

FUNCTION MAXIMA EVALUATION USING GENETIC ALGORITHM

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import java.io.*;
import java.lang.Math;
import java.util.*;
import javax.script.*;

class GA4
{
    static BufferedReader br=new BufferedReader(new InputStreamReader(System.in));
    static ScriptEngine engine = (new ScriptEngineManager()).getEngineByName("JavaScript");
    static int nvar=0,len=0,maxvar=10;
    static double x[][]=new double[maxvar][5]; //FIELDS: en preci lwrLim uprLim length
    static double n[];
    public static void main(String args[]) throws Exception
    {
        System.out.print("\nFunction\t: ");
        String fneq=br.readLine();
        //EXAMPLE : fneq=21.5+x1*Math.sin(4*Math.PI*x1)+x2*Math.sin(20*Math.PI*x2);
        int i,j,gen;
        for(i=0;i<10;i++)
            x[i][0]=0;
        final int len=fnvr(fneq);
        System.out.print("Population Size\t: ");
        final int pop_size=Integer.parseInt(br.readLine());
        System.out.print("Crossover Prob.\t: ");
        final double pc=Double.parseDouble(br.readLine());
        System.out.print("Mutation Prob.\t: ");
        final double pm=Double.parseDouble(br.readLine());
        System.out.print("Generations\t: ");
        final int gtot=Integer.parseInt(br.readLine());

        String chr[]=new String [pop_size],tmp;
        //initialize 1st generation
        for(i=0;i<pop_size;i++)
        {
            tmp="";
            for(j=0;j<len;j++)
            {
                if(Math.random()>0.5) tmp+="1";
                else tmp+="0";
            }
            chr[i]=tmp;
        }
        System.out.println("\nSTART OF GENETIC ITERATION");
        int eop,pos,c1=0,c2=0;
        double r,ns,best[]=new double [nvar+2];
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double pqr[]=new double [pop_size];
n=new double[nvar];
String chrt[]=new String [pop_size]; //temp Str arr : next gen chromosomes
//main looping for each generation
for(gen=0;gen<gtot;gen++)
{
    System.out.println("Generation : "+(gen+1));
    for(i=0;i<pop_size;i++)
    {
        pos=0;
        for(j=0;j<nvar;j++)
        {
            n[j]=B2D(chr[i].substring(pos,pos+(int)x[j][4]),j);
            pos+=(int)x[j][4];
        }
        if(i==0)
        {
            pqr[i]=evalt(fneq);
            if(pqr[i]>best[0])
            {
                best[0]=pqr[i];
                best[1]=gen+1;
                for(j=0;j<nvar;j++)
                    best[j+2]=n[j];
            }
        }
        else
        {
            pqr[i]=pqr[i-1]+evalt(fneq);
            if((pqr[i]-pqr[i-1])>best[0])
            {
                best[0]=pqr[i]-pqr[i-1];
                best[1]=gen+1;
                for(j=0;j<nvar;j++)
                    best[j+2]=n[j];
            }
        }
    }
    System.out.println("Total Value = "+pqr[pop_size-1]);
    //select chromosomes for next generation
    for(i=0;i<pop_size;i++)
        pqr[i]/=pqr[pop_size-1]; //cumulative prob. Roulette Wheel
    for(i=0;i<pop_size;i++)
    {
        r=Math.random();
        for(j=0;j<pop_size;j++)
        {
            if(r<pqr[j])
            {
                chrt[i]=chr[j];
            }
        }
    }
}

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        break;
    }
}
//Crossover
ns=0;
for(i=0;i<pop_size;i++)
{
    if(Math.random()<pc)
    {
        pqr[i]=1;
        ns++;
    }
    else pqr[i]=0;
}
if(ns%2!=0) //odd no. of selected chromosome selected for pairing
    ns=Math.random()*(pop_size-ns);
for(i=0;i<pop_size;i++) //select extra chromosome for crossover pairing
{
    if(pqr[i]==0) ns--;
    if(ns==0)
    {
        pqr[i]=1;
        break;
    }
}
//perform Crossover
for(i=0;i<pop_size;i++)
{
    if(pqr[i]==0) continue;
    eop=0;
    if(eop==0)
    {
        c1=i;
        eop=1;
    }
    else if(eop==1)
    {
        c2=i;
        eop=2;
    }
    if(eop==2)
    {
        pos=(int)Math.random()*len;
        chrt[c1]=chrt[c1].substring(0,pos)+chrt[c2].substring(pos);
        chrt[c2]=chrt[c2].substring(0,pos)+chrt[c1].substring(pos);
        eop=0;
    }
}
}
//Mutation

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        for(i=0;i<pop_size*len;i++)
        {
            if(Math.random()<pm)
            {
                j=i/len;
                pos=i%len;
                if(chrt[j].charAt(pos)=='0')
                    chrt[j]=chrt[j].substring(0,pos)+"1"+chrt[j].substring(pos);
                else
                    chrt[j]=chrt[j].substring(0,pos)+"0"+chrt[j].substring(pos);
            }
        }
        for(i=0;i<pop_size;i++)
            chr[i]=chrt[i];
    }
    //Display Best Values
    System.out.println("\nBest Value : "+best[0]);
    System.out.println("Generation : "+best[1]);
    System.out.println("Values   :");
    for(i=2;i<nvar+2;i++)
        System.out.println("x" +(i-1) + " \t: " +best[i]);
}
//Method to evaluate string expression using JavaScript
public static double evalt(String fneq)throws Exception
{
    double fn=0;
    for(int i=0;i<nvar;i++)
    {
        engine.put(("x" +(i+1)),n[i]);
    }
    fn=Double.parseDouble(engine.eval(fneq)+"");
    return fn;
}
//Method to convert binary chromosome sequence to its numerical value
public static double B2D(String s, int vn)
{
    double n=0.0;
    for(int i=0;i<s.length();i++)
        if(s.charAt(i)=='1') n+=Math.pow(2,(s.length()-i-1));
    n=x[vn][2]+n*(x[vn][3]-x[vn][2])/(Math.pow(2,s.length()-1));
    return n;
}
//Method to evaluate expression variables and their parameters
public static int fnvr(String fneq)throws Exception
{
    int i,j,len=0;
    for(i=0;i<fneq.length();i++)
    {
        if(fneq.charAt(i)=='x')
        {

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int idx=Integer.parseInt(fneq.charAt(i+1)+"");
idx--;
if(x[idx][0]==0)
{
    System.out.println("Variable\t: "+(idx+1));
    System.out.print("Precision\t: ");
    x[idx][1]=Double.parseDouble(br.readLine());
    System.out.print("Lower Limit\t: ");
    x[idx][2]=Double.parseDouble(br.readLine());
    System.out.print("Upper Limit\t: ");
    x[idx][3]=Double.parseDouble(br.readLine());
    x[idx][0]=1;
    nvar++;
    for(j=1;;j++)
    {
        if(Math.pow(2,j)>(x[idx][1]*(x[idx][3]-x[idx][2])))
        {
            x[idx][4]=j;
            break;
        }
    }
}
}
}
for(i=0;i<nvar;i++)
    len+=x[i][4];
return (len);
}
}

```

TERMINAL WINDOW :

Function	: 21.5+x1*Math.sin(4*Math.PI*x1)+x2*Math.sin(20*Math.PI*x2)
Variable	: 1
Precision	: 10000
Lower Limit	: -3.0
Upper Limit	: 12.1
Variable	: 2
Precision	: 10000
Lower Limit	: 4.1
Upper Limit	: 5.8
Population Size	: 20
Crossover Prob.	: 0.25
Mutation Prob.	: 0.01
Generations	: 10000

START OF GENETIC ITERATION

Generation : 1

Total Value = 409.79377257070024

Generation : 2

Total Value = 445.9614442942778

Generation : 3

Total Value = 459.3753003342473

Generation : 4

Total Value = 448.226443550758

Generation : 5

Total Value = 473.81885531853123

Generation : 6

Total Value = 509.41613116246873

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Generation : 6914

Total Value = 598.3270407702545

Generation : 6915

Total Value = 598.6102671938445

Generation : 6916

Total Value = 579.8202426974135

Generation : 6917

Total Value = 603.6994231395291

Generation : 6918

Total Value = 601.9279309504764

Generation : 6919

Total Value = 606.0494566642751

Generation : 6920

Total Value = 545.9536540717735

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Generation : 9997

Total Value = 605.2871812340687

Generation : 9998

Total Value = 582.4194264851458

Generation : 9999

Total Value = 610.8088485016723

Generation : 10000

Total Value = 585.362350731463

Best Value : 38.84702806965913

Generation : 6917.0

Values :

x1 : 11.623918243096325

x2 : 5.724771874141667