Package 'genetic.algo.optimizeR'

October 10, 2024

Title Genetic Algorithm Optimization
Version 0.3.2
Description Genetic algorithm are a class of optimization
 algorithms inspired by the process of natural selection and genetics.
 This package is for learning purposes and allows users to optimize
 various functions or parameters by mimicking biological evolution
 processes such as selection, crossover, and mutation. Ideal for tasks
 like machine learning parameter tuning, mathematical function
 optimization, and solving an optimization problem that involves finding
 the best solution in a discrete space.

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Type Package

URL https://danymukesha.github.io/genetic.algo.optimizeR/,
 https://github.com/danymukesha/genetic.algo.optimizeR

BugReports https://github.com/danymukesha/genetic.algo.optimizeR/issues

Imports dplyr, ggplot2, magrittr, rsconnect, stats, stringr, tinytex, biocViews, DiagrammeR

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crossover

Crossover of selected parents from the fitting population

Description

This function performs crossover between the selected individuals that fit the best based on the predefined condition(aim/objective).

e.g.: To optimize the function $f(x) = x^2 - 4x + 4$ to find the value of x that minimizes the function. x: represents a possible value the an individual from the population can have.

Usage

```
crossover(selected_parents, offspring_size)
```

Arguments

```
selected_parents
```

The list of selected individuals from the population.

offspring_size The number of offspring that the selected population should have.

Value

The output expected should be a list of offspring for the next generation.

Author(s)

Dany Mukesha

```
population <- c(1, 3, 0)

# Evaluate fitness
fitness <- evaluate_fitness(population)
print("Evaluation:")
print(fitness)

# Selection</pre>
```

evaluate_fitness 3

```
selected_parents <- selection(population, fitness, num_parents = 2)
print("Selection:")
print(selected_parents)

# Crossover
offspring <- crossover(selected_parents, offspring_size = 2)
print("Crossover:")
print(offspring)</pre>
```

evaluate_fitness

Evaluating the fitness of the population

Description

The function described below applies this equation:

$$f(x) = x^2 - 4x + 4$$

It assesses every individual x of a given population, to then provide an overview of how that population fits the equation.

Usage

```
evaluate_fitness(population)
```

Arguments

population

The list of individuals of the population.

Value

The output expected should be a list of f(x) values calculated from individuals in the population.

Author(s)

Dany Mukesha

```
# example of usage
population <- c(1, 3, 0)
# Evaluate fitness
genetic.algo.optimizeR::evaluate_fitness(population)</pre>
```

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Description

The function described below creates a population of individuals(values).

Usage

```
initialize_population(population_size, min = -100, max = 100)
```

Arguments

population_size

The number of individuals of the population.

min The minimum number an individual could have.

max The maximum number an individual could have.

Details

(Note: the values are random and the population should be highly diversified)

The space of x value is kept integer type and on range from 0 to 3, for simplification.

Value

The output expected should be a list of individuals of the population with the size indicated in the input.

Author(s)

Dany Mukesha

```
# example of usage
genetic.algo.optimizeR::initialize_population(population_size = 3, min = 0, max = 3)
```

mutation 5

mutation

Mutating the offspring

Description

The function described below mutates offspring from the selected individuals that fit the best based on the predefined condition(aim/objective).

e.g.: To optimize the function $f(x) = x^2 - 4x + 4$ to find the value of x that minimizes the function. x: represents a possible value the an individual from the population can have.

Usage

```
mutation(offspring, mutation_rate)
```

Arguments

```
offspring The list of offspring.

mutation_rate The probably of a single offspring to be modified/mutated.
```

Details

The mutation is needed to increase the diversity in the population and help the next generation close to the fitness.

Value

The output expected should be a list of mutated offspring.

Author(s)

Dany Mukesha

```
# example of usage
population <- c(1, 3, 0)

# Evaluate fitness.
fitness <- genetic.algo.optimizeR::evaluate_fitness(population)
print("Evaluation:")
print(fitness)

# Selection
selected_parents <- genetic.algo.optimizeR::selection(population, fitness, num_parents = 2)
print("Selection:")
print(selected_parents)

# Crossover
offspring <- genetic.algo.optimizeR::crossover(selected_parents, offspring_size = 2)</pre>
```

6 replacement

```
print("Crossover:")
print(offspring)

# Mutation
mutated_offspring <- genetic.algo.optimizeR::mutation(offspring, mutation_rate = 0)
# (no mutation in this example)
print(mutated_offspring)</pre>
```

replacement

Replacing non-selected individual(s)

Description

This function replace the individual(s) that was/were not selected (i.e. not the best fit) based on the predefined condition(aim/objective).

e.g.: To optimize the function $f(x) = x^2 - 4x + 4$ to find the value of x that minimizes the function. x: represents a possible value the an individual from the population can have.

Usage

```
replacement(population, offspring, num_to_replace)
```

Arguments

population The list of individuals of the population

offspring The list of offspring.

num_to_replace The number of selected individuals that should be replaced in the population.

Value

The output expected should be a list of selected individuals that fit the best with the predefined aim.

Author(s)

Dany Mukesha

```
# example of usage
population <- c(1, 3, 0)

# Evaluate fitness
fitness <- genetic.algo.optimizeR::evaluate_fitness(population)
print("Evaluation:")
print(fitness)

# Selection
selected_parents <- genetic.algo.optimizeR::selection(population, fitness, num_parents = 2)</pre>
```

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```
print("Selection:")
print(selected_parents)

# Crossover and mutation
offspring <- genetic.algo.optimizeR::crossover(selected_parents, offspring_size = 2)
mutated_offspring <- mutation(offspring, mutation_rate = 0) # (no mutation in this example)
print(mutated_offspring)

# Replacement
population <- genetic.algo.optimizeR::replacement(population, mutated_offspring, num_to_replace = 1)
print("Replacement:")
print(population)</pre>
```

selection

Selecting the fitting the population

Description

The function described below selects the individuals that fit the best based on the predefined condition(aim/objective).

e.g.: To optimize the function $f(x) = x^2 - 4x + 4$ to find the value of x that minimizes the function. x: represents a possible value the an individual from the population can have.

Usage

```
selection(population, fitness, num_parents)
```

Arguments

population The list of individuals of the population

fitness The list of individuals(value) obtained from the function of genetic.algo.optimizeR

namely 'evaluate_fitness'.

num_parents The number of selected individuals that fit the best with the predefined aim.

Value

The output expected should be a list of selected individuals that fit the best with the predefined condition(aim/objective).

Author(s)

Dany Mukesha

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```
# example of usage
library(genetic.algo.optimizeR)

population <- c(1, 3, 0)

# Evaluate fitness
fitness <- genetic.algo.optimizeR::evaluate_fitness(population)
print("Evaluation:")
print(fitness)

# Selection
selected_parents <- genetic.algo.optimizeR::selection(population, fitness, num_parents = 2)
print("Selection:")
print(selected_parents)</pre>
```

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