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measures. These models  
allow the specification  
of both fixed and  
random effects. The  
fixed effects describe  
how the population  
means differ across  
subject characteristics,  
whereas the random  
effects capture the  
variability among  
subjects or other units.

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become very common in  
biology and recent  
developments involve  
the manipulation of the  
variance-covariance  
matrix for random

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effects and residuals. To the best of my knowledge, within the frame of frequentist methods, the only freeware solution in R should be based on the `nlme` package, as the `lmer` package does not easily permit such manipulations.

Fitting 'complex' mixed models with 'nlme'.

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