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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LinkingTo Rcpp

RdMacros Rdpack

RoxygenNote 7.1.1.9001

R topics documented:

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 implementating 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*.

gpfso *Global Particle filter Stochastic Optimization*

Description

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

Usage

```
gpfso(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 $E[\mathrm{fn}(\theta, Y)]$ (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 $\sum_{i=1}^{n} \mathrm{fn}(\theta, y_i)$.
`resampling`	Resampling algorithm to be used. Resamping 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 $\bar{\theta}^N_{\mathrm{numit}}$ and $\tilde{\theta}^N_{\mathrm{numit}}$. The former is defined by $\bar{\theta}^N_{\mathrm{numit}} = \frac{1}{\mathrm{numit}} \sum_{t=1}^{\mathrm{numit}} \tilde{\theta}^N_t$ 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

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

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 `diag(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 $t = 1, \ldots, \mathrm{numit}$ 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=FALSE 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 resamling step is performed when $ESS_t <= Nc_{\text{ess}}$. 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 $\bar{\theta}^N_{\text{numit}}$
`T_par`	Value of $\tilde{\theta}^N_{\text{numit}}$
`T_hist`	Value of $\tilde{\theta}^N_t$ for $t = 1, ..., \text{numit}$ (only if trace=TRUE)
`ESS`	Value of the effective sample for $t = 1, ..., \text{numit}$ (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 funciton 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_{numit} and \tilde{\theta}^N_{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$T_hist[,k]),type='l', col='red')
  abline(h=mle[k])
}
##Example 2: Expected log-likelihood estimation in the Gaussian model
## Estimation of theta_star using gpfso
Est<-gpfso(y, N=100, fn=fn_toy, init=pi0, control=list(trace=TRUE))
## print \bar{\theta}^N_{numit} and \tilde{\theta}^N_{numit}
print(Est$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 `length(U)=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 $\{1, ..., N\}$, with `N=length(U)=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` *Stratified resampling*

Description

This function implements the stratified resampling algorithm descibed 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 `length(U)=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 $\{1, ..., N\}$, with `N=length(U)=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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