# Package 'SAMGEP'

January 20, 2025

Type Package

**Title** A Semi-Supervised Method for Prediction of Phenotype Event Times **Version** 0.1.0-1

**Description** A novel semi-supervised machine learning algorithm to predict phenotype event times using Electronic Health Record (EHR) data.

URL https://github.com/celehs/SAMGEP

BugReports https://github.com/celehs/SAMGEP/issues

License GPL-3

Encoding UTF-8

RoxygenNote 7.1.1

**Depends** R (>= 3.5.0)

**Imports** stats, mvtnorm, nlme, pROC, abind, nloptr, foreach, doParallel, parallel, Rcpp

LinkingTo Rcpp, RcppArmadillo

Suggests knitr, rmarkdown

VignetteBuilder knitr

LazyData true

NeedsCompilation yes

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**Repository** CRAN

Date/Publication 2021-01-06 10:00:02 UTC

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SAMGEP-package

SAMGEP: A Semi-supervised Method for Prediction of Phenotype Event Times Using the Electronic Health Record

#### Description

Semi-supervised Adaptive Markov Gaussian Embedding Process (SAMGEP) is a novel semi-supervised machine learning algorithm to predict phenotype event times using Electronic Health Record (EHR) data.

samgep

Semi-supervised Adaptive Markov Gaussian Process (SAMGEP)

#### Description

Semi-supervised Adaptive Markov Gaussian Process (SAMGEP)

#### Usage

```
samgep(
 dat_train = NULL,
 dat_test = NULL,
 Cindices = NULL,
 w = NULL,
 w0 = NULL,
  V = NULL,
 observed = NULL,
 nX = 10,
  covs = NULL,
  survival = FALSE,
 Estep = Estep_partial,
 Xtrain = NULL,
 Xtest = NULL,
  alpha = NULL,
  r = NULL,
  lambda = NULL,
  surrIndex = NULL,
  nCores = 1
)
```

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#### samgep

#### Arguments

dat_train	(optional if Xtrain is supplied) Raw training data set, including patient IDs (ID), healthcare utilization feature (H) and censoring time (C)
dat_test	(optional) Raw testing data set, including patient IDs (ID), a healthcare utilization feature (H) and censoring time (C)
Cindices	(optional if Xtrain is supplied) Column indices of EHR feature counts in dat_train/dat_test
W	(optional if Xtrain is supplied) Pre-optimized EHR feature weights
wØ	(optional if Xtrain is supplied) Initial (i.e. partially optimized) EHR feature weights
V	(optional if Xtrain is supplied) nFeatures x nEmbeddings embeddings matrix
observed	(optional if Xtrain is supplied) IDs of patients with observed outcome labels
nX	Number of embedding features (defaults to 10)
covs	(optional) Baseline covariates to include in model; not yet operational
survival	Binary indicator of whether target phenotype is of type survival (i.e. stays posi- tive after incident event) or relapsing-remitting (defaults to FALSE)
Estep	E-step function to use (Estep_partial or Estep_full; defaults to Estep_partial)
Xtrain	(optional) Embedded training data set, including patient IDs (ID), healthcare utilization feature (H) and censoring time (C)
Xtest	(optional) Embedded testing data set, including patient IDs (ID), healthcare uti- lization feature (H) and censoring time (C)
alpha	(optional) Relative weight of semi-supervised to supervised MGP predictors in SAMGEP ensemble
r	(optional) Scaling factor of inter-temporal correlation
lambda	(optional) L1 regularization hyperparameter for feature weight (w) optimization
surrIndex	(optional) Index (within Cindices) of primary surrogate index for outcome event
nCores	Number of cores to use for parallelization (defaults to 1)

#### Value

w\_opt Optimized feature weights (w)

r\_opt Optimized inter-temporal correlation scaling factor (r) alpha\_opt Optimized semi-supservised:supervised relative weight (alpha) lambda\_opt Optiized L1 regularization hyperparameter (lambda) margSup Posterior probability predictions of supervised model (MGP Supervised) margMix Posterior probability predictions of semi-supervised model (MGP Semi-supervised) margMix Posterior probability predictions of SAMGEP cumSup Cumulative probability predictions of semi-supervised model (MGP Supervised) cumSemisup Cumulative probability predictions of semi-supervised model (MGP Supervised) cumMix Cumulative probability predictions of SAMGEP

#### simdata

simdata

# Description

Click **HERE** to view details.

# Usage

simdata

### Format

An object of class list of length 3.

# Examples

str(simdata)

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