Introduction

Many engineering problems are characterized by several, often contradicting objectives, that have to be satisfied simultaneously. In many cases the standard approach of combining the objectives to a single one does not provide satisfactory results [1]. Evolutionary Algorithms are an attractive alternative in such cases [2], since they are a direct parallel search method and as such are able to obtain a Pareto trade-off surface between the objective function on a single run.

Strengthen Pareto Evolutionary Algorithm 2 (SPEA 2) is a Multi-Objective Evolutionary Algorithm (MOEA) developed by Computer Engineering (TIK), ETH Zurich, and reported to work well on many benchmark problems [3]. SPEA 2 is originally written in C, and developed under the PISA framework (Platform and Programming Language Independent Interface for Search Algorithms).

In order to allow easy use of SPEA 2 under MATLAB and avoid the file-based information exchange between the solver and the problem, the Institute of Control Systems, Hamburg University of Technology, combined SPEA 2 with the DTLZ variation function, modified the C files to interface with MATLAB and created an additional M file `moea.m` for user convenience.

Used abbreviations:

- MOEA: Multi-Objective Evolutionary Algorithm
- PISA: Platform and Programming Language Independent Interface for Search Algorithms
- SPEA 2: Strengthen Pareto Evolutionary Algorithm 2
1  Copyright and Legal Notes

PISA

Platform and Programming Language Independent Interface for Search Algorithms

Computer Engineering (TIK) ETH Zurich


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Combined MATLAB - C implementation of SPEA 2

Hamburg University of Technology, Institute of Control Systems

[http://www.tu-harburg.de/rts](http://www.tu-harburg.de/rts)

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2 Files and Options

The usage steps are described in Section 3.

spea2.dll

This is the compiled version of SPEA2 multi-objective optimization algorithm. It includes interface between MATLAB and C environment as well as the SPEA2 algorithm for selection and DTLZ functions for variation (mutation, uniform crossover and binary simulated crossover). The source codes for both selector (SPEA2) and variator (DTLZ) can be found on the PISA web-site.

The original PISA structure is removed and the selector and variator programs and combined to a single one.

The input arguments of spea2 are similar to those, described below. To see the full list, please read the source code of moea.m.

The normal stopping criteria of the function is the number of generations.

To recompile spea2 use:

```
mex spea2.c variator.c selector.c
```

moea.m

This program gets the user arguments, checks their validity and calls spea2, thus making the use of the program more intuitive.

The function call is in the form:

```
[genes, fitness] = moea('file_name', bounds, options)
```

file_name - name of the .M function, calculating the cost functions that should be minimized (!! SPEA2 always minimizes !!). The input arguments of this functions are the decision variables and the output arguments are the minimized functions.

bounds - \( N \times 2 \) matrix, containing the lower and upper boundaries for each decision variables, where N is the number of decision variables:

\[
\begin{align*}
\text{bounds} = [ & \text{Lower1, Upper1;} \\
& \text{Lower2, Upper2;} \\
& \text{...} \\
& \text{LowerN, UpperN}];
\end{align*}
\]

Where as \text{LowerK} and \text{UpperK} mean the correspondingly the lower and upper boundaries for decision variable K.
The number of decision variables is determined by the number of rows in bounds.

options - structure, containing the optimization options. options = moea() loads a predefined options set. That set can be easily modified to fit the user needs, by changing the values of the options, e.g., options.MaxGen = 100;

options.MaxGen - maximal number of generations (integer > 0);
options.PopSize - population size (integer > 0);
options.Dim - problem dimension (number of objective functions to be minimized simultaneously) (integer > 0);
options.Seed - seed for the random generator (real);
options.Tourn - number of individuals for Tournament selection (integer > 1);
options.i_mut_p - individual mutation probability (real ∈ [0, 1]);
options.i_rec_p - individual recombination probability (real ∈ [0, 1]);
options.v_mut_p - variable mutation probability (real ∈ [0, 1]);
options.v_swa_p - variable swap probability (real ∈ [0, 1]);
options.v_rec_p - variable recombination probability (real ∈ [0, 1]);
options.eta_mut - eta mutation (real ∈ [0, 1]);
options.eta_rec - eta recombination (real ∈ [0, 1]);
options.Display - if different then 0, then the number of remaining generations are shown during the optimization.

For more information on the meaning of the options, please refer to the PISA documentation.

3 Usage Steps

1. Install MOEA (copy in MATLAB\toolbox folder and set path to it). Depending on the MATLAB version delete either spea2.dll (MATLAB ver. 7.1+) or spea2.mex32 (all other). Alternatively you can re-compile SPEA2 yourself. For the purpose open the source folder in MATLAB and execute build_spea2.

2. Create your cost evaluation function (e.g. myFunct.m). The input parameter is a vector row and the function should return the results in a vector form (either row or column);

3. Define the optimization options. Call opt = moea() to get the standard options. Modify what needed, e.g. opt.MaxGen = 200; . Don’t forget to specify the correct problem dimension (number of cost functions), e.g. opt.Dim = 3;
4. Define boundaries for each design variable. The search is constrained within the boundaries, e.g.,

```matlab
bounds = [ 0 10; 0 2; -5 5 ];
```

The number of elements of `bounds` specifies the number of decision variables

5. Start the optimization

```matlab
[genes, fitness] = moea('myFunct', bounds, opt);
```

References

