

## An Evolution Program

### Objective

Solve the following optimization of the 0/1 Knapsack Problem.

### Given

- Weights,  $W_i$
- Profits,  $P_i$
- Capacity,  $C$ ,

### Find a binary vector

$$\vec{x} = \langle x_1, x_2, \dots, x_n \rangle, x_i \in \{0, 1\},$$

which maximizes

$$P(x) = \sum_i^n (x_i \cdot P_i),$$

subject to

$$\sum_i^n (x_i \cdot W_i) \leq C.$$

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### Test Data

The difficulty of the knapsack problem can be contributed to by the level of correlation of the Profits  $P_i$  and Weights  $W_i$ .

This project uses a weakly correlated test data set to provide some challenge while still remaining solvable using penalty methods.

For each  $i = 1..n$

$$W_i = rand.float(1, v),$$

$$P_i = \max(0, W_i + rand.float(-r, r))$$

### Test Data Parameters

Defaults shown.

- Number of items to choose from,  $n = 20$
- Upper bound for weight,  $v = 10$

- Profit correlation factor,  $r = 5$
  - Capacity,  $C = 40$
- 

## Example Menu

```

=====
                Problems To Solve
=====
        1 - Binary Knapsack
=====

Which problem? [1]:
=====
Enter Number of Items [20]:
=====
Enter Maximum Weight of One Item [10.0]:
=====
Enter Profit Correlation Factor [5.0]:
=====
Enter Knapsack Capacity [40.0]:

=====
                Penalty Methods
=====
        1 - Logarithmic
        2 - Absolute
        3 - Dynamic
=====

Which Penalty Method? [1]:
=====
Enter Rho for Logarithmic Penalty [1.3]:

=====
                Selection Mechanisms
=====
        1 - Proportional
        2 - Truncation
        3 - Tournament (Deterministic)
        4 - Tournament (Stochastic)
        5 - Linear Ranking
=====

```

```

=====
Which mechanism? [1]: 4
=====
Enter Probability of Fittest Winner [0.9]:

=====
                Crossover Methods
=====
        1 - Single Point
        2 - P-Uniform
        3 - Majority Voting
=====

=====
Which Crossover Method? [1]:
=====
Enter Probability of Crossover [0.65]:
=====
Population Size [30]:
=====
Probability of Mutation [0.05]:
=====
Max Generations [50]:
=====
Maximize [Y/n]:
=====
Single run? [y/N]:
=====
Collect stats? [Y/n]: Y
=====
Number of Runs [30]:

```

### Example Profits | Weights

Profits	Weights
[ 1.41120996	4.66918325]
[ 0.98976516	1.49829436]
[ 9.7947617	8.0968139 ]
[ 8.15973775	3.58574666]
[ 8.38648779	5.05315528]
[ 3.8183104	3.73521075]
[ 7.0378312	5.73759572]
[ 9.96465678	6.61430992]
[ 7.46429557	7.99097912]
[ 9.79907955	7.17617481]

```
[11.19767219  9.82844977]
[ 5.03030377  6.40734483]
[ 3.47146217  8.32571668]
[ 7.79765528  7.37780637]
[ 0.          1.24781212]
[ 5.58830843  9.13840496]
[ 5.53806313  5.04914366]
[ 0.          2.07032189]
[ 7.50751635  8.51770158]
[ 0.2206161   2.82023404]]
```

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## Example Default Options

```
Options: {
  "Crossover_Method": "<class 'binary_knapsack.crossover_method.single_point.SinglePoint'>",
  "Penalty_Method": "<class 'binary_knapsack.penalty_method.logarithmic.LogarithmicPenalty'>",
  "Select_Mechanism": "<class 'binary_knapsack.selection_mechanism.proportional.Proportional'>",
  "crossover_parameters": {
    "p_c": 0.65
  },
  "maximize": true,
  "p_m": 0.05,
  "penalty_parameters": {
    "rho": 1.3
  },
  "pop_size": 30,
  "problem_instance": "Profits\t\tWeights\n[[ 1.41120996  4.66918325]\n [ 0.98976516  1.49821412]
  "problem_parameters": {
    "capacity": 40.0,
    "dims": 20,
    "max_item_weight": 10.0,
    "profit_correlation_factor": 5.0
  },
  "selection_parameters": {},
  "t_max": 50
}
```

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## Representation

- Chromosomes (input vector  $\vec{x}$ ) made up of bit-string values, e.g.  $\langle 10010101101101100110 \rangle$ .
  - The  $i^{th}$  item is selected if and only if  $x_i = 1$ .
-

## Evolution Program Parameters

Defaults shown.

- Population size,  $N = 300$
  - Mutation probability,  $p_m = 0.05$
  - Maximum generations,  $t_{max} = 50$
- 

## Penalty Methods

### Logarithmic Penalty

I chose this static penalty method because of its ease of use and proven performance.

$$Penalty(x) = \log_2(1 + \rho \cdot (\sum_i^n x_i \cdot W_i - C))$$

### Parameters for Dynamic Penalty

- Scalar of degree of violation,  $\rho = 1.3$

### Results

Stats over 30 runs:

Best Overall: (Score: 56.96086516860768 :: [0 0 1 1 1 1 1 1 0 1 0 0 0 0 0 0 0 0 0] => cost

Mean Best Fitness: 53.06670943662913

Standard Deviation of Best Fitness: 1.771948255995935

Mean Generation Best Was Acheived: 31.5

Standard Deviation of Generations: 13.037765657248688

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### Absolute Penalty

Overage of any constraint leads to a zero fitness.

I chose this static penalty method because of its ease of use and to prove penalization works.

**Results** I could not get results due to the following error:

```
File ".\src\binary_knapsack\selection_mechanism\proportional.py", line 49, in _generate_pr
    assert self.sum_of_fitnesses > 0
           ~~~~~~
```

AssertionError

The absolute penalty method was not able to acheive feasible results.

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## Dynamic Penalty

As generations go by, the penalty for the same violation goes up.

$$Penalty(x) = (C * t)^{alpha} * \sum_j^m f_j(x)^{beta}$$

I chose this dynamic penalty method to test the claims that this method leads to premature convergence.

### Parameters for Dynamic Penalty

- Scalar of generation number,  $C = 0.5$
- Exponent of temporal harshening,  $alpha = 2.0$
- Exponent of degree of violation,  $beta = 2.0$

### Results

Stats over 30 runs:

Best Overall: (Score: 55.5352304177474 :: [0 1 0 1 1 0 1 1 0 1 1 0 0 0 0 0 0 0 0] => cost:

Mean Best Fitness: 52.74168473724467

Standard Deviation of Best Fitness: 1.3731685279839918

Mean Generation Best Was Acheived: 29.933333333333334

Standard Deviation of Generations: 13.01520478346597

This did not perform as well as logarithmic, and also did not acheive feasible results sometimes.

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## Selection Mechanisms

### Proportional Selection

An individual's probability of advancement equal to fitness score over sum of all fitness scores.

$$P(x_i) = f(x_i) / \sum_j^N f(x_j)$$

### Results

Stats over 30 runs:

Best Overall: (Score: 56.96086516860768 :: [0 0 1 1 1 1 1 1 0 1 0 0 0 0 0 0 0 0 0] => cost:

Mean Best Fitness: 53.09012691182008

Standard Deviation of Best Fitness: 1.4157700470325698  
Mean Generation Best Was Acheived: 30.3  
Standard Deviation of Generations: 12.607273033187365

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### Truncation Selection

The top  $\tau\%$  are selected for advancement. The next generation is sampled uniformly random with replacement from the top  $\tau\%$ .

#### Parameters for Truncation Selection

- Top tau,  $\tau = 0.4$

### Results

Stats over 30 runs:  
Best Overall: (Score: 56.96086516860768 :: [0 0 1 1 1 1 1 1 0 1 0 0 0 0 0 0 0 0] => cost:  
Mean Best Fitness: 51.930860569359396  
Standard Deviation of Best Fitness: 5.591743945371317  
Mean Generation Best Was Acheived: 26.866666666666667  
Standard Deviation of Generations: 16.390512160664443

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### Deterministic Tournament Selection

Pick two individuals from the current population uniformly randomly with replacement.

Advance the chromosome  $\vec{x}$  with the better fitness score.

Repeat  $N$  times.

### Results

Stats over 30 runs:  
Best Overall: (Score: 55.5352304177474 :: [0 1 0 1 1 0 1 1 0 1 1 0 0 0 0 0 0 0] => cost:  
Mean Best Fitness: 49.19210281339055  
Standard Deviation of Best Fitness: 8.154572784009153  
Mean Generation Best Was Acheived: 24.666666666666668  
Standard Deviation of Generations: 17.793881595150122

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### Stochastic Tournament Selection

Same as deterministic tournament selection, except the worse chromosome  $\vec{x}$  advances with a typically small random chance.

### Parameters for Stochastic Tournament Selection

- Probability of the higher performer winning,  $prob = 0.9$

### Results

Stats over 30 runs:

Best Overall: (Score: 56.96086516860768 :: [0 0 1 1 1 1 1 1 0 1 0 0 0 0 0 0 0 0] => cost

Mean Best Fitness: 46.870285204235365

Standard Deviation of Best Fitness: 5.924614709499177

Mean Generation Best Was Acheived: 13.533333333333333

Standard Deviation of Generations: 16.036901889773546

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### Linear Ranking Selection

Each chromosome  $\vec{x}$  is ranked from best to worst based on fitness score.

An individual's probability of advancement is proportional to its rank.

### Parameters for Linear Ranking Selection

- Expected # of copies of best  $\vec{x}$ ,  $max = 1.2$

### Results

Stats over 30 runs:

Best Overall: (Score: 54.89214401149931 :: [0 1 1 1 1 0 0 1 0 1 0 0 0 1 0 0 0 0 0] => cost

Mean Best Fitness: 47.97313785058804

Standard Deviation of Best Fitness: 2.9990674396662595

Mean Generation Best Was Acheived: 22.133333333333333

Standard Deviation of Generations: 15.622064168633058

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## Crossover Methods

### Single-Point Crossover

From two parents, randomly choose a cut-point and swap sides of each parent to create two children.

### Parameters for Single-Point Crossover

- Crossover probability,  $p_c = 0.65$



## Results

Stats over 30 runs:

Best Overall: (Score: 55.5352304177474 :: [0 1 0 1 1 0 1 1 0 1 1 0 0 0 0 0 0 0 0] => cost)

Mean Best Fitness: 52.81255744849033

Standard Deviation of Best Fitness: 1.328633982424607

Mean Generation Best Was Acheived: 30.33333333333332

Standard Deviation of Generations: 14.25794125702897

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## P-Uniform Crossover

From two parents and for each locus, randomly choose which parent to use at that locus. Choose the better parent with probability  $p$ . This process produces two children per set of parents.

### Parameters for P-Uniform Crossover

- Crossover probability,  $p_c = 0.65$
- Probability that the higher-performing parent donates their allele,  $p = 0.5$

## Results

Stats over 30 runs:

Best Overall: (Score: 56.96086516860768 :: [0 0 1 1 1 1 1 1 0 1 0 0 0 0 0 0 0 0 0] => cost)

Mean Best Fitness: 53.27474215638246

Standard Deviation of Best Fitness: 1.5440293311694724

Mean Generation Best Was Acheived: 30.466666666666665

Standard Deviation of Generations: 11.315868896770096

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## Majority Voting Crossover

From multiple parents and for each locus, deterministically choose the most common allele for that locus.

### Parameters for Majority Voting Crossover

- Crossover probability,  $p_c = 0.65$
- The number of parents to produce one child,  $parents = 4$

## Results

Stats over 30 runs:

Best Overall: (Score: 55.30528934249181 :: [0 0 0 1 1 0 0 1 0 1 1 0 0 1 0 0 0 0 0] => cost)

Mean Best Fitness: 50.370564380480324

Standard Deviation of Best Fitness: 2.406028580696979

Mean Generation Best Was Acheived: 31.833333333333332  
Standard Deviation of Generations: 14.758236871508586

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## Mutation Methods

### Gene-wise Mutation

Each bit ( $x_i$ ) of chromosome  $\vec{x}$  has equal but independent chance ( $p_m$ ) to mutate a small amount.

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## Termination Conditions

Simulate until one of the following has been met:

- Simulation reaches  $t_{max}$  generations.
  - Population fully converges.
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## Local Installation

From the project root, run `pip install -e src/`.

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See repository for full code.

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## Certification

I certify that this report is my own, independent work and that it does not plagiarize, in part or in full, any other work.

- Austin Hester