

Defining Effect Methods for Other Models

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The **effects** package in R is designed primarily to draw graphs that visualize a fitted response surface of a fitted model in problems with a linear predictor. Many modeling paradigms that can be fit with base R or contributed packages fit into this framework, including methods for linear, multivariate linear, and generalized linear models fit by the standard **lm** and **glm** functions and by the **svyglm** function in the **survey** package (Lumley, 2004); linear models fit by generalized least squares using the **gls** function in the **nlme** package (Pinheiro et al., 2016); multinomial regression models fit by **multinom** in the **nnet** package (Venables and Ripley, 2002); ordinal regression models using **polr** from the **MASS** package (Venables and Ripley, 2002) and **clm** and **clm2** from the **ordinal** package (Christensen, 2015); linear and generalized linear mixed models using the **lme** function in the **nlme** package (Pinheiro et al., 2016) and the **lmer** and **glmer** functions in the **lme4** package (Bates et al., 2015); and latent class models fit by **poLCA** in the **poLCA** package (Linzer and Lewis, 2011). This is hardly an exhaustive list of fitting methods that are based on a linear predictor, and we have been asked from time to time to write functions to use **effects** with this other fitting methods.

The default **Effect.default** may work with some modeling functions, as would objects of the class **gls** that we describe below in Section 1. This will work if your function recognizes the defaults for the arguments in the **sources** list described in Section 1. If the defaults don't work, you will need to create your own **Effect** method or call **Effect.default** with your own value of **sources**.

The **effect** package has five functions that create the information needed for drawing effects plots, **Effect**, **allEffects**, **effect** and **predictorEffect** and **predictorEffects**. To add new modeling to the package only **Effect** needs to be written; the package will take care of all the other functions.

All the functions described below are included in the **effects** package. These can be used as templates for adding methods for other modeling types.

1 Generalized Least Squares (nlme package)

The **gls** function in the **nlme** package (Pinheiro et al., 2018) fits linear models via generalized least squares. A call to **gls** creates an object of class **gls**. The following function will allow usage of such objects with the **effects** package.

```

Effect.gls <- function(focal.predictors, mod, ...){
  args <- list(
    type = "glm",
    call = mod$call,
    formula = formula(mod),
    family = family(mod),
    link = NULL,
    coefficients = coef(mod),
    vcov = as.matrix(vcov(mod)))
  Effect.default(focal.predictors, mod, ..., sources=args)
}

```

This function sets an argument `sources` that is then passed to the default `Effect.default`. The argument `focal.predictors` will be used to pass the focal predictors to other methods. The `mod` argument is the name of the regression object that has been created. The `...` argument allows passing other arguments to the default method.

The argument `sources` to `Effect.default` contains the information needed in a list of up to five arguments:

type The `effects` package has three basic modeling functions: `type = "glm"`, the default, is used for functions with a univariate response and a linear predictor and possibly a link function. This class includes linear models, generalized linear models, robust regression, generalized least squares fitting, linear and generalized linear mixed effects models and many others. The `type = "polr"` is used for ordinal regression models, as in the `polr` function in the `MASS` package, and similar methods in Section 6. The `type = "multinom"` for multinomial log-linear models as fit by the `multinom` function in `nnet`, and to polytomous latent class models created with the `polCA` function in the `polCA` package.

call The `Effect.default` method uses the `call` to harvest additional arguments that it needs, such as `data`, `subset` and `family`. These are used to compute fitted values, and to interpret the terms in the formula for the linear predictor. The default is `mod$call` for S3 objects and `mod@call` for S4 objects.

formula In most cases the formula for the linear predictor is returned by `formula(mod)`, the default, but if this is not the case the value of this argument should be the value of the formula for fixed effects.

family and **link** GLM-like models include a `family` specifying both an error distribution and a link function. If the call `family(mod)` returns a family, you do not need to specify either of these values. The `betareg` function, on the other hand, fixes the error distribution as a Beta distribution, but it does include a link function using a `link` argument. In this case, you must specify the link function; see the `betareg` example in Section 5

below. If you have a family, but it is not returned by `family(mod)`, use the `family` argument to specify it.

At present, non-canonical families like the negative binomial with two parameters in the variance function are not supported. This may be changed in the future.

coefficients In many cases the (fixed-effect) coefficient estimates are returned by `coef(mod)`, the default, but if this is not the case then the value of this argument should be the estimates of the coefficients in the linear predictor. The functions in the **effects** package do not use estimates of random effects.

vcov In many cases the estimated covariance matrix of the (fixed-effect) coefficient estimates is returned by `vcov(mod)`, the default, but if this is not the case then the value of this argument should be the estimated covariance matrix of the (fixed-effect) coefficient estimates in the linear predictor.

Since the values of all the arguments in **sources** are default values for the `gls` function, there is no need to have written the `Effect.gls` method, as the default method would work.

2 Mixed Effects with `lme` (`nlme` package)

The `lme` function in the `nlme` package (Pinheiro et al., 2018) fits linear mixed models. The required function for fitted objects from this function to be used with **effects** functions is

```
Effects.lme <- function(focal.predictors, mod, ...){
  args <- list(
    call = mod$call,
    formula = mod$call$fixed,
    coefficients = mod$coefficients$fixed,
    vcov = mod$varFix)
  Effect.default(focal.predictors, mod, ..., sources=args)
}
```

The `formula`, `coefficients` and `vcov` arguments do not have the default values, but can be obtained from the fitted model `mod` as shown. The `link` entry in **sources** is not used by `lme` and is given its default value of `NULL`. The `call` argument is set to be its default value.

3 Mixed Effects with the `lmer` (`lme4` package)

The `lme4` package (Bates et al., 2015) fits linear and generalized linear mixed effects models with the `lmer` and `glmer` functions, respectively. The same **Effect** function can be used for `lmer` and `glmer` models.

The following method is a little more complicated because it contains an additional argument `KR` to determine if the Kenward-Roger coefficient covariance matrix is to be used to compute effect standard errors. The default is `FALSE` because the computation is very slow. If `KR = TRUE`, the function also checks if the `pbkrtest` package is present.

```
Effect.merMod <- function(focal.predictors, mod, ..., KR=FALSE){
  if (KR && !requireNamespace("pbkrtest", quietly=TRUE)){
    KR <- FALSE
    warning("pbkrtest is not available, KR set to FALSE")
  }
  fam <- family(mod)
  args <- list(
    call = mod@call,
    coefficients = lme4::fixef(mod),
    vcov = if (fam == "gaussian" && fam$link == "identity" && KR)
      as.matrix(pbkrtest::vcovAdj(mod)) else as.matrix(vcov(mod))
  )
  Effect.default(focal.predictors, mod, ..., sources=args)
}
```

Because `lmer` is an S4 object (tested using the `isS4` function), the default for `call` is `mod@call`, and this argument would have been set automatically had we not included it in the above function. The `coefficient` for an object created by a call to `lmer` or `glmer` are not returned by `coef(mod)`, so the value of `coefficients` is the value returned by `lme4::fixef(mod)`. The `vcov` estimate contains its estimated variance covariance matrix of the fixed effects.

The `formula` for a mixed-effects model in the `lme4` package specifies both the linear predictor in the mean function and the linear predictor(s) in the variance functions in terms with parentheses and and vertical bars such as `(1 + age | subject)`. The `effects` code will automatically remove any terms like these in any formula.

4 Robust Linear Mixed Models (robustlmm package)

The `rlmer` function in the `robustlmm` package (Koller, 2016) fits linear mixed models with a robust estimation method. As `rlmer` closely parallels the `lmer` function, an object created by `rlmer` is easily used with `effects`:

```
Effect.rlmerMod <- function(focal.predictors, mod, ...){
  args <- list(
    coefficients = lme4::fixef(mod)
  )
  Effect.default(focal.predictors, mod, ..., sources=args)
}
```

5 Beta Regression

The `betareg` function in the `betareg` package (Grün et al., 2012) fits regressions with a link function but with Beta distributed errors.

```
Effect.betareg <- function(focal.predictors, mod, ...){
  coef <- mod$coefficients$mean
  vco <- vcov(mod)[1:length(coef), 1:length(coef)]
  args <- list(
    call = mod$call,
    formula = formula(mod),
    coefficients = coef,
    link = mod$link$mean,
    vcov = vco)
  Effect.default(focal.predictors, mod, ..., sources=args)
}
```

The Beta distributions require estimation of the parameters of the Beta, but these are not used by `Effect`. The relevant coefficients and covariance matrix are extracted by the first two lines of the function. This method has a link function specified by the `link` argument, but no family, so the `link` argument is added to `sources`.

6 Ordinal Models (ordinal package)

Proportional odds logit and probit regression models fit with the `polr` function in the `MASS` package (Venables and Ripley, 2002) are supported in the `effects` package. The `ordinal` package, (Christensen, 2015) contains three functions that are very similar to `polr`. The `clm` and `clm2` functions allow more link functions and a number of other generalizations. The `clmm` function allows including random effects.

6.1 clm

```
Effect.clm <- function(focal.predictors, mod, ...){
  if (requireNamespace("MASS", quietly=TRUE)){
    polr <- MASS::polr}
  if(mod$link != "logit")
    stop("Effects only supports the logit link")
  if(mod$threshold != "flexible")
    stop("Effects only supports the flexible threshold")
  if(is.null(mod$Hessian)){
    message("\nRe-fitting to get Hessian\n")
    mod <- update(mod, Hess=TRUE)}
  numTheta <- length(mod$Theta)
  numBeta <- length(mod$beta)
```

```

or <- c( (numTheta+1):(numTheta + numBeta), 1:(numTheta))
args <- list(
  type = "polr",
  coefficients = mod$beta,
  vcov = as.matrix(vcov(mod)[or, or]))
Effect.default(focal.predictors, mod, ..., sources=args)
}

```

This function first checks that the MASS package is available. Since the `clm` function allows suppressing the computation of the Hessian, the function checks and computes it if needed to get the estimated covariance matrix. The `clm` function orders the parameters in the order (threshold parameters, linear predictor parameters), so the next few lines identify the elements of `vcov` that are needed by `Effects`. Since the `polr` function does not allow thresholds other than `flexible`, we don't allow them either. Similarly, we have only implemented effects for the default `logit` link.

6.2 `clm2`

```

Effect.clm2 <- function(focal.predictors, mod, ...){
  if (requireNamespace("MASS", quietly=TRUE)){
    polr <- MASS::polr}
  if(is.null(mod$Hessian)){
    message("\nRe-fitting to get Hessian\n")
    mod <- update(mod, Hess=TRUE)}
  if(mod$link != "logistic")
    stop("Effects only supports the logit link")
  if(mod$threshold != "flexible")
    stop("Effects only supports the flexible threshold")
  numTheta <- length(mod$Theta)
  numBeta <- length(mod$beta)
  or <- c( (numTheta+1):(numTheta + numBeta), 1:(numTheta))
  args <- list(
    type = "polr",
    formula = mod$call$location,
    coefficients = mod$beta,
    vcov = as.matrix(vcov(mod)[or, or]))
  Effect.default(focal.predictors, mod, ..., sources=args)
}

```

The syntax for `clm2` is not the same as `clm`, so a separate method is required.

6.3 `clmm`

This function allows for random effects in an ordinal model.

```

Effect.clmm <- function(focal.predictors, mod, ...){
  if (requireNamespace("MASS", quietly=TRUE)){
    polr <- MASS::polr}
  if(is.null(mod$Hessian)){
    message("\nRe-fitting to get Hessian\n")
    mod <- update(mod, Hess=TRUE)}
  if(mod$link != "logit")
    stop("Only the logistic link is supported by Effects")
  if(mod$threshold != "flexible")
    stop("Only threshold='flexible' supported by Effects")
  numTheta <- length(mod$Theta)
  numBeta <- length(mod$beta)
  or <- c( (numTheta+1):(numTheta + numBeta), 1:(numTheta))
  skip <- length(unique(model.frame(mod)[,1])) - 1
  vcov <- matrix(NA, nrow=numBeta + skip, ncol=numBeta + skip)
  sel <- rownames(vcov(mod)) %in% names(mod$beta)
  vcov[1:numBeta, 1:numBeta] <- vcov(mod)[sel, sel]
  args <- list(
    type = "polr",
    formula = fixFormula(as.formula(mod$formula)),
    coefficients = mod$beta,
    vcov = as.matrix(vcov))
  Effect.default(focal.predictors, mod, ..., sources=args)
}

```

Complications here come from getting the right elements of `vcov(mod)` corresponding to the fixed effects.

6.4 Others

The `poLCA` function in the `poLCA` package (Linzer and Lewis, 2011) fits polytomous variable latent class models, which uses the multinomial effects plots.

The `svyglm` function in the `survey` package (Lumley, 2004, 2016) fits generalized linear models using survey weights.

References

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