

Model-Based Evolutionary Algorithms

Part 2: Linkage Tree Genetic Algorithm

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Evolutionary Algorithms

- Population-based, stochastic search algorithms
- **Exploitation**: selection
- **Exploration**: mutation & crossover

Model-Based Evolutionary Algorithms

- Population-based, stochastic search algorithms
- **Exploitation**: selection
- **Exploration**:
 - 1 Learn a (probabilistic) model from selected solutions
 - 2 Generate new solutions from the model (& population)

Gene-pool Optimal Mixing Evolutionary Algorithm

- Population-based, stochastic search algorithms
- **Exploitation**: selection (by replacement)
- **Exploration**:
 - 1 Learn a Family-Of-Subsets model
 - 2 Generate new solutions through optimal mixing

GOMEA: design objectives

- 1 Be able to efficiently **learn** dependency information (= linkage) between variables
- 2 Be able to efficiently **decide** between competing building blocks
- 3 **Transfer** all optimal building blocks from the parents to the offspring solution

Family Of Subsets (FOS) model

- Key idea is to identify **groups** of **problem variables** that together make an important contribution to the quality of solutions.
- These variable groups are interacting in a **non-linear** way and should be processed as a block = **building block**

FOS \mathcal{F}

Dependency structure generally called a **Family Of Subsets** (FOS).

- Let there be ℓ **problem variables** $x_0, x_1, \dots, x_{\ell-1}$.
- Let S be a set of all variable **indices** $\{0, 1, \dots, \ell - 1\}$.
- A FOS \mathcal{F} is a **set of subsets** of the set S .
- FOS \mathcal{F} is a **subset** of the **powerset** of S ($\mathcal{F} \subseteq \mathcal{P}(S)$).

Example Family Of Subsets (FOS) models:

- **Univariate** FOS structure

$$\mathcal{F} = \{\{0\}, \{1\}, \{2\}, \{3\}, \{4\}, \{5\}, \{6\}, \{7\}, \{8\}, \{9\}\}$$

- **Marginal Product** FOS Structure

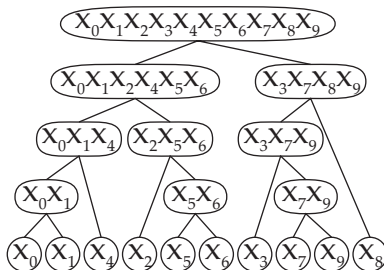
$$\mathcal{F} = \{\{0, 1, 2\}, \{3\}, \{4, 5\}, \{6, 7, 8, 9\}\}$$

- **Linkage Tree** FOS Structure

$$\begin{aligned} \mathcal{F} = & \{\{7, 5, 8, 6, 9, 0, 3, 2, 4, 1\}, \\ & \{7, 5, 8, 6, 9\}, \{0, 3, 2, 4, 1\}, \{7\}, \{5, 8, 6, 9\}, \\ & \{0, 3, 2, 4\}, \{1\}, \{5, 8, 6\}, \{9\}, \{0, 3\}, \{2, 4\}, \\ & \{5, 8\}, \{6\}, \{0\}, \{3\}, \{2\}, \{4\}, \{5\}, \{8\}\} \end{aligned}$$

Linkage Tree

- Problem variables in subset are considered to be **dependent** on each other but become **independent** in a child subset.
- \approx **Path** through dependency space, from **univariate** to **joint**.
- Linkage tree has ℓ **leaf** nodes (= single problem variables) and $\ell - 1$ **internal** nodes.



Linkage Tree Learning

- Start from **univariate** structure.
- Build linkage tree using **bottom-up** hierarchical clustering algorithm.
- **Similarity** measure:
 - ① Between individual variables X and Y : **mutual information** $I(X, Y)$.

$$I(X, Y) = H(X) + H(Y) - H(X, Y)$$

- ② Between cluster groups X_{Fi} and X_{Fj} : **average pairwise linkage** clustering (= unweighted pair group method with a arithmetic mean: UPGMA).

$$I^{UPGMA}(X_{Fi}, X_{Fj}) = \frac{1}{|X_{Fi}||X_{Fj}|} \sum_{X \in X_{Fi}} \sum_{Y \in X_{Fj}} I(X, Y).$$

$(H(X), H(Y), H(X, Y))$ are the marginal and joint entropies)

Linkage Tree Learning

- This agglomerative hierarchical clustering algorithm is computationally **efficient**.
- Only the mutual information between pairs of variables needs to be computed once, which is a $O(\ell^2)$ operation.
- The bottom-up hierarchical clustering can also be done in $O(\ell^2)$ computation by using the *reciprocal nearest neighbor chain* algorithm.
- note: commonly used bottom-up hierarchical clustering algorithms (*hclust* and *agnes* in R) have $O(\ell^3)$ complexity.

Optimal Mixing EA

- FOS linkage models specify the linked variables.
- A subset of the FOS is used as **crossover mask**
- Crossover is **greedy**: only **improvements** (or **equal**) are accepted.
- Each generation a new FOS model is build from selected solutions.
- For each solution in the population, **all subsets** of the FOS are tried with a **donor** solution randomly picked from the population
- **Recombinative OM (ROM)** and **Gene-pool OM (GOM)**
 - ▶ ROM is **GA-like**: select **single** donor solution to perform OM with
 - ▶ GOM is **EDA-like**: select **new** donor solution for each subset in OM

Gene-pool Optimal Mixing EA

GOMEA ()

```
Pop ← InitPopulation()
while NotTerminated(Pop)
  FOS ← BuildFOS(Pop)
  forall Sol ∈ Pop
    forall SubSet ∈ FOS
      Donor ← Random(Pop)
      Sol ← GreedyRecomb(Sol, Donor, SubSet, Pop)
return Sol
```

GreedyRecomb (Sol, Donor, SubSet, Pop)

```
NewSol ← ReplaceSubSetValues(Sol, SubSet, Donor)
if ImprovementOrEqual(NewSol, Sol)
  then Sol ← NewSol
return Sol
```

Recombinative Optimal Mixing EA

ROMEAE ()

```
Pop ← InitPopulation()
while NotTerminated(Pop)
    FOS ← BuildFOS(Pop)
    forall Sol ∈ Pop
        Donor ← Random(Pop)
        forall SubSet ∈ FOS
            Sol ← GreedyRecomb(Sol, Donor, SubSet, Pop)
return Sol
```

GreedyRecomb (Sol, Donor, SubSet, Pop)

```
NewSol ← ReplaceSubSetValues(Sol, SubSet, Donor)
if ImprovementOrEqual(NewSol, Sol)
    then Sol ← NewSol
return Sol
```

Optimal Mixing

- Characteristic of **Optimal Mixing** is the use of intermediate function evaluations (inside variation)
- Can be regarded as **greedy improvement** of existing solutions
- Coined **Optimal Mixing** because better instances for substructures are immediately accepted and not dependent on noise coming from other parts of the solution
- Building block competition **no longer** a stochastic decision making problem that requires a sizable minimal population size
- Population sizes in GOMEA much **smaller** than in GAs or EDAs.

Linkage Tree Genetic Algorithm

- The LTGA is an instance of **GOMEA** that uses a Linkage Tree as FOS model
- **Each generation** a new **hierarchical cluster tree** is build.
- For each solution in population, traverse **tree** starting at the top.
- Nodes (= clusters) in the linkage tree used as **crossover masks**.
- Select random donor solution, and its values at the crossover mask **replace** the variable **values** from the **current** solution.
- Evaluate new solution and **accept** if better/equal, otherwise **reject**.

Convergence model

Univariate FOS model on onemax problem

- ℓ : string length
- n : population size
- $p(t)$: proportion bit '1' at generation t
- $q(t)$: proportion bit '0' at generation t

Bit '0' only survive if parent and donor both have a '0' at that index:

- $q(t + 1) = q^2(t)$
- $p(t) = 1 - [1 - p(0)]^{2^t}$

Number of function evaluations FE :

- In 1 generation:

$$FE = 2 p(t) ([1 - p(t)] \times \ell \times n$$

- After g generations:

$$FE = \sum_{t=0}^g 2 p(t) ([1 - p(t)] \times \ell \times n$$

- After convergence g_{conv} :

$$FE = 2 [1 - p(0)] \times \ell \times n$$

- Initial random population ($p(0) = 0.5$):

$$FE = \ell \times n$$

 \Rightarrow

$$O(\ell \log \ell)$$

$$\begin{aligned}
& \sum_{t=0}^g p(t)([1 - p(t)]) \\
= & \sum_{t=0}^g q(t)([1 - q(t)]) \\
= & q(0)[1 - q(0)] + q(1)[1 - q(1)] + \dots + q(g)[(1 - q(g))] \\
= & q(0) - q(1) + q(1) - q(2) + \dots - q(g) + q(g) - q^2(g) \\
= & q(0) - [q(0)]^{2^{(g+1)}}
\end{aligned}$$

$$g_{conv} : \rightarrow q(0)$$

Minimal population size

Need to have at least one bit '1' at each index:

$$\text{Prob}[\text{success}] = [1 - (1 - p(0))^n]^\ell$$

$$\approx 1 - \ell [1 - p(0)]^n$$

$$1 - 0.01 = 1 - \ell \left[1 - \frac{1}{2}\right]^n$$

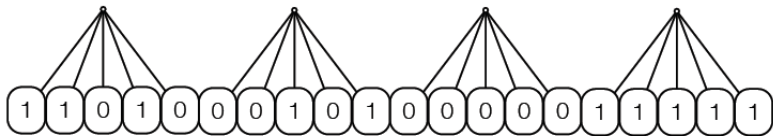
$$n = \log_2(100\ell)$$

$$n = O(\log \ell)$$

Deceptive Trap Function

Interacting, non-overlapping, deceptive groups of variables.

$$f_{\text{DT}}(x) = \sum_{i=0}^{l-k} f_{\text{DT}}^{\text{sub}}(x_{(i, \dots, i+k-1)})$$



Nearest-neighbor NK-landscape

- **Overlapping**, neighboring random subfunctions

$$f_{\text{NK-S1}}(x) = \sum_{i=0}^{l-k} f_{\text{NK}}^{\text{sub}}(x_{(i, \dots, i+k-1)}) \quad \text{with } f_{\text{NK}}^{\text{sub}}(x_{(i, \dots, i+k-1)}) \in [0..1]$$

- eg. 16 subsfcts, length $k = 5$, overlap $o = 4 \Rightarrow$ stringlength $\ell = 20$

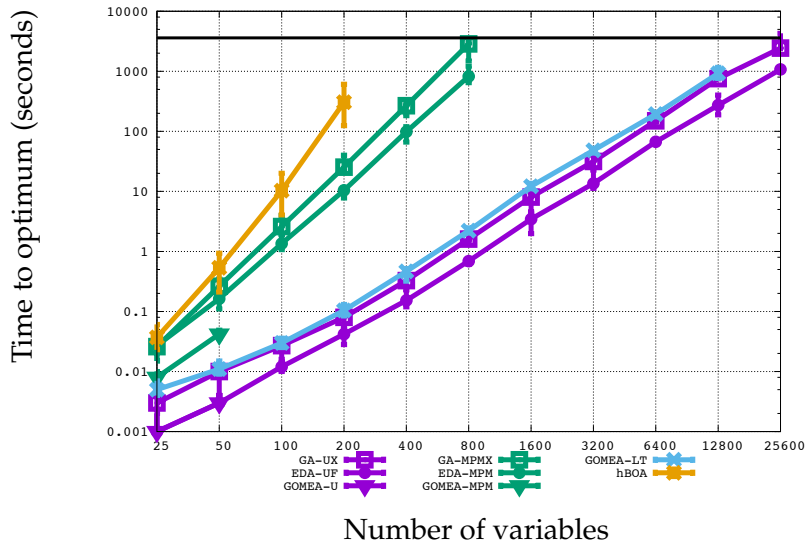


- **Global optimum** computed by dynamic programming
- Benchmark function: **structural information is not known !**
- \Rightarrow **Randomly shuffled** variable indices.

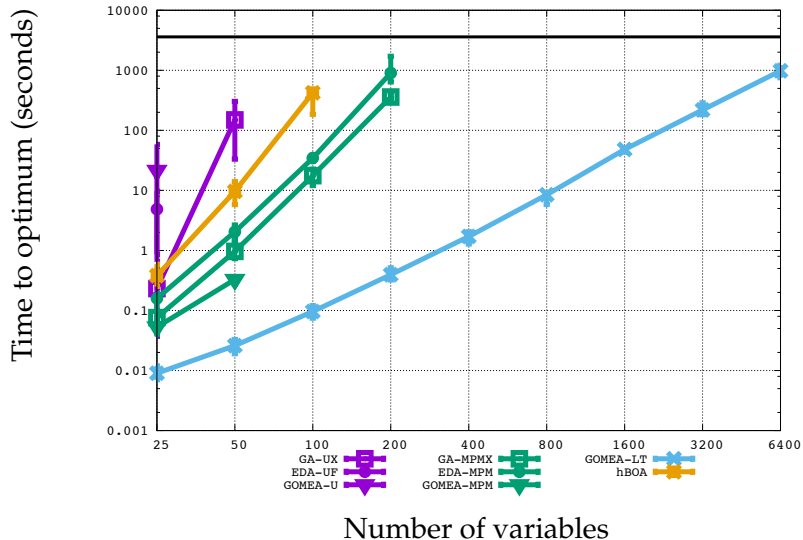
Experiments

- Compare GA, EDA, and GOMEA while each are learning the Marginal Product (MP) *FOS* structure, and GOMEA learning the Linkage Tree (LT) as *FOS* structure.
- Note:
 - ▶ EDA using MP = Extended Compact GA (ECGA).
 - ▶ GOMEA using LT = Linkage Tree Genetic Algorithm (LTGA).
 - ▶ hBOA = EDA learning a Bayesian network each generation.

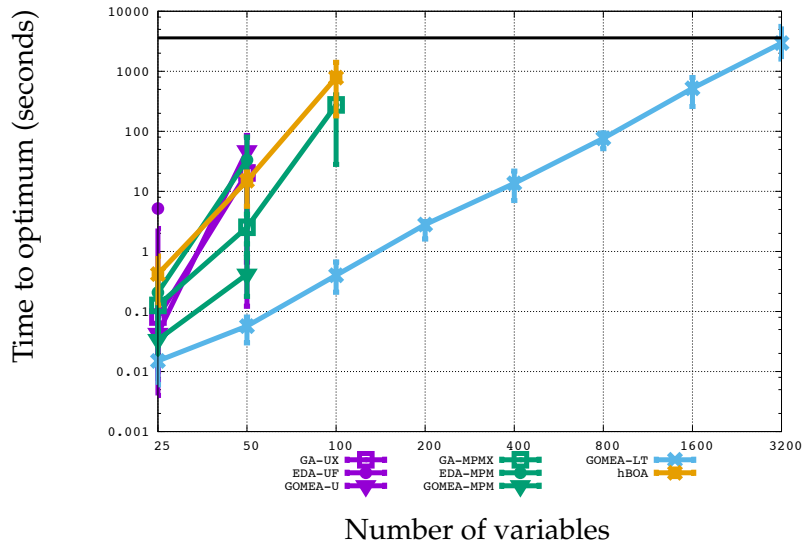
Experiments - Onemax



Experiments - Deceptive trap



Experiments - Overlapping NK



Experiments

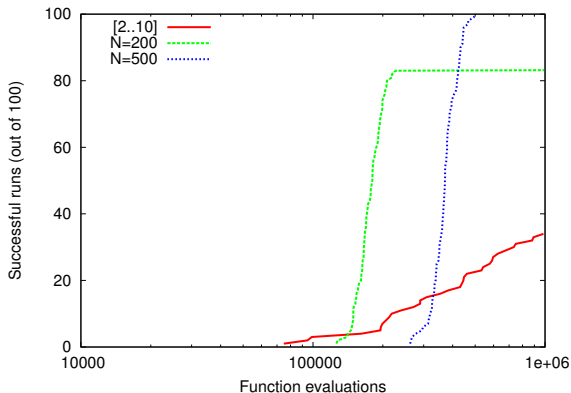
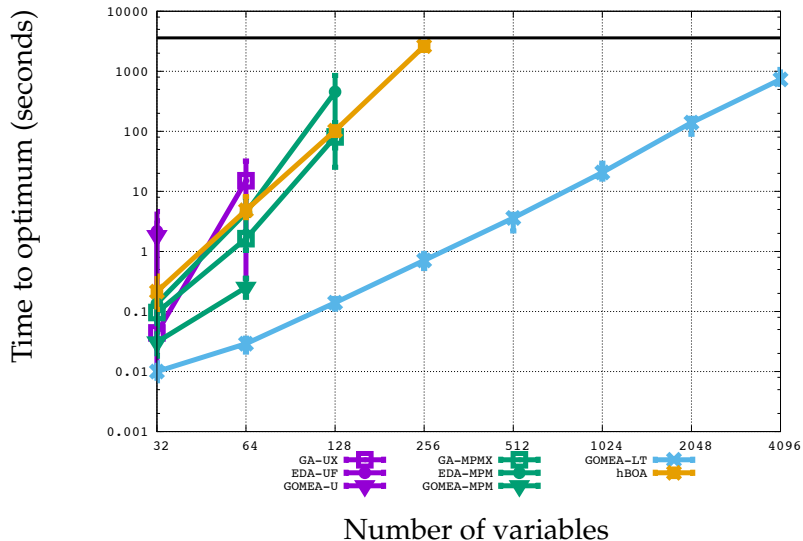


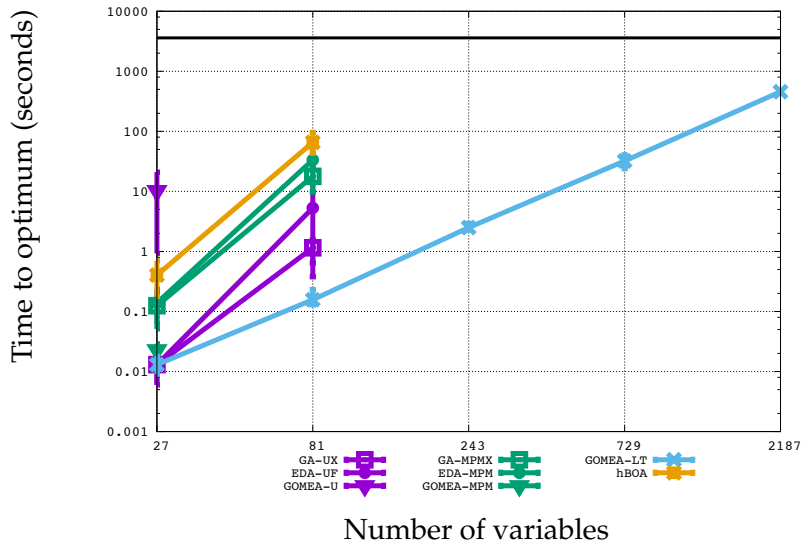
Figure: LTGA vs. ILS: 100 NK problems

Iterated Local Search: perturbation size each time randomly picked between 2 and 10 bits (= better than any fixed value).

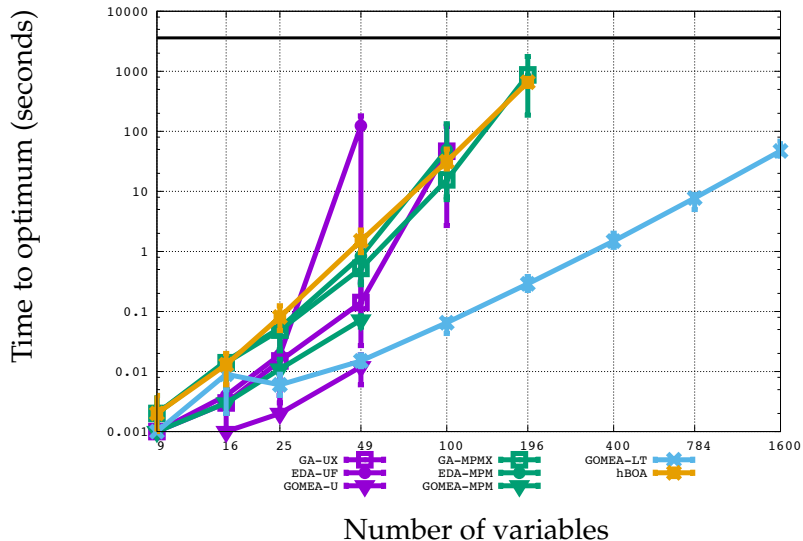
Experiments - HIFF



Experiments - HTrap



Experiments - MAX-CUT 2D square grid



Conclusions¹

- “Blind” Evolutionary Algorithms are **limited** in their capability to **detect** and **exploit** partial solutions (building blocks).
- Optimal Mixing Evolutionary Algorithms efficiently **learn** important building blocks and efficiently **decide** between competing building blocks
- **Linkage Tree** appears to be good compromise between FOS model complexity and search efficiency.

¹http://homepages.cwi.nl/~bosman/source_code.php