

Computer Lab 6

Computational Statistics

Linköpings Universitet, IDA, Statistik

2017/03/14

Kurskod och namn:	732A90 Computational Statistics
Datum:	2017/03/06–2017/03/26
Delmomentsansvarig:	Krzysztof Bartoszek
Instruktioner:	<p>This computer laboratory is part of the examination for the Computational Statistics course</p> <p>Create a group report on the solutions to the lab as a .PDF file.</p> <p>Be concise and do not include unnecessary printouts and figures produced by the software and not required in the assignments.</p> <p>All R code should be included as an appendix into your report.</p> <p>A typical lab report should 2–4 pages of text plus some amount of figures plus appendix with codes.</p> <p>In the report reference ALL consulted sources and disclose ALL collaborations.</p> <p>The report should be handed in via LISAM (or alternatively in case of problems e-mailed to krzysztof.bartoszek@liu.se), by 23:59 26 March 2017 at latest.</p> <p>The report can be written in English or Swedish.</p> <p>There is no seminar related to this Lab.</p>

Question 1: Genetic algorithm

In this assignment, you will try to perform one-dimensional maximization with the help of a genetic algorithm.

1. Define the function

$$f(x) := \frac{x^2}{e^x} - 2 \exp(-(9 \sin x)/(x^2 + x + 1))$$

2. Define the function `crossover()`: for two scalars x and y it returns their “kid” as $(x+y)/2$.

3. Define the function `mutate()` that for a scalar x returns the result of the integer division $x^2 \bmod 30$. (Operation `mod` is denoted in R as `%%`).
4. Write a function that depends on the parameters `maxiter` and `mutprob` and:
 - (a) Plots function f in the range from 0 to 30. Do you see any maximum value?
 - (b) Defines an initial population for the genetic algorithm as $X = (0, 5, 10, 15, \dots, 30)$.
 - (c) Computes vector `Values` that contains the function values for each population point.
 - (d) Performs `maxiter` iterations where at each iteration
 - i. Two indexes are randomly sampled from the current population, they are further used as parents (use `sample()`).
 - ii. One index with the smallest objective function is selected from the current population, the point is referred to as victim (use `order()`).
 - iii. Parents are used to produce a new kid by crossover. Mutate this kid with probability `mutprob` (use `crossover()`, `mutate()`).
 - iv. The victim is replaced by the kid in the population and the vector `Values` is updated.
 - v. The current maximal value of the objective function is saved.
 - (e) Add the final observations to the current plot in another colour.
5. Run your code with different combinations of `maxiter`= 10, 100 and `mutprob`= 0.1, 0.5, 0.9. Observe the initial population and final population. Conclusions?

Question 2: EM algorithm

The data file `physical.csv` describes a behavior of two related physical processes $Y = Y(X)$ and $Z = Z(X)$.

1. Make a time series plot describing dependence of Z and Y versus X . Does it seem that two processes are related to each other? What can you say about the variation of the response values with respect to X ?
2. Note that there are some missing values of Z in the data which implies problems in estimating models by maximum likelihood. Use the following model

$$Y_i \sim \exp(X_i/\lambda), \quad Z_i \sim \exp(X_i/(2\lambda))$$

where λ is some unknown parameter.

The goal is to derive an EM algorithm that estimates λ .

3. Implement this algorithm in R, use $\lambda_0 = 100$ and convergence criterion “stop if the change in λ is less than 0.001”. What is the optimal λ and how many iterations were required to compute it?
4. Plot $E[Y]$ and $E[Z]$ versus X in the same plot as Y and Z versus X . Comment whether the computed λ seems to be reasonable.