

Package ‘PFoptim’

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Type Package

Title Global Stochastic Optimization using a Particle Filter Algorithm

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Description This package implements the G-PFSO (Global Particle Filter Stochastic Optimization) algorithm of Gerber and Douc (2021) for finding the global minimizer of a function defined through an expectation. Informally speaking, G-PFSO can be seen as a particle and derivative-free version of stochastic gradient methods.

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PFoptim-package

*The 'PFoptim' package: summary information***Description**

The package provides an implementation of the G-PFSO (Global Particle Filter Stochastic Optimization) algorithm of Gerber and Douc (2021) for finding the global minimizer of a function defined through an expectation. In addition, a function for implementing the SSP resampling algorithm (Gerber et al. 2019) and a function for implementing the Stratified resampling algorithm are also provided.

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References

Gerber M, Chopin N, Whiteley N (2019). "Negative association, ordering and convergence of resampling methods." *The Annals of Statistics*, **47**(4), 2236–2260.

Gerber M, Douc R (2021). "A global stochastic optimization particle filter algorithm." *arXiv preprint arXiv:2007.04803*.

gpfs

*Global Particle filter Stochastic Optimization***Description**

This function implements the G-PFSO (Global Particle Filter Stochastic Optimization) algorithm of Gerber and Douc (2021) for minimizing either the function $\theta \mapsto E[\text{fn}(\theta, Y)]$ from i.i.d. realizations y_1, \dots, y_n of Y or the function $\theta \mapsto \sum_{i=1}^n \text{fn}(\theta, y_i)$, where θ is a vector of dimension d .

Usage

```
gpfs(obs, N, fn, init, numit, resampling=c("SSP", "STRAT", "MULTI"), ..., control= list())
```

Arguments

obs Either a vector of observations or a matrix of observations (the number of rows being the sample size).

N Number of particles. The parameter N must be greater or equal to 2.

fn	function for a single observation. If theta is an N by d matrix and y is a single observation (i.e. y is a scalar if obs is a vector and a vector if obs is a matrix) then fn(theta,y) must be a vector of length N. If some rows of theta are outside the search space then the corresponding entries of the vector fn(theta,y) must be equal to Inf.
init	Function used to sample the initial particles such that init(N) is an N by d matrix (or alternatively a vector of length N if d=1).
numit	Number of iterations of the algorithm. If numit is not specified then G-PFSO estimates the minimizer of the function $E[\text{fn}(\theta, Y)]$ (in which case the observations are processed sequentially and numit is equal to the sample size). If numit is specified then G-PFSO computes the minimizer of the function $\sum_{i=1}^n \text{fn}(\theta, y_i)$.
resampling	Resampling algorithm to be used. Resampling should be either "SSP" (SSP resampling), "STRAT" (stratified resampling) or "MULTI" (multinomial resampling).
...	Further arguments to be passed to fn.
control	A list of control parameters. See details.

Details

Note that arguments after ... must be matched exactly.

G-PFSO computes two estimators of the minimizer of the objective function, namely the estimators $\bar{\theta}_{\text{numit}}^N$ and $\tilde{\theta}_{\text{numit}}^N$. The former is defined by $\bar{\theta}_{\text{numit}}^N = \frac{1}{\text{numit}} \sum_{t=1}^{\text{numit}} \bar{\theta}_t^N$ and converges to a particular element of the search space at a faster rate than the latter, but the latter estimator can find more quickly a small neighborhood of the minimizer of the objective function.

By default the sequence $(t_p)_{p \geq 0}$ is taken as

$$t_p = t_{p-1} + \lceil \max(A t_{p-1}^\varrho \log(t_{p-1}), B) \rceil$$

with $A=B=1$, $\varrho = 0.1$ and $t_0 = 5$. The value of A, B, ϱ and t_0 can be changed using the control argument (see below).

The control argument is a list that can supply any of the following components:

alpha: Parameter α of the learning rate $t^{-\alpha}$, which must be a strictly positive real number. By default, alpha=0.5.

Sigma: Scale matrix used to sample the particles. Sigma must be either a d by d covariance matrix or a strictly positive real number. In this latter case the scale matrix used to sample the particles is $\text{diag}(\text{Sigma}, d)$. By default, Sigma=1.

trace: If trace=TRUE then the value of $\tilde{\theta}_t$ and of the effective sample size ESS_t for all $t = 1, \dots, \text{numit}$ are returned. By default, trace=FALSE.

indep: If indep=TRUE and Sigma is a diagonal matrix or a scalar then the Student's t-distributions have independent components. By default, indep=TRUE and if Sigma is a not a diagonal matrix this parameter is ignored.

A: Parameter A of the sequence $(t_p)_{p \geq 0}$ used by default (see above). This parameter must be strictly positive.

- B:** Parameter B of the sequence $(t_p)_{p \geq 0}$ used by default (see above). This parameter must non-negative.
- varrho:** Parameter varrho of the sequence $(t_p)_{p \geq 0}$ used by default (see above). This parameter must be in the interval (0,1).
- t0:** Parameter t_0 of the the sequence $(t_p)_{p \geq 0}$ used by default (see above). This parameter must be a non-negative integer.
- nu:** Number of degrees of freedom of the Student's t-distributions used at time $t \in (t_p)_{\geq 0}$ to generate the new particles. By default nu=10
- c_ess:** A resampling step is performed when $ESS_t \leq N_{c_{ess}}$. This parameter must be in the interval (0,1] and by default c_ess=0.7.

Value

A list with the following components:

B_par	Value of $\bar{\theta}_{\text{numit}}^N$
T_par	Value of $\hat{\theta}_{\text{numit}}^N$
T_hist	Value of $\hat{\theta}_t^N$ for $t = 1, \dots, \text{numit}$ (only if trace=TRUE)
ESS	Value of the effective sample for $t = 1, \dots, \text{numit}$ (only if trace=TRUE)

References

Gerber M, Douc R (2021). “A global stochastic optimization particle filter algorithm.” *arXiv preprint arXiv:2007.04803*.

Examples

```
#Definition of fn
fn_toy<-function(theta, obs){
  test<-rep(0,nrow(theta))
  test[theta[,2]>0]<-1
  ll<-rep(-Inf,nrow(theta))
  ll[test==1]<-dnorm(obs,mean=theta[test==1,1], sd=theta[test==1,2],log=TRUE)
  return(-ll)
}
#Generate data y_1,...,y_n
n<-10000 #sample size
theta_star<-c(0,1) #true parameter value
y<-rnorm(n,mean=theta_star[1], sd=theta_star[2])
d<-length(theta_star)
#Define init function to be used
pi0<-function(N){
  return(cbind(rnorm(N,0,5), rexp(N)))
}
##Example 1: Maximum likelihood estimation in the Gaussian model
##true value of the MLE
mle<-c(mean(y),sd(y))
## use gpfso to compute the MLE
Est<-gpfso(y, N=100, fn=fn_toy, init=pi0, numit=20000, control=list(trace=TRUE))
```

```

## print  $\bar{\theta}^{N_{\text{numit}}}$  and  $\tilde{\theta}^{N_{\text{numit}}}$ 
print(Est$B_par)
print(Est$T_par)
##assess convergence
par(mfrow=c(1,2))
for(k in 1:2){
  plot(Est$T_hist[,k],type='l', xlab="iteration", ylab="estimated value")
  lines(cumsum(Est$T_hist[,k])/1:length(Est$T_hist[,k]),type='l', col='red')
  abline(h=mle[k])
}
##Example 2: Expected log-likelihood estimation in the Gaussian model
## Estimation of theta_star using gpfso
Est<-gpfso(y, N=100, fn=fn_toy, init=pi0, control=list(trace=TRUE))
## print  $\bar{\theta}^{N_{\text{numit}}}$  and  $\tilde{\theta}^{N_{\text{numit}}}$ 
print(Est$B_par)
print(Est$T_par)
##assess convergence
par(mfrow=c(1,2))
for(k in 1:2){
  plot(Est$T_hist[,k],type='l', xlab="iteration", ylab="estimated value")
  lines(cumsum(Est$T_hist[,k])/1:length(Est$T_hist[,k]),type='l', col='red')
  abline(h=theta_star[k])
}

```

SSP_Resampler

*SSP resampling***Description**

This function implements the SSP resampling algorithm (Gerber et al. 2019).

Usage

```
SSP_Resampler(U,W)
```

Arguments

W A vector of normalized weights.

U A vector of points in (0,1) such that $\text{length}(U)=\text{length}(W)$.

Details

For efficiency reasons, SSP_Resampler does not perform checks on the supplied arguments.

Value

A vector of length N with elements in the set $\{1, \dots, N\}$, with $N=\text{length}(U)=\text{length}(W)$.

References

Gerber M, Chopin N, Whiteley N (2019). “Negative association, ordering and convergence of resampling methods.” *The Annals of Statistics*, **47**(4), 2236–2260.

Examples

```
N<-100
W<-rbeta(N,0.5,2)
W<-W/sum(W)
J<-SSP_Resampler(runif(N),W)
```

Stratified_Resampler	<i>Stratified resampling</i>
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Description

This function implements the stratified resampling algorithm described see e.g. in Section 9.6 of Chopin and Papaspiliopoulos (2020)

Usage

```
Stratified_Resampler(U,W)
```

Arguments

W	A vector of normalized weights.
U	A vector of points in (0,1) such that $\text{length}(U)=\text{length}(W)$.

Details

For efficiency reasons, Stratified_Resampler does not perform checks on the supplied arguments.

Value

A vector of length N with elements in the set $\{1, \dots, N\}$, with $N=\text{length}(U)=\text{length}(W)$.

References

Chopin N, Papaspiliopoulos O (2020). *An introduction to sequential Monte Carlo*, volume 4. Springer.

Examples

```
N<-100
W<-rbeta(N,0.5,2)
W<-W/sum(W)
J<-Stratified_Resampler(runif(N),W)
```

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