



## ASSIGNMENT 3

Subject : Linked-list Implementation

Advisor : : Res. Assist. Selim YILMAZ

Programming Language: C

Due Date: 02.01.2020 — 18:59:59

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# INTRODUCTION / AIM:

In this assignment, we are expected to gain knowledge on C language including linked-list implementation and dynamic memory allocation. Prior knowledge on basic C syntax is assumed.

## PROBLEM DEFINITION:

Our task is, storing selection, mutate, population, xover files on program. Program arguments are problem size, population size, generation size.

The detailed version of your GA implementation is given as follows:

Read command line arguments;

Load all the parameters including selection, mutation, crossover;

Initialize population w.r.t. population file;

Evaluate the fitness and ranking of each chromosome;

Sort the population in ascending order;

Store the best chromosome;

Print the population with the best chromosome found so far;

repeat

    Apply selection;

    Apply crossover;

    Apply mutation;

    Evaluate the fitness and ranking of each chromosome;

    Sort the population in ascending order;

    Update the best chromosome if found better;

    Print the population with the best chromosome found so far;

until MAX GEN is reached;

return best chromosome;

## STRUCTS THAT I USED:

```
struct dna {  
    int dna;  
    struct dna *next;  
};
```

dna struct stores:

- dna(gene)

-link of next dna

```

struct gen {
    int fitness;
    struct dna *dna_data;
    struct gen *next;
};

```

gen struct stores:

- fitness
- link of next dna
- link of next gen
- #its chrosome's struct

## FUNCTIONS THAT I USED:

**void readFiles(struct gen \*\*head, int dna\_size, int pop\_size, char bestChromosome[])**

Function read 4 file which are population, selection, xover, mutate. Function call addGen() then for each line in selection file it call evolution(),doMutate(),calculateFitness() functions.

**void addGen(struct gen \*\*head, char genList[], int n)**

Function creates new nodes of gen then call addDnaToGen() function for each dna.

**void addDnaToGen(struct gen \*head, int dna)**

For each dna function creates new node of dna.

**void sort(struct gen \*\*head)**

Function sort the Dna's for their fitness.

**void calculateFitness(struct gen \*head,int n, int generation, char bestChromosome[])**

Function calculate the fitness of dna's.

**void doMutate(struct gen \*head, int mutateIndex)**

Function mutates the gen.

**void doXover(struct gen \*\*head, int genIndex1, int genIndex2, int dnaIndex1, int dnaIndex2)**

Function cross the gens of given dna's.

**void evolution(struct gen \*head, char selectionLine[], char xoverLine[], int pop\_size)**

Function calls format the selection and xoverline then call doXover()

**void Display(struct gen \*head, int generation, char bestChromosome[])**

It's print function

**void freeList(struct gen\* head)**

It's free the structs

**int main(int argc,char\* argv[])**

Main gets arguments, then call readFiles()

# MY ALGORITHM

My program starts with readFiles() function. "readFiles()" read the files line by line. First while population file==NULL, readFiles() call addGen. And Then AddGen() calls addDnaToGen(). Then readFiles() call evolution(), doMutate(), calculateFitness() functions respectively. evolution() calls doXover(). calculateFitness() call sort() then Display(). So I crate my first generation with my files first line. When readFiles() get other line, this steps repeated by program.