

Variable	Example	Type of Regression	R function / R function for mixed models	
Continuous	Quality of Life, linear scales	linear	lm()	
			<pre>lmer(), glmmTMB()</pre>	
Binary	Success yes/no	binary logistic	glm(family=binomial)	
			<pre>glmer(*), glmmTMB(*)</pre>	
Binary (weighted)	Success yes/no, with weights	quasi-binary logistic	glm(family=quasibinomial)	
			<pre>glmmPQL(family="quasibinomial")</pre>	
Trials (or proportion of <i>counts</i>)	20 successes out of 30 trials	logistic¹	<pre>glm(cbind(trial, success), family=binomial)</pre>	
			<pre>glmer(*), glmmTMB(*)</pre>	
Count data	Number of usages, counts of events	Poisson	glm(family=poisson)	
			<pre>glmer(*), glmmTMB(*)</pre>	
Count data, with	Number of usages, counts of events (with higher variance than mean of response)	negative binomial	glm.nb()	
excess zeros or overdispersion			<pre>glmer.nb(), glmmTMB(family=nbinom)</pre>	
Count data with very many zeros (inflation)	see count data, but response is modelled as mixture of Bernoulli & Poisson (two sources of zeros)	zero-inflated	zeroinfl()	
			glmmTMB(ziformula, family=poisson)	
Count data, with	Number of usages,	zero-inflated	zeroinfl(dist="negbin")	
very many zeros (inflation) and overdispersion	counts of events (with higher variance than mean of response)	negative binomial	glmmTMB(ziformula, family=nbinom)	
	see count data, but only for positive counts (hurdle component models zero-counts)	hurdle (Poisson)	hurdle()	
Count data, zero- truncated			<pre>glmmTMB(family=truncated_poisson)</pre>	
Count data, zero-	see "Count data, zero- truncated", but with higher variance than mean of response	hurdle (neg. binomial)	vglm(family=posnegbinomial)	
truncated and overdispersion			<pre>glmmTMB(family=truncated_nbinom)</pre>	
Proportion / Ratio (without zero and one)	Percentages, proportions of continuous data	Beta ¹	betareg()	
			<pre>glmmTMB(family=beta)</pre>	
Proportion / Ratio (including zero and one)	Percentages, proportions of continuous data	Beta-Binomial, zero-inflated Beta, ordered Beta ²	<pre>BBreg(), betabin(), ordbetareg(), vglm(family=betabinomial)</pre>	
			<pre>glmmTMB(ziformula, family=beta_family/ betabinomial/ ordbeta), ordbetareg()</pre>	





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Oudinal	Likert scale,	ordinal, pro- portional odds	polr(), clm()
Ordinal	worse/ok/better		<pre>clmm(), mixor(), MCMCglmm()</pre>
Cumulative, multinomial	No natural order of categories, like	cumulative link, multinomial	<pre>multinom(), clm(),bracl(), brmultinom()</pre>
	red/green/blue		<pre>clmm(), mixor(), MCMCglmm()</pre>
Continuous right alrawed	Financial data, reaction	Gamma	glm(family=Gamma)
Continuous, right-skewed	times		<pre>glmer(*), glmmTMB(*)</pre>
(Semi-)Continuous, (right)	Financial data, probably	Tweedie	<pre>glm(family=tweedie), cpglm()</pre>
skewed, probably spike at zero (zero-inflation)	exponential dispersion of variance		cpglmm(),glmmTMB(*)
(Semi-)Continuous,	Normal distribution,	Tobit	<pre>censReg(), tobit()</pre>
skewed, zero-inflation	negative values censored and stacked on zero		semLme()
Continuous, but truncated or outliers		truncated	<pre>censReg(), tobit(), vglm(family=tobit)</pre>
Continuous, but	wildlife populations,	log-transformed, non-linear	<pre>glm(family=Gaussian("log"), nls()</pre>
exponential growth	financial investments		<pre>glmmTMB(*), nlmer()</pre>
Proportion / Ratio with > 2 categories	Biomass partitioning in plants (ratio of leaf, stem and root mass)	Dirichlet	DirichReg()
	Survival-analysis, time until	Cox	coxph()
Time-to-Event	event/death occurs	(proportional hazards)	coxme()

^{*} Indicates same family-option for mixed models as for their non-multilevel (glm) counterparts.

Last update: 2023/09/19



¹ Note that ratios or proportions from *count data*, like **cbind(successes, failures)**, are modelled as logistic regression with **glm(cbind(successes, failures), family=binomial())**, while ratios from *continuous data* (where the response ranges from 0 to 1) are modelled using beta-regression.

² Usually, zero-inflated models are used when 0 or 1 come from a separate process or category. However, when the 0/1 values are most consistent with censoring rather than with a separate category/process, the *ordered beta regression* is probably a better choice (i.e., 0 mean "below detection", not "something qualitatively different happened") (Source: https://twitter.com/bolkerb/status/1577755600808775680)