Bayesian Estimation of Monotone Single Index Models

Description and Intended Functionality

This package intends to perform a Markov Chain Monte Carlo algorithm to estimate a monotone Single index models of the form:

$$y = g(x^{\top}\beta) + \epsilon$$

where y: response, x: p-dimensional predictors, β : p-dimensional coefficient vector, ϵ : independent N(0, σ_{ϵ}^2) random errors, g: unknown monotone link function. To construct a prior on g, we consider the basis expansion: $g(x) = \sum_{l=0}^{L} \xi_l \, \psi_l(x)$, where $\psi_l(x) = \int_{-1}^{x} h_l(t) \, \mathrm{d}t$, h_l is a B-Spline basis function of order 2. Then, g is monotonically increasing if and only if $\xi_l \geq 0 \, \forall \, l$, thereby enforcing the monotonicity constraint in an equivalent way.

The monotoneSIM function rescales the covariates and β so that $|x^{\top}\beta| \leq 1$. Then an MCMC algorithm is performed to generate samples from the conditional posteriors of the unknown parameters, ξ , β , σ_{ϵ}^2 . The obtained sample of β is back-scaled to get β corresponding to the original covariates. The function also returns the estimated monotone function g for a grid of x values based on ξ obtained from the algorithm.

Installation

To install this package from Github, run the following in your R console:

devtools::install github("das-snigdha/monotoneSIM")

To Do

The following tasks shall be performed during the remaining of the semester:

- Perform Compatibility Checks on user supplied input.
- Select equispaced knots, u_0, u_1, \ldots, u_L between -1 and 1, if not supplied.
- Select grid of x values, grid.x between -1 and 1 of length size.grid.x, if not supplied
- Perform scaling of covariate matrix X and supplied beta.init.
- Perform back-scaling on obtained beta, to get coefficients corresponding to unscaled X.
- Create a vignette to demonstrate the usage, if time permits.