

README: Implementation for “*Addressing both variable selection and misclassified responses with parametric and semiparametric methods*”

Hui Guo, Grace Y. Yi*, and Boyu Wang

Overview

This repository contains the implementation of the methods proposed in the manuscript submitted to *Bernoulli*, titled:

“*Addressing both variable selection and misclassified responses with parametric and semiparametric methods*”

by Hui Guo, Grace Y. Yi *, and Boyu Wang.

The code includes both parametric and semiparametric methods for addressing response misclassification and performing variable selection in binary classification settings.

Directory Structure

- `train/`: Core implementations
 - `param.py`: Parametric method implementation
 - `semi.py`: Semiparametric method implementation
 - `path_following.py`: Approximate path-following (APF) method from Liu and Zhang (2014)
 - `test.py`: Evaluation script
 - `utils.py`: Supporting utilities
- `run.py`: Entry point for running the full method pipeline.
- `example.ipynb`: Jupyter notebook demonstrating:
 - Synthetic data simulation
 - End-to-end usage of the proposed methods

Usage

You can invoke the main functionality via `run.py`, specifying inputs and configuration arguments as needed.

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Required Data Inputs

- **Z**: Main-study covariate matrix.
 - `numpy` array of shape (n, p)
- **Y_star**: Noisy binary responses in main-study data.
 - `numpy` array of shape $(n,)$
- **Z_val**: Covariates in validation data.
 - `numpy` array of shape (n_v, p)
- **Y_val**: True labels in validation data.
 - `numpy` array of shape $(n_v,)$
- **Y_star_val**: Noisy labels in validation data.
 - `numpy` array of shape $(n_v,)$
- **discrete_idx**: List of indices for discrete features.
 - `list`
 - Each element takes values in $\{0, 1, \dots, p - 1\}$
 - default: `[]`
- **Z_test**: Covariates in test data. Optional.
 - `numpy` array of shape (n_{test}, p)
 - default: `None`
- **Y_test**: True labels in test data. Optional.
 - `numpy` array of shape $(n_{test},)$
 - default: `None`
- **test**: Boolean flag indicating whether to test.
 - `bool`
 - `True` or `False`
 - default: `False`

Model Setup

- **link_func**: Link function.
 - `str`
 - `'logit'` or `'probit'`
 - default: `'logit'`
- **penalty**: Penalty type.
 - `str`
 - `'l1'`, `'scad'`, or `'mcp'`
 - default: `'scad'`
- **use_intercept**: Boolean flag indicating whether to include intercept.

- `bool`
 - `True` or `False`
 - default: `True`
- **criterion**: Model selection criterion.
 - `str`
 - `'gcv'` or `'bic'`
 - default: `'gcv'`
- **model_running**: Method type.
 - `str`
 - `'param'` or `'semi'`
 - default: `'semi'`
- **densityType**: Density estimation method (for semiparametric method).
 - `str`
 - `'Kernel'` and `'pcaKernel'`
 - default: `'pcaKernel'`

Hyperparameters

- **eta**: Decreasing coefficient for the sequence of regularization parameters.
 - `float` in $[0.9, 1)$
 - default: `0.91`
- **R**: Projection radius.
 - positive `float`
 - default: `0.91`
 - If set to `None`, the algorithm computes it automatically
- **L**: Initial learning rate.
 - small positive `float`
 - default: `0.05`
- **N_iter**: Number of iterations (outer loop).
 - `int`
 - default: `5`
- **max_loop**: Maximum number of loops.
 - `int`
 - default: `20`