

Package ‘microbats’

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Type Package

Title An Implementation of Bat Algorithm in R

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Description A nature-inspired metaheuristic algorithm based on the echolocation behavior of microbats that uses frequency tuning to optimize problems in both continuous and discrete dimensions. This R package makes it easy to implement the standard bat algorithm on any user-supplied function. The algorithm was first developed by Xin-She Yang in 2010 (<DOI:10.1007/978-3-642-12538-6_6>, <DOI:10.1109/CINTI.2014.7028669>).

Depends R (>= 3.2.1)

License GPL (>= 2)

URL <https://github.com/stathwang/microbats>

LazyData TRUE

RoxygenNote 5.0.1

NeedsCompilation no

Repository CRAN

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bat_optim

*Bat Algorithm***Description**

The function `bat_optim` implements a nature-inspired metaheuristic algorithm that deals with both continuous and discrete optimization problems. The algorithm is based on the echolocation behavior of microbats and uses frequency tuning.

Usage

```
bat_optim(D, NP, N_Gen, A, r, Qmin, Qmax, Lower, Upper, FUN, ...)
```

Arguments

D	integer: the dimension of the search variables
NP	integer: the population size, typically between 10 and 40
N_Gen	integer: the number of generations, or iterations
A	numeric; loudness, between 0 and 1, can be constant or decreasing
r	numeric; pulse rate, must be positive, can be constant or decreasing
Qmin	minimum frequency
Qmax	maximum frequency
Lower	lower bound of the search variables
Upper	upper bound of the search variables
FUN	the objective function to optimize, must be supplied by a user
...	optional arguments to FUN

Details

`bat_optim` implements the standard bat algorithm in three robust steps. The first step is to initialize the parameters of algorithm to generate and evaluate the initial population from which to determine the best solution.

Secondly, a population of virtual microbats are moved in a d -dimensional search or solution space according to the updating rules of the algorithm: each bat is encoded with a velocity and a location at each iteration in the search space. The location is a solution vector, and the current best solution is achieved.

Then the current best solution is improved using random walks. The new solution is evaluated and updated. See *References* below for more details.

In essence, frequency tuning acts as mutation to vary the solutions locally; hence, increasing the range of frequencies leads to a global search. The mutation, compared with genetic algorithms, has no crossover but depends on loudness and pulse emission. So technically, varying loudness and pulse emission rates can also make the search intensive approaching the global optimality.

One of the advantages of the bat algorithm is that it can converge very quickly at the initial stage and can switch from exploration to exploitation when the optimality is approaching.

Value

Returns a list of four values: minimum fitness, population of solutions, fitness, best solution(s)

Author(s)

Seong Hyun Hwang, Rachel Myoung Moon

References

[1] Yang, X.-S. "A new metaheuristic bat-inspired algorithm." Nature inspired cooperative strategies for optimization (NICSO 2010). Springer Berlin Heidelberg, 2010. 65-74.

[2] Fister, I. Jr., Fister, I., Yang, X.-S., Fong, S., Zhuang, Y. "Bat algorithm: Recent advances." IEEE 15th International Symposium on Computational Intelligence and Informatics (CINTI), IEEE, 2014. 163-167.

Examples

```
# find the x-value that gives the minimum of the quadratic function y = x^2 - 3x
# x should then be 1.5
quad_func <- function(D, sol) {
  val = 0
  for (i in 1:D) {
    val <- val + sol[i] * sol[i] - sol[i] * 3
  }
  return(val)
}

# run a simulation using the standard bat algorithm
set.seed(5) # for reproducive results
fit <- bat_optim(D = 1, NP = 20, N_Gen = 100, A = 0.5, r = 0.5,
                Qmin = 0, Qmax = 2, Lower = -10, Upper = 10, FUN = quad_func)
x <- fit$best_solution
```

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