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Comparing Optimization Algorithms in the Fitting of Linear Mixed Models: Evaluating Speed and Accuracy using lme4 in R and lmm in Julia

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Abstract

The **Timings** package allows for the comparison of several optimizers, in both R and Julia, used in the fitting of various linear mixed models. In R the optimizers are called by lmer from the **lme4** package (version 1.1-8). In Julia the optimizers are called by lmm from **MixedModels** package. From the **Timings** package, conclusions regarding an optimizers relative speed, accuracy and general effectiveness of different optimizers paired with different types of models (ranging from simple to complex) can easily be drawn and interpretted.

There are differences in the model formulations in **lme4** and in **MixedModels**. The numerical representation of the model in **lme4** and the method of evaluating the optimizers, described in this paper, is the same for all models. In **MixedModels** there are specialized representations for some model forms, such as models with a single grouping factor for the random effects. Some of the specialized representations allow for evaluation of the gradient of the objects, which can enhance convergence (but, interestingly, sometimes can impede convergence).

Keywords: optimizers, mixed models, linear mixed models, lme4, lmm, R, Julia.

1. Introduction

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2. Methods

To provide consistency we have copied all the data sets used in the timings to the **Timings** package itself. We have done all timings on the same computer. This computer has a relatively recent Intel processor and we used the Intel Math Kernel Library (MKL) with Julia. We attempted to use Revolution R Open (RRO) as the R implementation as it can be configured with **MKL**. However, we ran into version problems with this so we used the standard Ubuntu version of R linked against OpenBLAS, which is also multi-threaded.

Variables were renamed in the pattern:

- Y the response
- A, B, ... categorical covariates
- G,H, I, ... grouping factors for random effects
- \bullet U, V, ... (skipping Y) continuous covariates

The timing results are saved in JSON (JavaScript Object Notation) files in the directory accessible as

```
> system.file("JSON",package="Timings")
```

within **R**. The directory name will end with ./Timings/inst/JSON/ in the package source directory, for example the result of cloning the github repository. There is one .json file for each data set. Each such file contains results on timings of one or more models.

The **Timings** package for R provides a **retime** function that takes the name of one of these JSON files and, optionally, the name of a file with the updated timings. Similarly there are some source files for **Julia** retimings.

```
> include("../julia/retime.jl")
> retime("../JSON/Alfalfa.json","/tmp/Alfalfa.json")
> retime("../JSON/Alfalfa.json","/tmp/Alfalfa.json")
```

INCLUDE SUMMARY TABLE USING res.rda

The timing was repeated so that compilation time is not included in the results. This repetition is only needed once per session.

A careful examination of these results shows that the main differences in the **Julia** timings (the **R** timings are merely reported, not evaluated) are that the LN_BOBYQA and LD_MMA optimizers

are much faster in the second run. This is because much of the code needs to be compiled the first time that a derivative-free optimizer and a derivative-based optimizer are used.

The names of the optimizers used with lmm are those from the **NLopt** package for Julia. Names that begin with LD_ are gradient-based methods. Names that begin with LN_ are derivative-free methods. There is one other derivative-free method, LN_PRAXIS, available in the **NLopt** package but, for some reason, it can hang on very simple problems like this. Frequently we omit it.

The optimizers used with lmer include the Nelder_Mead optimizer built into the lme4 package, the bobyqa optimizer from the minqa package, the derivative-free optimizers from the nloptr package and several optimizers from the optimx package.

The optimx:bobyqa optimizer is just a wrapper around bobyqa (bounded optimization by quadratic approximation) from the minqa package and should provide results similar to those from the bobyqa optimizer. For some reason the number of function evaluations is not reported for the version in optimx.

The optimizers from **nloptr** (i.e. those whose names begin with NLOPT_LN_) use the same underlying code as do the similarly named optimizers in the **NLopt** package for Julia. The number of iterations to convergence should be similar for the same underlying code, although not nessarily exactly the same because the evaluation of the objective in R and in Julia may produce slightly different answers. Also the convergence criteria in the Julia version are more strict than those in the R version

Also shown are the value of the criterion (negative twice the log-likelihood, lower is better) achieved, the elapsed time and the number of function and gradient evaluations. The nopt value is the number of parameters in the optimization problem. mtype is the model type in the Julia code. There are special methods for solving the penalized least squares (PLS) problem, and for evaluating the objective and its gradient when there is only one grouping factor for the random effects. The model type is called PLSOne.

The Alfalfa example is a particularly easy one and all of the optimizerws converge to an objective value close to -10.81023 in less than 0.6 seconds.

3. Results

For the Alfalfa data there is not much of a burden in refitting the model with all the Julia optimizers just to get the table shown above. But other examples can take an hour or more to converge and we don't really need to refit them every time. The tabulate.jl file contains a function optdir to create a DataFrame from the results of all the model fits.

```
> include("../julia/tabulate.jl")
> res = optdir("../JSON");
\
```

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25	"LD_CCSAQ" "mm0"	69588	-	6	١	0.0	1	4.8286		1.1034
26	"LD_CCSAQ" "Oxboys"	234	-	3	١	136.788	1	0.0169		5.3039
27	"LD_CCSAQ" "PBIB"	60	-	1	١	0.0	1	0.0014		8.6177
28	"LD_CCSAQ" "Penicillin"	144	1	2	١	0.0	1	0.0131	I	15.3763
29	"LD_CCSAQ" "Semiconductor"	48	-	1	١	0.0	1	0.0012		1.133
30	"LD_CCSAQ" "SIMS"	3691	-	3	١	3.60856	1	0.134		2.3154
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3.1. Proportion Converged

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3.2. Reliability

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4. Conclusions

5. References

Note all optimization packages used, lmm, lme4, jsonlite,

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