

Lab 8

Optimization via Gene Expression

Function(x):

return $\sum (x^{**2})$

Initialize - Population (pop^{size}, num-genes, lower, upper)

pop \leftarrow random (lower, upper, (pop^{size}, num-genes))

return pop

Evaluate-fitness (pop, function):

fitness \leftarrow zeros (pop.shape[0])
 for i in range pop.shape[0]:
 fitness[i] \leftarrow function (pop[i])
 return fitness

selection (pop, fitness, num-selected):

prob \leftarrow fitness / fitness.sum()
 indices \leftarrow random.choice (range(len(pop)),
 size = num-selected, p = prob)

selected-pop \leftarrow pop[indices]

return selected-pop

Crossover (selected-pop, crossover):

new-pop \leftarrow []
 num-individuals \leftarrow len(selected-pop)

for i in range 0 to num-individuals-1, step 2:

$p1, p2 \leftarrow \text{selected-pop}[i], \text{selected-pop}[i+1]$
 if $\text{len}(p1) > 1$ and $\text{random} < \text{crossover}$:

$\text{crossover-point} \leftarrow \text{random}(1, \text{len}(p1) - 1)$
 $\text{child1} \leftarrow \text{concatenate}(p1[:\text{crossover-point}], p2[\text{crossover-point}:])$

$\text{child2} \leftarrow \text{concatenate}(p2[:\text{crossover-point}], p1[\text{crossover-point}:])$

$\text{new-pop.extend}([p1, p2])$

if $\text{num-individuals} \% 2 \neq 1$:

$\text{new-pop.append}(\text{selected-pop}[-1])$

return new-pop

Mutation(pop, rate, lower, upper):

for i in range(pop.shape[0]):

if $\text{random} < \text{rate}$:

$\text{gene} \leftarrow \text{random}(0, \text{pop.shape}[1] - 1)$
 $\text{pop}[i, \text{gene}] \leftarrow \text{random-uniform}(\text{lower}, \text{upper})$

return pop

Gene-expression(individual, function):

return function(individual)

Gene-Expression-Algorithm (pop-size, num-genes, lower, upper, max-gen, mutation-rate, crossover-rate, function):

pop \leftarrow initialize-population (pop-size, num-genes, lower, upper)

best-sol \leftarrow NIL

best-fit $\leftarrow \infty$

for gen in 0 to max-gen:

fitness \leftarrow evaluate-fitness (pop, function)
min-fitness \leftarrow fitness.min()

if min-fitness < best-fit:

best-fit \leftarrow min-fitness

best-sol \leftarrow pop [min(fitness)]

selected-pop \leftarrow selection (pop, fitness, pop-size // 2)

offspring-pop \leftarrow crossover (selected-pop, crossover-rate)

pop \leftarrow mutation (offspring-pop, mutation-rate, lower, upper)

print (Generation, fitness)

return best-sol, best-fit

Inputs :

pop-size = 50

num-genes = 1

lower = -5

upper = 5

max-gen = 100

mutation-rate = 0.1

crossover-rate = 0.7