lme4: past, present, future

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- 1 Past
- 2 Present
 - The package itself
 - The larger context
- 3 Future

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Past

■ nlme:

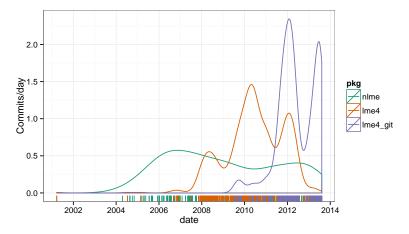
- v. 3.1.1, Nov-1999
- Pinheiro and Bates (2000)
- v. 3.1.50: move to R SVN, Apr-2004
- becomes Recommended 2007?
- current 3.1.109, Apr-2013

■ lme4:

- initial ChangeLog entry, Jul-2002
- v. 0.2.1, Jun-2003
- move to R-forge SVN, Nov-2007
- 1me4a ca. Aug-2009
- $lme4Eigen \rightarrow lme4$: Mar-2012
- move to github, Nov-2012
- v. 1.0-0 Aug-2013 ???

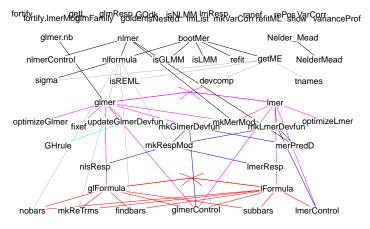


Past 0



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Code structure (mvbutils::foodweb)





Tiers

I low-level computational/matrix representation;
PWRSS solutions
 (Eigen package; Julia)
II low-level computation:
PWRSS iteration/step-halving, Laplace/AGQ
III nonlinear optimization over θ, (β)
IV API; interface, methods, modularization

V downstream packages

Features

- I fill-reducing permutations (speed, for large problems; crossed random effects)
- | GLMMs (step-halving, Laplace, AGQ)
- III fit on constrained scale; default is derivative-free with box constraints (Nelder-Mead, BOBYQA), but flexible
- IV profiling; parametric bootstrap; predict and simulate; deviance function and modularity

Issues

What have we (lme4 authors) been doing?

- fixing glitches and achieving consistency . . .
- ...especially for GLMMs (e.g. updating weights)
- numerical stability of fits (e.g. boundary issues)
- evaluation and environments (e.g. call-modification)

The package itself

Metrics

- 7600 lines of R code, 3270 of C++ code
- ≈ 80 downstream packages (Imports/Suggests/Depends); 433 recursive links (tools::dependsOnPkgs)
- for 9 June 4 August 2013, 4020 downloads from http://probability.ca/cran (rank 44/4866)

Competitors (deterministic)

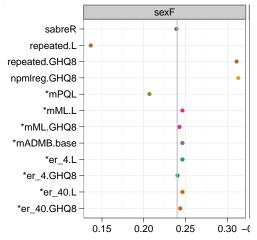
package	features	
glmmADMB*	negative binomial I/II, zero-inflated/hurdle,	
glmmML	non-Normal RE, bootstrap	
npmlreg	non-parametric RE (by EM)	
hglm	hierarchical GLMs	
lmm	(LMMs only)	
${ t repeated}^*$		
sabreR		
MASS::glmmPQL	R-side effects (???)	
_	·	

(*=off-CRAN)

Non-R: AS-REML/Genstat (flexible variance), SAS, MLWiN, HLM, GLAAMM/Stata, . . .

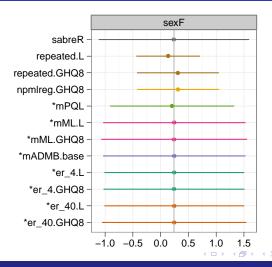


Comparison with Zhang et al. (2011) (1)



The larger context

Comparison Zhang et al. (2011) (2)



The larger context

Competitors (stochastic)

package	features	
bernor*	Monte Carlo expectation-maximization	
MCMCglmm	multivariate/multitype, flexible variance, MCMC	zero-inflated/hurdle,
INLA*		
Non-R: BUGS/JAGS/Stan (and R2*)		



The larger context

Environment

- CRAN administrators
 - C++/Rcpp/RcppEigen issues
- Infrastructure maintainers (reference classes, Rcpp)
- downstream package maintainers
 - variation in coding/packaging style, statistical philosophy
 - what should core 1me4 provide?
- users (r-sig-mixed-models, Stack Overflow/Cross Validated)
 - variation in sophistication/needs
 - latent demand?



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Near future

- cleanup and tweaks, e.g.
 - finish neg binom models (IV)
 - better input/output tests (III, IV)
 - low-hanging parallelization
- fix GLMM edge cases (PWRSS failures) (II)
- evaluate/improve performance, esp. on large data (I)
- write JSS paper(s)



Wishlist

- R-side effects (IV)
- flexible RE variance (e.g. spatial/phylogenetic GLMMs) (IV)
- speed optimization (I)
- working MCMC sampler (IV)
- AGQ, postVar for complex models (II, IV)
- improve bootMer (merBoot features) (IV)
- improve GLMMs that use estimated scale parameters (III, IV)
- lightweight stored models (IV)
- simulate/predict with newdata, newpar(a)ms
- systematic (G)LMM comparison across packages/settings



Zhang, H., Lu, N., et al., 2011. Statistics in Medicine. ISSN 1097-0258. doi:10.1002/sim.4265.

Past