

Knapsack Problem

10.27

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My parameters:

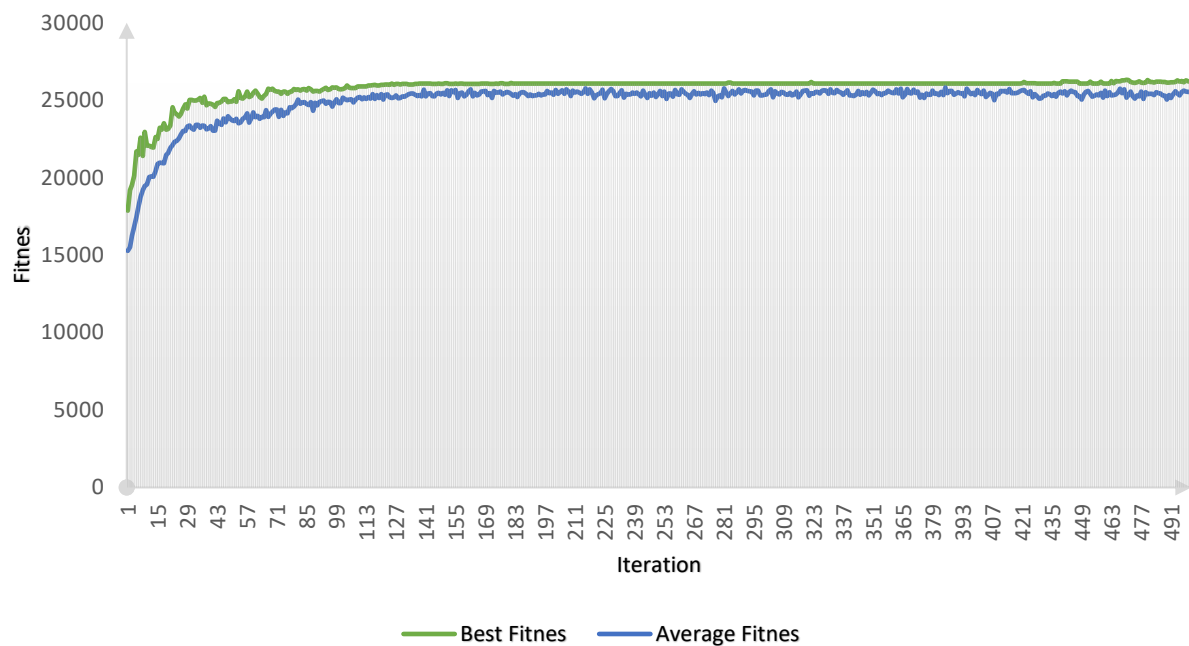
A (size of knapsack) = 10000

limit of n = 350

Random generated w_i and x_i :

Volume	1	2	3	3	9	6	10	5	7	2	6	1	1	6	4	1	1	9	6	6
Price	9	4	10	7	7	10	1	7	7	3	1	7	5	6	4	3	8	10	9	9

Graphic: iteration from fitness



Tables with 5 best chromosomes:

1 Iteration

Best Five chromosome

```
238 221 78 155 32 211 16 298 51 60 44 331 219 46 230 129 138 65 22 113
275 24 49 250 106 58 77 317 104 201 34 281 6 95 64 35 208 6 239 91
232 11 26 341 7 256 267 47 112 332 48 246 259 23 61 156 284 22 50 63
324 74 202 201 35 29 100 229 190 92 14 151 126 16 23 91 73 106 275 6
347 296 331 141 69 83 53 101 3 35 68 97 145 36 11 64 60 118 33 230
```

100 Iteration

Best Five chromosome

```
342 348 341 312 10 282 19 101 3 199 3 345 345 6 52 320 313 21 125 247
342 345 341 312 10 282 19 101 3 199 3 345 345 6 52 320 313 21 125 247
342 345 341 312 10 282 19 101 3 263 3 316 345 6 52 320 313 21 125 247
342 345 341 312 10 282 19 94 3 310 3 316 345 6 75 320 313 17 107 247
342 345 341 312 10 282 19 94 3 263 3 316 345 6 52 320 313 21 125 247
```

200 Iteration

Best Five chromosome

342	345	347	312	10	282	19	101	3	199	3	345	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	345	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	345	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	345	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	345	345	13	55	320	346	21	125	247

300 Iteration

Best Five chromosome

342	345	347	312	10	282	19	101	3	199	3	346	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	346	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	346	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	346	345	13	55	320	346	21	125	247
342	345	347	312	10	282	19	101	3	199	3	346	345	13	55	320	346	21	125	247

500 Iteration

Best Five chromosome

342	345	347	312	10	282	2	101	3	198	12	349	345	28	61	317	346	21	125	247
342	345	347	312	10	282	2	101	17	198	3	349	345	13	61	317	346	21	125	247
342	345	347	312	12	282	2	101	3	198	12	349	345	20	61	317	346	21	125	247
342	345	347	312	12	282	2	101	3	198	3	349	345	20	61	317	346	21	125	247
342	345	347	312	12	282	2	101	3	198	3	349	345	20	61	317	346	21	125	247

Source Code

```
package brestterespol;

import java.util.Random;

class GeneticAlgo {

    public enum SelectionType {
        TOURNEY, ROULETTE_WHEEL, TRUNCATING
    }

    public enum CrossingType {
        UNIFORM, TWO_POINT_RECOMBINATION, ELEMENTWISE_RECOMBINATION, ONE_ELEMENT_EXCHANGE
    }

    private SelectionType slctp; //Тип Селекции
    private CrossingType crstp; //Тип Скрещивания
    private static int genomLength = 20; //Длина хромосомы
    private static int generationCount = 10000000; //Кол-во поколений
    private static int individualCount = 40; //Кол-во Геномов (Индивидов, Особей) в поколении
    private static int limitVolumes = 10000; //ограничение по общему объему продуктов
    private static int limitPrice = 10; //ограничение по цене продукта
    private static int limitVolume = 10; //ограничение по объему, 1 продукта
    private int limitCount = 350; //лимит по количеству
    private int[][] population; //популяция. где каждый бит в особи = количество данного продукта
    int[] price;
    int[] volume;
    int[] sortFitnes;

    public GeneticAlgo(String s, String c) {
        slctp = SelectionType.valueOf(s);
        crstp = CrossingType.valueOf(c);
        population = new int[individualCount + 1][genomLength];
        volume = new int[genomLength];
    }
}
```

```

        price = new int[genomLength];
        sortFitnes = new int[individualCount];
    }

    public void run() {
        this.generateFirstGeneration();
        for (int i = 0; i < generationCount; i++) {
            this.selection();
        }
    }

    private void generateFirstGeneration() {
        Random rnd = new Random();
        for (int j = 0; j < genomLength; j++) {
            volume[j] = Math.abs(rnd.nextInt()) % limitVolume + 1;
            price[j] = Math.abs(rnd.nextInt()) % limitPrice + 1;
        }
        for (int i = 0; i < individualCount; i++) {
            do {
                for (int j = 0; j < genomLength; j++) {
                    population[i][j] = Math.abs(rnd.nextInt()) % limitCount;
                }
            } while (!getLimit(i));
        }
    } //генерация первого поколения

    private void selection() {
        int[][] genomListOffsprings = new int[individualCount][genomLength];
        Random rndd = new Random();
        switch (this.slctp) {
            case TRUNCATING: {
                this.sort();
                this.print();
                for (int i = 0; i < GeneticAlgo.individualCount; i++) {
                    do {
                        genomListOffsprings[i] = this.crossing((Math.abs(rndd.nextInt())) %
individualCount / 2, (Math.abs(rndd.nextInt())) % individualCount / 2);
                        population[individualCount] = genomListOffsprings[i];
                    } while (!getLimit(individualCount));
                }
                System.arraycopy(genomListOffsprings, 0, population, 0,
GeneticAlgo.individualCount);
                break;
            }
            default:
                break;
        }
    } //Процедура селекции

    private void sort() {
        for (int i = 0; i < GeneticAlgo.individualCount; i++) {
            sortFitnes[i] = fitnes(i);
        }
        for (int i = 0; i < GeneticAlgo.individualCount; i++) {
            int[] genom;
            for (int j = i; j < GeneticAlgo.individualCount; j++) {
                if (sortFitnes[i] < sortFitnes[j]) {
                    int cur = sortFitnes[i];
                    sortFitnes[i] = sortFitnes[j];
                    sortFitnes[j] = cur;
                    genom = population[i];
                    population[i] = population[j];
                    population[j] = genom;
                }
            }
        }
    }
}

```

```

}

private int fitnes(int nomber) {
    int ftns = 0;
    for (int j = 0; j < genomLength; j++) {
        ftns += price[j] * population[nomber][j];
    }
    return (getLimit(nomber)) ? ftns : 0;
} //Фитнес функция

private void print() {
    System.out.print(sortFitnes[0]);
    System.out.print(":");
    System.out.print(avgfitnes());
    System.out.println();
}

private int[] crossing(int a, int b) {
    int[] vec = new int[genomLength];
    Random rand = new Random();
    switch (crstp) {
        case UNIFORM: {
            for (int i = 0; i < genomLength; i++) {
                boolean n = rand.nextBoolean();
                vec[i] = (n) ? population[b][i] : population[a][i];
            }
            break;
        }
        default:
            break;
    }
    //мутация
    for (int i = 0; i < genomLength; i++) {
        int t = Math.abs(rand.nextInt()) % limitCount;
        vec[i] = ((Math.abs(rand.nextInt()) % 20) == 0) ? (vec[i] == t) ? (t + 1) % limitCount
: t : vec[i];
    }
    return vec;
} //Процедура скрещивания

private boolean getLimit(int n) {
    int sum = 0;
    for (int i = 0; i < genomLength; i++) {
        sum += population[n][i] * volume[i];
    }
    return (sum < limitVolumes);
}

private int getVolume(int i) {
    int sum = 0;
    for (int j = 0; j < genomLength; j++) {
        sum += population[i][j] * volume[j];
    }
    return sum;
}

private int avgfitnes() {
    int sum = 0;
    for (int i = 0; i < individualCount; i++) {
        sum += sortFitnes[i];
    }
    return sum / individualCount;
}

}

public class BrestTerespol {

```

```
public static void main(String[] args) {  
    GeneticAlgo p = new GeneticAlgo("TRUNCATING", "UNIFORM");  
    p.run();  
}  
}
```