

Project 1 - Genetic Algorithm

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Abstract

My Genetic Algorithm works well for most functions, except Rastrigin and Schwefel function. I tested all the crossover methods, except "Permutation". Among the crossover methods, I find that 1 point crossover works the best for most of the functions.

1 Description for Genetic Algorithm

Generally, my Genetic Algorithm used Steady-State type that:

- generate a population
- have enough loops
- for each loop
 1. tournament select two parents
 2. crossover (mainly by 1 point crossover)
 3. mutate 2 children
 4. replace two loser (tournament selection) by two children
- finally check the fitness of each individual and find the best

1.1 Fitness function: just math, no need to explain

- Sphere:

```
double evaluate(double x[X_I])
{
    double z = 0;
    double tmp;
    int i;
    for(i = 0 ; i < X_I; i++ )
    {
        tmp = x[i];
        z += tmp*tmp;
    }

    return z;
}
```

- Rosenbrock

```
double evaluate(double x[X_I])
{
    double z = 0;
    int i;
    for(i = 0; i < X_I - 1; i++) {
```

```

        z += 100*pow(x[i+1]-pow(x[i],2) , 2) + pow(x[i] - 1, 2);
    }

    return z;
}

```

- Rastrigin

```

double evaluate(double x[X_I])
{
    double z = 10*X_I;
    int i;
    for(i = 0; i < X_I; i++){
        z += pow(x[i] , 2) - 10*cos(2*M_PI*x[i]);
    }
    return z;
}

```

- Schwefel

```

double evaluate(double x[X_I])
{
    double z = 418.9829*X_I;
    int i;
    for(i = 0; i < X_I; i++){
        z += x[i]*sin(sqrt(abs(x[i])));
    }
    return z;
}

```

- Ackley

```

double evaluate(double x[X_I])
{
    double z = 20+exp(1);
    double a = 0;
    double b = 0;
    int i;
    for(i = 0; i<X_I;i++){
        a += pow(x[i] , 2);
        b += cos(2*M_PI*x[i]);
    }
    z = z - 20*exp(-0.2*sqrt(a/X_I)) - exp(b/X_I);

    return z;
}

```

- Griewangk

```

double evaluate(double x[X_I])
{
    double z = 1;
    double a = 0;
    double b = 1;

```

```

    int i;
    for(i = 0; i < X_I; i++) {
        a += pow(x[i],2) / 4000.00;
        b *= cos(x[i]/(i+1));
    }
    z = z + a - b;

    return z;
}

```

1.2 Generate initial random solutions

- ```

// Generate a population
for 0 to Population step 1
 // Generate each individual
 for 0 to 30 step 1
 // Take random values in range for each vector
 // Calculate each individuals fitness, put into an array

```

## 1.3 Crossover

Once we get the parents, we will generate two children by crossover. I did 4 crossover methods, listed below:

- one point crossover

```

void crossover(double father[X_I],
 double mother[X_I], double children[2][X_I])
{
 int cut = rand()%X_I; // get a random cut/point
 int i;
 for(i = 0; i < cut; i++){ // keep the front half
 children[0][i] = father[i];
 children[1][i] = mother[i];
 }
 for(; i < X_I; i++){ // swap/cross the back half
 children[1][i] = father[i];
 children[0][i] = mother[i];
 }
}

```

- two points crossover

```

void crossover(double father[X_I],
 double mother[X_I], double children[2][X_I])
{
 // get two random cuts/points
 int cut1 = rand()%X_I;
 int cut2 = rand()%(X_I - cut1);
 int i;

```

```

 for(i = 0; i < cut1; i++){
 children[0][i] = father[i];
 children[1][i] = mother[i];
 }
 for(; i < cut1+cut2; i++){ // only swap the middle
 children[1][i] = father[i];
 children[0][i] = mother[i];
 }
 for(; i < X_I; i++){
 children[0][i] = father[i];
 children[1][i] = mother[i];
 }
}

```

- uniform crossover

```

void crossover(double father[X_I],
 double mother[X_I], double children[2][X_I])
{
 double tmp= 0;
 int P = 0;
 int i;
 for(i = 0; i < X_I; i++) {
 // for each value, we have 30% chance to swap that "gene"
 if((P = rand()%100) < 30) {
 tmp = children[0][i];
 children[0][i] = children[1][i];
 children[1][i] = tmp;
 }
 }
}

```

- arithmetic

```

void crossover(double father[X_I],
 double mother[X_I], double children[2][X_I])
{
 double tmp = 0;
 int i;
 for(i = 0; i < X_I; i++) {
 // get the avg of two values and get two same children
 tmp = (children[0][i] + children[1][i]) / 2;
 children[0][i] = tmp;
 children[1][i] = tmp;
 }
}

```

## 1.4 Mutation

Mutate the children by “creep”. For each vector in child, we have 50percent chance to mutate the value by +/- a small value (0.001 for Rosenbrock; 0.01 for Sphere, Rastrigin; 0.1 for Ackley; 1 for Schwefel, Griewangk).

- ```
void mutate(double child[X_I])
{
    for 0 to 30 step by 1
        if rand()%10 < 5 // 50% chance
            // 50% + and 50% -
            child[i] += (rand()%10 < 5 ? -1 : 1)*INTERVAL;
}
```

1.5 GeneticAlgorithm

The major algorithm part. It will finally return the best fitness, also pass the reference of the best individual (solution) so far.

- ```
/* Steady-State */
double GeneticAlgorithm(double resultA[X_I])
{
 // Generate a population
 // Calculate each individuals fitness, put into an array

 while(count < population*100) // as Dr. Soul suggested
 {
 // select father by Tournament
 // select mother by Tournament (diff from father)

 // crossover father and mother to get two children

 // Mutate both children

 // select two losers by Tournament
 // Replace two losers by two children
 }

 /* Finally, check the fitness array */
 bestEval <- fitness[0](inital);
 for 0 to population step 1
 if fitness[i] < bestEval
 bestEval <- fitness[i];
 best = i // record the position;

 // copy best solution to resultA

 return bestEval;
}
```

## 1.6 Selection

This is a tournament selection function which select a winner/loser (depend on the index) from a sample of N random individuals (N is also a random number). Finally the function returns a int which is the index of that individual among population array.

```

 /* Tournament selection */
int selection(int good_poor, double fitness[POP])
/* good_poor is the index to determe we are select a winner or loser*/
{
 // get a random winner and calc his fitness
 winner <- rand()%POP;
 winner_fitness <- fitness[winner];

 // generate a random N
 N <- rand()%POP ;
 // loop to get the best
 for 0 to N step 1
 temp <- rand%POP
 if(good_poor == GOOD && fitness[temp] < winner_fitness){
 // this is for winner
 winner_fitness <- fitness[temp];
 winner <- temp;
 } else if(good_poor == POOR && fitness[temp] > winner_fitness){
 // this is for loser
 winner_fitness <- fitness[temp];
 winner <- temp;
 }

 return winner; // the index among population
}

```

## 2 Results

### 2.1 Sphere

| Crossover           | 1point     | 2points    | uniform    | arithmetic |
|---------------------|------------|------------|------------|------------|
| Population          | 50         | 50         | 50         | 50         |
| Mutation Interval   | 0.01       | 0.01       | 0.01       | 0.01       |
| Running Times(*POP) | 100        | 100        | 100        | 100        |
| 1st Best Fitness    | 187.526800 | 148.298600 | 142.681100 | 148.242200 |
| 1st Avg Fitness     | 258.182810 | 259.975670 | 258.766226 | 265.070316 |
| Final Best Fitness  | 0.002500   | 0.001200   | 2.557200   | 1.217325   |
| Final Avg Fitness   | 0.003112   | 0.001756   | 5.150842   | 2.173404   |

## 2.2 Rosenbrock

| Crossover           | 1point       | 2points      | uniform      | arithmetic   |
|---------------------|--------------|--------------|--------------|--------------|
| Population          | 100          | 100          | 100          | 100          |
| Mutation Interval   | 0.001        | 0.001        | 0.001        | 0.001        |
| Running Times(*POP) | 2000         | 2000         | 2000         | 2000         |
| 1st Best Fitness    | 6633.415331  | 6537.239168  | 6105.194947  | 5716.116572  |
| 1st Avg Fitness     | 14865.619920 | 14462.804456 | 14392.547199 | 14755.587222 |
| Final Best Fitness  | 0.192033     | 4.180114     | 159.677838   | 141.739331   |
| Final Avg Fitness   | 0.196815     | 4.183143     | 209.094239   | 148.872279   |

## 2.3 Rastrigin

My genetic algorithm doesn't work for Rastrigin function. The best result I get will stay (little change) at some point, even I add more running times. But the fitness function I write should be correct. I have no idea about it.

## 2.4 Schwefel

For this function, I get the same trouble as Rastrigin function.

## 2.5 Ackley

| Crossover           | 1point    | 2points   | uniform   | arithmetic |
|---------------------|-----------|-----------|-----------|------------|
| Population          | 50        | 50        | 50        | 50         |
| Mutation Interval   | 1         | 1         | 1         | 1          |
| Running Times(*POP) | 1000      | 1000      | 1000      | 1000       |
| 1st Best Fitness    | 18.924080 | 18.885888 | 18.775362 | 18.762705  |
| 1st Avg Fitness     | 19.334254 | 19.379073 | 19.375065 | 19.308400  |
| Final Best Fitness  | 1.711187  | 1.841785  | 2.637531  | 2.467152   |
| Final Avg Fitness   | 2.319135  | 2.409010  | 14.853659 | 15.595730  |

## 2.6 Griewangk

| Crossover           | 1point     | 2points    | uniform    | arithmetic |
|---------------------|------------|------------|------------|------------|
| Population          | 50         | 50         | 50         | 50         |
| Mutation Interval   | 1          | 1          | 1          | 1          |
| Running Times(*POP) | 100        | 100        | 100        | 100        |
| 1st Best Fitness    | 533.451750 | 612.249000 | 594.165750 | 518.436000 |
| 1st Avg Fitness     | 929.957840 | 915.618320 | 906.391795 | 900.314490 |
| Final Best Fitness  | 0.277369   | 2.207000   | 6.057750   | 4.453250   |
| Final Avg Fitness   | 0.391046   | 2.354080   | 11.263200  | 10.335357  |



### **3 Conclusion**

My Genetic Algorithm should be successful, but still need more researches to figure out the trouble I get on Rastrigin and Schwefel functions.