KU LEUVEN

Extending Idefix package

Intermediate presentation

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0 Outline

- Introduction
- 2 Modified Fedorov Algorithm
- 3 Kullback-Leibler criterion
- 4 Coordinate Exchange Algorithm
- 6 Simulation study

Outline

- Introduction

1 What is *idefix* R package for?

- ► To create optimal designs for discrete choice experiments (DCEs) based on the multinomial logit model (MNL) and
- Individually adapted designs for the mixed multinomial logit model (MIXL).
- Available on CRAN (v 0.3.3).

Discrete choice experiments

DCEs are composed by:

- Nominal or Ordinal response variable
- Choice sets
- Alternatives within each choice set
- Attributes and levels

1 Multinomial Logit Model

- ► Choice design matrix $\mathbf{X} = [\mathbf{x}_{js}']$, where \mathbf{x}_{js} is a $k \times 1$ vector of attributes for profile j in choice set s.
- ▶ Respondent's utility is $u_{js} = \mathbf{x}_{js}' \boldsymbol{\beta} + \epsilon_{js}$, where $\boldsymbol{\beta}$ is a vector of parameters and ϵ_{js} is an i.i.d. extreme value error term.
- lacktriangle Probability a respondent chooses alternative j in choice set s is

$$p_{js} = \frac{e^{\mathbf{x}'_{js}\beta}}{\sum_{t=1}^{J} e^{\mathbf{x}'_{ts}\beta}}$$

Information Matrix is

$$\mathbf{M}(\mathbf{X}, \boldsymbol{\beta}) = N \sum_{s=1}^{S} \mathbf{X}_{s}' (\mathbf{P}_{s} - \mathbf{p}_{s} \mathbf{p}_{s}') \mathbf{X}_{s}$$

where \mathbf{X}_s is the design matrix of choice set s, $\mathbf{p}_s = [p_{1s}, \cdots, p_{Js}]$ and $\mathbf{P}_s = diag[p_{1s}, \cdots, p_{Js}]$ and N is the number of respondents.

1 D-optimality

- ▶ In OLS is defined as $D = |\mathbf{X}'\mathbf{X}|$
- \blacktriangleright In MNL is defined adopting the prior distribution of $oldsymbol{eta}$

$$D_B = \int_{\mathcal{R}^k} \left\{ \det \left(\mathbf{M}^{-1}(\mathbf{X}, \boldsymbol{\beta}) \right) \right\}^{1/k} \pi(\boldsymbol{\beta}) d\boldsymbol{\beta}$$

Where k is the number of unknown parameters in the model and $\pi(\beta)$ is the prior distribution of β . This criterion is also called Bayesian D-optimality criterion or just D_B .

1 Mixed Multinomial Logit model

- MNL models assume that the respondents have the same preferences, β , for the attributes studied in the experiment.
- MIXL models assume that the individual preferences, β_n , follow a certain distribution across respondents $(\beta_n \sim f(\mu_\beta, \sigma_\beta))$.
 - ullet Probability a respondent chooses alternative j in choice set s is

$$p_{js}^* = \int p_{js}(\boldsymbol{\beta}) f(\boldsymbol{\beta}) d\beta$$

Where $p_{js}(\beta)$ is defined as in the MNL model.

MIXL model assumes that respondents choose according to an MNL model, but each with different preferences.

1 Individually Adapted designs

The proper name of the methodology is *Individually adapted* sequential Bayesian design. It consists in two stages:

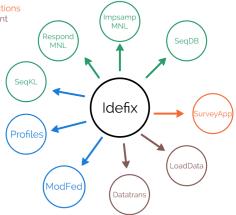
- ▶ **Initial static stage:** use a common initial prior distribution $\pi(\beta)$ for all respondents. It is used to generate an initial design.
- ▶ Adaptative sequential stage: the prior information is updated sequentially after each response, and each choice set is constructed using the updated prior. Therefore, each respondent will have a different design.

Notes:

- Any criterion can be used to select a new choice set.
- ▶ Different algorithms can be used to select a new choice set. Here the Modified Fedorov Algorithm is used.

1 Current state of the package

- In Blue MNL functions
- In Green MIXL functions
- In Orange real survey functions
- In Brown Data management functions



1 **Objectives**

- Improve processing time of the **Modified Fedorov algorithm** by implementing some parts of the algorithm in C++.
- Implement the **KL** criterion and compare it with the function that is already available in the package.
- Implement the Coordinate Exchange algorithm to create optimal designs.
- Make a **simulation study** to compare processing times and optimality of designs between the Modified Fedorov algorithm, the Coordinate Exchange algorithm and the use of DB and KL criteria.
- Reorganize some functions inside the package, remove possible redundancy in code and implement parts of the code in C++.

Outline

- 2 Modified Fedorov Algorithm

2 Modified Fedorov Algorithm

- Point exchange algorithm
- Algorithm steps:
 - 1 Create a random initial design from candidate set.
 - 2 For each row of this design
 - 1 Exchange this row with each row from the candidate set. Resulting in N different designs.
 - 2 Compute optimality criterion for each modified design and choose the best value.
 - 3 update initial design and start again with the following row
 - 3 Repeat this process until no differences are found between the initial design and the final.

2 Modified Fedorov Algorithm

Example

- ▶ Design $3^3/2/8 \Rightarrow$ Design matrix 16×6 (dummy coding).
- ightharpoonup Candidate set has $3^3 = 27$ rows/profiles
- In each iteration $16 \times 27 = 432$ information matrices and determinants are computed to find the best optimal design. Assume that only one iteration is needed.
- ▶ But, the information matrix needs draws from the prior of β . Assuming just 10 draws, the number of information matrices and determinants is 4320.
- But, different random initial designs to avoid local optima. Assuming 10 initial designs, the final number of information matrices and determinants is 43200.

First activity: Improve processing time in individually adapted designs.

SeqDB function selects the next DB-efficient choice set given parameter values and an initial design.

Example

Considering $3^3/2/8$ design:



How to make it faster?

Using Hadley Wickham approach in his book Advanced R:

- 1 Find the biggest bottleneck (the slowest part of the code).
- 2 Try to eliminate it (you may not succeed but that is ok).
- 3 Repeat until your code is **fast enough**.

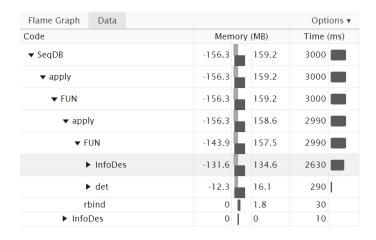
But, how to make it faster?

- Using faster functions in R and avoiding loops using vectorized functions.
- ▶ Implementing parts of the code in C++.

Find the biggest bottleneck

- ▶ $4 \times 3 \times 2/2/8$ design.
- ▶ 10 draws from β distribution.
- Pre-defined initial design and alternatives chosen (1st stage of IASB approach).

Profiling SeqDB function



Implementation in C++

- Use of Rcpp package
- ▶ Use of Rcpp Armadillo: C++ linear algebra library

Implementation in C++

- ► Use of Rcpp package
- Use of Rcpp Armadillo: C++ linear algebra library

```
InfoDes <- function(par, des, n.alts) {
  group <- rep(seq(1, nrow(des) / n.alts, 1), each = n.alts)
  # probability
  u <- des %*% diag(par)
  u <- .rowSums(u, m = nrow(des), n = length(par))
  p <- exp(u) / rep(rowsum(exp(u), group), each = n.alts)
  # information matrix
  info. des <- crossprod(des * p, des) - crossprod(rowsum( des * p, group))
  return(info.des)
}</pre>
```

10 lines of code

Implementation in C++

- Use of Rcpp package
- Use of Rcpp Armadillo: C++ linear algebra library

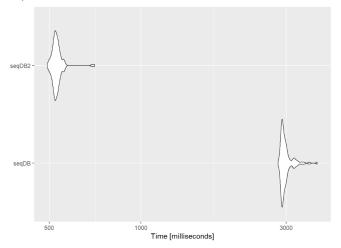
```
InfoDes <- function(par, des, n.alts)</pre>
  group \leftarrow rep(seq(1, nrow(des) / n.alts, 1), each = n.alts)
  # probability
  u <- des %*% diag(par)
  u <- .rowSums(u, m = nrow(des), n = length(par))
  p <- exp(u) / rep(rowsum(exp(u), group), each = n.alts)</pre>
  # information matrix
  info.des <- crossprod(des * p. des) - crossprod(rowsum( des * p. group))
  return(info.des)
       [[Rcpp::depends(RcppArmadillo)]]
    # include <RcppArmadillo.h>
```

```
using namespace Rcpp;
// [[Rcpp::export]]
NumericMatrix InfoDes_cpp(NumericVector par. NumericMatrix des.
                              double n_alts) {
  int i = 0:
  NumericVector group(des.nrow());
```

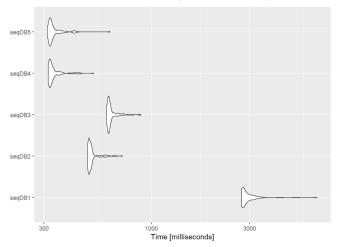
10 lines of code

> 117 of

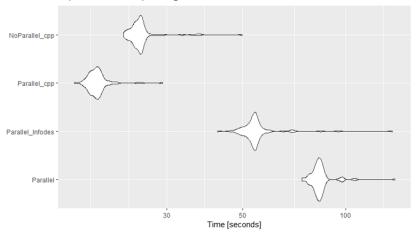
Result: Implementation in C++ is almost 6x faster.



Find next bottleneck and improve it (multiple times)



What about parallel computing?



Second activity: Improve processing time in MNL designs

- The hardest work had already been done
 - Find bottlenecks
 - Improve functions in C++
- ModFed processing time was also improved by using the same functions as in SeqDB.

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- ► The hardest work had already been done
 - Find bottlenecks
 - Improve functions in C++
- ModFed processing time was also improved by using the same functions as in SeqDB.

What was needed to complete this task with success?

- Understand how Modified Fedorov algorithm works.
- Learn version control (Git and GitHub).
- ▶ Learn how C++ and C++ armadillo work in R.
- Learn about benchmarking and parallel computing.

Outline

- Kullback-Leibler criterion

- It was developed under individually adapted designs for the MIXL.
- It is an alternative to D-optimal criterion.
- It is faster to compute and it provides equally efficient designs.
- It is based on the Kullback-Leibler information:

$$KL(f,g) = \int f(x)log\frac{f(x)}{g(x)}dx$$

Where f and g are continuous densities of X.

- KL is non-negative or zero (f(x) = g(x))
- KL increases as the densities become more divergent
- KL is not symmetric, $KL(f,g) \neq KL(g,f)$

Implementation in DCEs

- ▶ To select next choice set, maximize the KL between the current posterior of β and the updated posterior one can obtain with the additional response from the next choice set.
- Since there are multiple alternatives, the expectation over all possible choices is maximized.

$$KLP = \sum_{j=1}^{J} \pi(y_{jsn}|\mathbf{y}_n^{s-1})KL\left[f(\boldsymbol{\beta})_n|\mathbf{y}_n^{s-1}), f(\boldsymbol{\beta})_n|\mathbf{y}_n^{s-1}, y_{jsn})\right]$$

Where s is the next choice set, n is a particular respondent and j is the chosen alternative. The densities $f(\pmb{\beta})_n|\mathbf{y}_n^{s-1})$ and $f(\pmb{\beta})_n|\mathbf{y}_n^{s-1},y_{jsn})$ are the updated posteriors and $\pi(y_{jsn}|\mathbf{y}_n^{s-1})$ is the posterior weighted choice probabilities for the alternatives in the choice set s, given the previous responses.

Implementation in the package

ightharpoonup Applying KL definition, KLP can be written as

$$KLP = \sum_{j=1}^{J} \pi(y_{jsn}|\mathbf{y}_n^{s-1})[\log \pi(y_{jsn}|\mathbf{y}_n^{s-1}) - \sum_{j=1}^{J} \pi(y_{jsn}|\mathbf{y}_n^{s-1})]$$

$$\int \log p_{jsn}(\boldsymbol{\beta}_n) f(\boldsymbol{\beta}_n | \mathbf{y}_n^{s-1} d\boldsymbol{\beta}_n)]$$

- lackbox Modified Fedorov algorithm is used, but instead of using D-optimality criterion KLP is used.
- Simulations in R are not consistent with results obtained in the paper that proposed the criterion.

Third activity: Check why SeqKL function is not working

- Check code of simulations done in the paper that proposed the criterion.
 - Made in SAS. Proc IML.
 - 490 lines of code. No comments, no indentation.
- Check code of implementation in R
- Comparison of results from each function in both implementations.
- List differences.
- Discussion.

4 Outline

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- 3 Kullback-Leibler criterion
- 4 Coordinate Exchange Algorithm
- Simulation study

4 Coordinate Exchange Algorithm

- Coordinate exchange algorithm
 - Point exchange algorithm: Compute optimality criterion $\prod_{j=1}^{J} l_j$ times for each row.
 - Here: Compute optimality criterion $\sum_{j=1}^{J} l_j$ times for each row.
- Algorithm steps:
 - 1 Create a random initial design
 - 2 For each row of this design
 - 1 Take the first attribute in the row, evaluate the optimality criterion over all the levels of that attribute.
 - 2 If the optimality criterion of any of these levels is better than the current, then it is replaced.
 - 3 Repeat with the remaining attributes in the row.
 - 3 Repeat this process until no differences are found between the initial design and the final.

4 **Coordinate Exchange Algorithm**

Example

- ▶ Design $3^3/2/8 \Rightarrow$ Design matrix 16×6 (dummy coding).
- No Candidate set is needed.
- In each iteration $16 \times 9 = 144$ information matrices and determinants are computed to find the best optimal design. Assume that only one iteration is needed.
- ▶ But, the information matrix needs draws from the prior of β . Assuming just 10 draws, the number of information matrices and determinants is 1440.
- But, different random initial designs to avoid local optima. Assuming 10 initial designs, the final number of information matrices and determinants is 14400.

As a reminder, in Modified Fedorov the final number was 43200.

Coordinate Exchange Algorithm

Fourth activity: Implement the Coordinate Exchange algorithm

- Implementation with only categorical factors/attributes.
- Implementation with continuous attributes.
- 3 Implementation with both categorical and continuous.
- Improve processing time, if possible (parallel computing, C++).

Note:

D-optimality criterion is going to be used, so the implementation of the information matrix in C++ is also going to be used here.

Outline

- Simulation study

5 Simulation study

- ▶ The idea is to compare the processing time of Modified Fedorov algorithm and the Coordinate Exchange algorithm.
 - Determine the scenarios where one outperforms the other.
 - Determine in which situations parallel computing is needed.
- \triangleright Compare efficiency of designs found with D-optimality criterion and KL criterion
 - Determine the scenarios where one outperforms the other.

5 **Objectives**

- Improve processing time of the **Modified Fedorov algorithm** by implementing some parts of the algorithm in C++. \checkmark
- Implement the **KL** criterion and compare it with the function that is already available in the package. \checkmark
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5 **Optimality criteria**

- \triangleright To obtain precise estimates of β
 - D-optimality: minimize the determinant of the variance-covariance matrix of β
 - A—optimality: minimize the trace of the variance-covariance matrix of β
- ► To obtain precise response predictions
 - G-optimality: minimize the maximum prediction variance
 - V—optimality: minimize the average prediction variance

Note:

These criteria are based on the information matrix, which depends on the unknown values in β through the probabilities p_{is} . Therefore, a Bayesian strategy that integrates the design criteria over a prior parameter distribution $\pi(\beta)$ is adopted. Usually, the prior is a multivariate normal distribution.