
<code>inSide/in.out</code>	point-in-polygon test
<code>jagam</code>	Generate JAGS/Nimble code
<code>new.name</code>	Generate a variable name
<code>place.knots</code>	Place knots evenly
<code>rmvn</code>	Generate multivariate normal deviates

Extra help

<code>?gam.models</code>	Fitting fancy models
<code>?linear.functionals</code>	
<code>?random.effects</code>	
<code>?mgcv.FAQ</code>	frequently asked questions
<code>?mgcv.parallel</code>	Info on parallelisation
<code>?missing.data</code>	
<code>?choose.k</code>	How to select basis size
<code>?one.se.rule</code>	

Other packages

`scam`
`gratia`
`mgcViz`
`qgam`
`gamm4`