

AFSSEN

Adaptive Function-on-Scalar Smoothing Elastic Net

AFSSEN is a methodology that simultaneously select important variables and produce smooth estimates of their parameters in a function-on-scalar linear model with sub-Gaussian errors and high-dimensional predictors.

AFSSEN.R

We have option to control sparsity and smoothness separately with using two penalty parameters λ_H and λ_K . We aim to estimate a smooth version of β to minimize the following target function.

$$L_\lambda(\beta) = \frac{1}{2N} \|\mathbf{Y} - \mathbf{X}\beta\|_{\mathbb{H}}^2 + \frac{\lambda_K}{2} \sum_{i=1}^I \|\mathbf{L}(\beta_i)\|_{\mathbb{K}}^2 + \lambda_H \sum_{i=1}^I \tilde{\mathbf{w}}_i \|\beta_i\|_{\mathbb{H}}$$

The following **AFSSEN()** function helps us to estimate the smooth β and find the significant predictors:

```
AFSSEN<- function(X,Y, T_domain = seq(0, 1, length = 50),
  # X (a N*I numerical matrix) N = #observations I = #all_predictors
  # Y = Y_ful=XB+E (a N*m matrix of pointwise evaluation!) m = #time_points
  type_kernel="exponential",param_kernel=8,
  thres=0.02, number_non_zeros=20,
  ratio_lambda_H=0.01, number_lambda_H=100, num_lambda_H_NONad=50,
  lambda_H, lambda_K,
  early_CV=0, early_CV_thres=0.001, max_ite_nadp=10,
  max_ite_adp=30, max_ite_final=50, target_inc=1,
  proportion_training_set=0.75, verbose=FALSE, fold_ad=10)
```

Figure 1:

and here is the functionality of each parameters:

Parameter	functionality	Details
X	Numerical design matrix	It should be a N*I matrix (N= #observations ; I =#predictors)
Y	Matrix of pointwise evaluation for observations on T_domain	It should be a N*m matrix where
T_domain	Time domain for evaluation of Y and generating kernel	Default : T_domain = seq(0,1,m=50)
type_kernel	Type of kernel	'exponential', 'gaussian', 'sobolev'
param_kernel	Kernel parameter	In all types, the time domain is seq(0,1,50)
thres	Stopping criteria: β increment threshold $\ \beta^{\mathbf{T}} - \beta^{\mathbf{T}-1}\ _{\mathbb{H}} < \mathbf{thres}$	
number_non_zeros	Stopping Criteria: Kill switch ; number of nonzero predictors	
ratio_lambda_H	$\frac{\lambda_{Hmax}}{\lambda_{Hmin}}$	
number_lambda_H	Generate number of log-equally spaced in $[\lambda_{Hmin}, \lambda_{Hmax}]$	
num_lambda_H_NONad	Number of λ_H in non-adaptive step	

Parameter	functionality	Details
lambda_H	You have option to insert a vector of λ_H	If you want to make the log-equally spaced by ratio_lambda_H and number_lambda_H , set lambda_H=numeric()
lambda_K	Vector of λ_K	
early_CV	0 or 1 : applying the “early_CV_thres” stopping criteria or not	
early_CV_thres	Stopping Criteria: Breaking point in CV plot	$\frac{\ CV(h-1, k) - CV(h, k)\ }{CV(h-1, k)} < \text{early_CV_thres}$
max_ite_nadp	Stopping Criteria: Maximum iteration of coordinate descent alg. in non-adaptive step	
max_ite_adp	Stopping Criteria: Maximum iteration of coordinate descent alg. in adaptive step	
max_ite_final	Stopping Criteria: Maximum iteration of coordinate descent algorithm for the optimum λ_H and λ_K	
target_inc	Stopping Criteria: 0 or 1 : if target function is increased, stop!	
proportion_training_set	Proportion of training set for estimation in non-adaptive step	
fold_ad	Number of fold for using CV in adaptive steps to find optimum λ_H and λ_K and then estimation	