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The environmental variance model. Fitting mixed models has become very common in biology and recent developments involve the manipulation of the variance-covariance matrix for random effects and

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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'. Example #1 | R ...

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