Probabilistic Model-Building Genetic Algorithms

a.k.a. Estimation of Distribution Algorithms a.k.a. Iterated Density Estimation Algorithms

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Foreword

Motivation

- □ Genetic and evolutionary computation (GEC) popular.
- □ Toy problems great, but difficulties in practice.
- □ Must design new representations, operators, tune, ...

This talk

- □ Discuss a promising direction in GEC.
- $\hfill\square$ Combine machine learning and GEC.
- Create practical and powerful optimizers.

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Overview

- Introduction
 - Black-box optimization via probabilistic modeling.
- Probabilistic Model-Building GAs
 - □ Discrete representation
 - $\hfill\square$ Continuous representation
 - □ Computer programs (PMBGP)
 - Permutations
- Conclusions

Problem Formulation

- Input
 - □ How do potential solutions look like?
 - □ How to evaluate quality of potential solutions?
- Output
 - \Box Best solution (the optimum).
- Important
 - $\hfill\square$ No additional knowledge about the problem.

Why View Problem as Black Box?

Advantages

- □ Separate problem definition from optimizer.
- \Box Easy to solve new problems.
- □ Economy argument.

Difficulties

- □ Almost no prior problem knowledge.
- □ Problem specifics must be learned automatically.
- □ Noise, multiple objectives, interactive evaluation.

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Typical Situation

Previously visited solutions + their evaluation:

#	Solution	Evaluation
1	00100	1
2	11011	4
3	01101	0
4	10111	3

Question: What solution to generate next?

Representations Considered Here

- Start with
 - □ Solutions are n-bit binary strings.
- Later
 - Real-valued vectors.
 - Program trees.
 - Permutations

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Many Answers

- Hill climber
 - $\hfill\square$ Start with a random solution.
 - □ Flip bit that improves the solution most.
 - □ Finish when no more improvement possible.
- Simulated annealing
 - □ Introduce Metropolis.
- Probabilistic model-building GAs
 - □ Inspiration from GAs and machine learning (ML).

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Probabilistic Model-Building GAs



...replace crossover+mutation with learning and sampling probabilistic model

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Implicit vs. Explicit Model

- GAs and PMBGAs perform similar task
 - □ Generate new solutions using probability distribution based on selected solutions.
- GAs
 - Variation defines implicit probability distribution of target population given original population and variation operators (crossover and mutation).
- PMBGAs
 - Explicit probabilistic model of selected candidate solutions is built and sampled.

Other Names for PMBGAs

- Estimation of distribution algorithms (EDAs) (Mühlenbein & Paass, 1996)
- Iterated density estimation algorithms (IDEA) (Bosman & Thierens, 2000)

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What Models to Use?

- Start with a simple example
 Probability vector for binary strings.
- Later
 - □ Dependency tree models (COMIT).
 - □ Bayesian networks (BOA).
 - □ Bayesian networks with local structures (hBOA).

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Probability Vector

- Assume *n*-bit binary strings.
- Model: Probability vector $p=(p_1, ..., p_n)$
 - \square p_i = probability of 1 in position *i*
 - $\hfill\square$ Learn p: Compute proportion of 1 in each position.
 - \Box Sample p: Sample 1 in position *i* with prob. p_i

Example: Probability Vector

(Mühlenbein, Paass, 1996), (Baluja, 1994)



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Probability Vector PMBGAs

- PBIL (Baluja, 1995)
 - □ Incremental updates to the prob. vector.
- Compact GA (Harik, Lobo, Goldberg, 1998)
 - Also incremental updates but better analogy with populations.
- UMDA (Mühlenbein, Paass, 1996)
 What we showed here.
- DEUM (Shakya et al., 2004)
- All variants perform similarly.

Probability Vector Dynamics

- Bits that perform better get more copies.
- And are combined in new ways.
- But context of each bit is ignored.
- Example problem 1: Onemax

$$f(X_1, X_2, \dots, X_n) = \sum_{i=1}^n X_i$$

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Probability Vector on Onemax



Probability Vector: Ideal Scale-up

- O(n log n) evaluations until convergence
 - □ (Harik, Cantú-Paz, Goldberg, & Miller, 1997)
 - □ (Mühlenbein, Schlierkamp-Vosen, 1993)
- Other algorithms
 - □ Hill climber: O(n log n) (Mühlenbein, 1992)
 - \Box GA with uniform: approx. O(n log n)
 - \Box GA with one-point: slightly slower

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When Does Prob. Vector Fail?

- Example problem 2: Concatenated traps
 - □ Partition input string into disjoint groups of 5 bits.
 - □ Groups contribute via trap (ones=number of ones):

 $trap(ones) = \begin{cases} 5 & \text{if } ones = 5\\ 4 - ones & \text{otherwise} \end{cases}$

- \Box Concatenated trap = sum of single traps
- □ Optimum: String 111...1

Trap-5



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Probability Vector on Traps



Why Failure?

Onemax:

- $\hfill\square$ Optimum in 111...1
- \Box 1 outperforms 0 on average.
- Traps: optimum in 11111, but
 f(0****) = 2
 f(1****) = 1.375
- So single bits are misleading.

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How to Fix It?

- Consider 5-bit statistics instead 1-bit ones.
- Then, 11111 would outperform 00000.
- Learn model
 - □ Compute p(00000), p(00001), ..., p(11111)
- Sample model
 - $\hfill\square$ Sample 5 bits at a time
 - □ Generate 00000 with p(00000), 00001 with p(00001), ...

Correct Model on Traps: Dynamics



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Good News: Good Stats Work Great!

- Optimum in O(n log n) evaluations.
- Same performance as on onemax!

Others

- \square Hill climber: O(n⁵ log n) = much worse.
- \Box GA with uniform: O(2ⁿ) = intractable.
- \Box GA with k-point xover: O(2ⁿ) (w/o tight linkage).

Challenge

- If we could learn and use relevant context for each position
 - $\hfill\square$ Find non-misleading statistics.
 - $\hfill\square$ Use those statistics as in probability vector.
- Then we could solve problems decomposable into statistics of order at most k with at most O(n²) evaluations!
 - \square And there are many such problems (Simon, 1968).

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What's Next?

- COMIT
 Use tree models
- Extended compact GA
 - □ Cluster bits into groups.
- Bayesian optimization algorithm (BOA)
 Use Bayesian networks (more general).

Beyond single bits: COMIT



How to Learn a Tree Model?

Mutual information:

$$I(X_i, X_j) = \sum_{a, b} P(X_i = a, X_j = b) \log \frac{P(X_i = a, X_j = b)}{P(X_i = a)P(X_j = b)}$$

- Goal
 - □ Find tree that maximizes mutual information between connected nodes.
 - □ Will minimize Kullback-Leibler divergence.
- Algorithm
 - □ Prim's algorithm for maximum spanning trees.

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Prim's Algorithm

- Start with a graph with no edges.
- Add arbitrary node to the tree.
- Iterate
 - $\hfill\square$ Hang a new node to the current tree.
 - Prefer addition of edges with large mutual information (greedy approach).
- Complexity: O(n²)

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Variants of PMBGAs with Tree Models

- COMIT (Baluja, Davies, 1997)
 Tree models.
- MIMIC (DeBonet, 1996)
 Chain distributions.
- BMDA (Pelikan, Mühlenbein, 1998)
 Forest distribution (independent trees or tree)

Beyond Pairwise Dependencies: ECGA

- Extended Compact GA (ECGA) (Harik, 1999).
- Consider groups of string positions.



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Learning the Model in ECGA

- Start with each bit in a separate group.
- Each iteration merges two groups for best improvement.



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How to Compute Model Quality?

- ECGA uses minimum description length.
- Minimize number of bits to store model+data:

 $MDL(M,D) = D_{Model} + D_{Data}$

• Each frequency needs (0.5 log *N*) bits:

$$D_{Model} = \sum_{g \in G} 2^{|g|-1} \log N$$

Each solution X needs -log p(X) bits:

$$D_{Data} = -N \sum_{X} p(X) \log p(X)$$

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Sampling Model in ECGA

- Sample groups of bits at a time.
- Based on observed probabilities/proportions.
- But can also apply population-based crossover similar to uniform but w.r.t. model.

Building-Block-Wise Mutation in ECGA

- Sastry, Goldberg (2004); Lima et al. (2005)
- Basic idea
 - □ Use ECGA model builder to identify decomposition
 - $\hfill\square$ Use the best solution for BB-wise mutation
 - □ For each k-bit partition (building block)
 - Evaluate the remaining 2^{k-1} instantiations of this BB
 - Use the best instantiation of this BB
- Result (for order-k separable problems)
 - □ BB-wise mutation is $O(\sqrt{k} \log n)$ times faster than ECGA!
 - □ But only for separable problems (and similar ones).

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What's Next?

- We saw
 - □ Probability vector (no edges).
 - \Box Tree models (some edges).
 - □ Marginal product models (groups of variables).

Next: Bayesian networks

 $\hfill\square$ Can represent all above and more.

Bayesian Optimization Algorithm (BOA)

- Pelikan, Goldberg, & Cantú-Paz (1998)
- Use a Bayesian network (BN) as a model.
- Bayesian network
 - $\hfill\square$ Acyclic directed graph.
 - □ Nodes are variables (string positions).
 - □ Conditional dependencies (edges).
 - □ Conditional independencies (implicit).

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Example: Bayesian Network (BN)

- Conditional dependencies.
- Conditional independencies.



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BOA



Learning BNs

- Two things again:
 - \Box Scoring metric (as MDL in ECGA).
 - $\hfill\square$ Search procedure (in ECGA done by merging).

Learning BNs: Scoring Metrics

- Bayesian metrics
 - □ Bayesian-Dirichlet with likelihood equivallence

 $BD(B) = p(B) \prod_{i=1}^{n} \prod_{\pi_i} \frac{\Gamma(m'(\pi_i))}{\Gamma(m'(\pi_i) + m(\pi_i))} \prod_{x_i} \frac{\Gamma(m'(x_i, \pi_i) + m(x_i, \pi_i))}{\Gamma(m'(x_i, \pi_i))}$

Minimum description length metrics
 Bayesian information criterion (BIC)

$$BIC(B) = \sum_{i=1}^{n} \left(-H(X_i \mid \Pi_i)N - 2^{|\Pi_i|} \frac{\log_2 N}{2} \right)$$

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Learning BNs: Search Procedure

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- Start with empty network (like ECGA).
- Execute primitive operator that improves the metric the most (greedy).
- Until no more improvement possible.
- Primitive operators
 - □ Edge addition (most important).
 - □ Edge removal.
 - □ Edge reversal.

Learning BNs: Example



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BOA and **Problem** Decomposition

- Conditions for factoring problem decomposition into a product of prior and conditional probabilities of small order in Mühlenbein, Mahnig, & Rodriguez (1999).
- In practice, approximate factorization sufficient that can be learned automatically.

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Learning makes complete theory intractable.

BOA Theory: Population Sizing

Initial supply (Goldberg et al., 2001)

 Have enough stuff to combine.
 O(2^k)

 Decision making (Harik et al, 1997)

 Decide well between competing partial sols.
 O(√n log n)

 Drift (Thierens, Goldberg, Pereira, 1998)

 Don't lose less salient stuff prematurely.
 O(n)

 Model building (Pelikan et al., 2000, 2002)

 Find a good model.

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BOA Theory: Num. of Generations

- Two extreme cases, everything in the middle.
- Uniform scaling
 - □ Onemax model (Muehlenbein & Schlierkamp-Voosen, 1993)

 $O\left(\sqrt{n}\right)$

Exponential scaling

Domino convergence (Thierens, Goldberg, Pereira, 1998)

O(n)

Good News: Challenge Met!

- Theory
 Population sizing (Pelikan et al., 2000, 2002)
 Initial supply.
 Decision making.
 Drift.
 Model building.
 Number of iterations (Pelikan et al., 2000, 2002)
 Uniform scaling.
 Exponential scaling.
- BOA solves order-k decomposable problems in O(n^{1.55}) to O(n²) evaluations!

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BOA Siblings

- Estimation of Bayesian Networks Algorithm (EBNA) (Etxeberria, Larrañaga, 1999).
- Learning Factorized Distribution Algorithm (LFDA) (Mühlenbein, Mahnig, Rodriguez, 1999).

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Another Option: Markov Networks

- MN-FDA, MN-EDA (Santana; 2003, 2005)
- Similar to Bayes nets but with undirected edges.



Yet Another Option: Dependency Networks

- Estimation of dependency networks algorithm (EDNA)
 - □ Gamez, Mateo, Puerta (2007).
 - □ Use dependency network as a model.
 - □ Dependency network learned from pairwise interactions.
 - □ Use Gibbs sampling to generate new solutions.
- Dependency network
 - □ Parents of a variable= all variables influencing this variable.
 - □ Dependency network can contain cycles.

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Model Comparison



From single level to hierarchy

- Single-level decomposition powerful.
- But what if single-level decomposition is not enough?
- Learn from humans and nature
 - □ Decompose problem over multiple levels.
 - Use solutions from lower level as basic building blocks.
 - □ Solve problem hierarchically.

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Hierarchical Decomposition



Three Keys to Hierarchy Success

- Proper decomposition
 - □ Must decompose problem on each level properly.
- Chunking
 - □ Must represent & manipulate large order solutions.
- Preservation of alternative solutions
 - Must preserve alternative partial solutions (chunks).

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Hierarchical BOA (hBOA)

- Pelikan & Goldberg (2000, 2001)
- Proper decomposition
 - $\hfill\square$ Use Bayesian networks like BOA.
- Chunking
 - $\hfill\square$ Use local structures in Bayesian networks.
- Preservation of alternative solutions.
 - □ Use restricted tournament replacement (RTR).
 - \Box Can use other niching methods.

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Local Structures in BNs

- Look at one conditional dependency.
 2^k probabilities for k parents.
- Why not use more powerful representations for conditional probabilities?

⁴ X.	X_2X_3	$P(X_1=0 X_2X_3)$
	00	26 %
	01	44 %
A2 A3	10	15 %
	11	15 %

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Local Structures in BNs

- Look at one conditional dependency.
 2^k probabilities for k parents.
- Why not use more powerful representations for conditional probabilities?



Restricted Tournament Replacement

- Used in hBOA for niching.
- Insert each new candidate solution x like this:
 - □ Pick random subset of original population.
 - \Box Find solution y most similar to x in the subset.
 - \Box Replace y by x if x is better than y.

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Hierarchical Traps: The Ultimate Test

- Combine traps on more levels.
- Each level contributes to fitness.
- Groups of bits map to next level.



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hBOA on Hierarchical Traps



PMBGAs Are Not Just Optimizers

- PMBGAs provide us with two things
 - $\hfill\square$ Optimum or its approximation.
 - □ Sequence of probabilistic models.
- Probabilistic models
 - $\hfill\square$ Encode populations of increasing quality.
 - \Box Tell us a lot about the problem at hand.
 - □ Can we use this information?

Efficiency Enhancement for PMBGAs

- Sometimes O(n²) is not enough
 - □ High-dimensional problems (1000s of variables)
 - □ Expensive evaluation (fitness) function
- Solution
 - □ Efficiency enhancement techniques

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Efficiency Enhancement Types

- 7 efficiency enhancement types for PMBGAs
 - □ Parallelization
 - □ Hybridization
 - □ Time continuation
 - □ Fitness evaluation relaxation
 - □ Prior knowledge utilization
 - □ Incremental and sporadic model building
 - □ Learning from experience

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Promising Results with Discrete PMBGAs

- Artificial classes of problems
- Physics
- Bioinformatics
- Computational complexity and AI
- Others

Multi-objective PMBGAs

- Methods for multi-objective GAs adopted
 - Multi-objective mixture-based IDEAs (Thierens, & Bosman, 2001)
 - Another multi-objective BOA (from SPEA2 and mBOA) (Laumanns, & Ocenasek, 2002)
 - Multi-objective hBOA (from NSGA-II and hBOA) (Khan, Goldberg, & Pelikan, 2002) (Pelikan, Sastry, & Goldberg, 2005)
 - Regularity Model Based Multiobjective EDA (RM-MEDA) (Zhang, Zhou, Jin, 2008)

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Results: Artificial Problems

- Decomposition
 - □ Concatenated traps (Pelikan et al., 1998).
- Hierarchical decomposition
 Hierarchical traps (Pelikan, Goldberg, 2001).
- Other sources of difficulty
 - □ Exponential scaling, noise (Pelikan, 2002).

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BOA on Concatenated 5-bit Traps

hBOA on Hierarchical Traps



Results: Physics

- Spin glasses (Pelikan et al., 2002, 2006, 2008) (Hoens, 2005) (Santana, 2005) (Shakya et al., 2006)
 - $\hfill\square$ ±J and Gaussian couplings
 - $\hfill\square$ 2D and 3D spin glass
 - □ Sherrington-Kirkpatrick (SK) spin glass
- Silicon clusters (Sastry, 2001)
 Gong potential (3-body)

hBOA on Ising Spin Glasses (2D)



Results on 2D Spin Glasses

- Number of evaluations is $O(n^{1.51})$.
- Overall time is $O(n^{3.51})$.
- Compare O(n^{3.51}) to O(n^{3.5}) for best method (Galluccio & Loebl, 1999)
- Great also on Gaussians.

hBOA on Ising Spin Glasses (3D)



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hBOA on SK Spin Glass



Results: Computational Complexity, AI

- MAXSAT, SAT (Pelikan, 2002)
 - $\hfill\square$ Random 3CNF from phase transition.
 - □ Morphed graph coloring.
 - $\hfill\square$ Conversion from spin glass.
- Feature subset selection (Inza et al., 2001) (Cantu-Paz, 2004)

Results: Some Others

- Military antenna design (Santarelli et al., 2004)
- Groundwater remediation design (Arst et al., 2004)
- Forest management (Ducheyne et al., 2003)
- Nurse scheduling (Li, Aickelin, 2004)
- Telecommunication network design (Rothlauf, 2002)
- Graph partitioning (Ocenasek, Schwarz, 1999; Muehlenbein, Mahnig, 2002; Baluja, 2004)
- Portfolio management (Lipinski, 2005, 2007)
- Quantum excitation chemistry (Sastry et al., 2005)
- Maximum clique (Zhang et al., 2005)
- Cancer chemotherapy optimization (Petrovski et al., 2006)
- Minimum vertex cover (Pelikan et al., 2007)
- Protein folding (Santana et al., 2007)
- Side chain placement (Santana et al., 2007)

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Discrete PMBGAs: Summary

- No interactions
 Univariate models; PBIL, UMDA, cGA.
- Some pairwise interactions
 Tree models; COMIT, MIMIC, BMDA.
- Multivariate interactions
 Multivariate models: BOA, EBNA, LFDA.
- Hierarchical decomposition
 hBOA

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Discrete PMBGAs: Recommendations

Easy problems

□ Use univariate models; PBIL, UMDA, cGA.

- Somewhat difficult problems
 Use bivariate models; MIMIC, COMIT, BMDA.
- Difficult problems
 - □ Use multivariate models; BOA, EBNA, LFDA.
- Most difficult problems
 - □ Use hierarchical decomposition; hBOA.

Real-Valued PMBGAs

- New challenge
 - $\hfill\square$ Infinite domain for each variable.
 - □ How to model?
- 2 approaches
 - □ Discretize and apply discrete model/PMBGA
 - □ Create model for real-valued variables
 - Estimate pdf.

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PBIL Extensions: First Step

- SHCwL: Stochastic hill climbing with learning (Rudlof, Köppen, 1996).
- Model
 - □ Single-peak Gaussian for each variable.
 - □ Means evolve based on parents (promising solutions).
 - □ Deviations equal, decreasing over time.
- Problems
 - No interactions.
 - $\hfill\square$ Single Gaussians=can model only one attractor.
 - □ Same deviations for each variable.

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Use Different Deviations

- Sebag, Ducoulombier (1998)
- Some variables have higher variance.
- Use special standard deviation for each



Use Covariance

- Covariance allows rotation of 1-peak Gaussians.
- EGNA (Larrañaga et al., 2000)
- IDEA (Bosman, Thierens, 2000)



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How Many Peaks?

- One Gaussian vs. kernel around each point.
- Kernel distribution similar to ES.
- IDEA (Bosman, Thierens, 2000)



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Mixtures: Between One and Many

- Mixture distributions provide transition between one Gaussian and Gaussian kernels.
- Mixture types
 - □ Over one variable.
 - Gallagher, Frean, & Downs (1999).
 - □ Over all variables.
 - Pelikan & Goldberg (2000).
 - Bosman & Thierens (2000).
 - □ Over partitions of variables.
 - Bosman & Thierens (2000).
 - Ahn, Ramakrishna, and Goldberg (2004).

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Real-Coded BOA (rBOA)

- Ahn, Ramakrishna, Goldberg (2003)
- Probabilistic Model
 - □ Underlying structure: Bayesian network
 - □ Local distributions: Mixtures of Gaussians
- Also extended to multiobjective problems (Ahn, 2005)

Mixed BOA (mBOA)

- Mixed BOA (Ocenasek, Schwarz, 2002)
- Local distributions
 - \square A decision tree (DT) for every variable.
 - $\hfill\square$ Internal DT nodes encode tests on other variables
 - Discrete: Equal to a constant
 - Continuous: Less than a constant
 - □ Discrete variables:
 - DT leaves represent probabilities.
 - Continuous variables:
 - DT leaves contain a normal kernel distribution.

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Aggregation Pheromone System (APS)

- Tsutsui (2004)
- Inspired by aggregation pheromones
- Basic idea
 - $\hfill\square$ Good solutions emit aggregation pheromones
 - New candidate solutions based on the density of aggregation pheromones
 - Aggregation pheromone density encodes a mixture distribution

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Adaptive Variance Scaling

- Adaptive variance in mBOA
 - □ Ocenasek et al. (2004)
- Normal IDEAs
 - □ Bosman et al. (2006, 2007)
 - □ Correlation-triggered adaptive variance scaling
 - Standard-deviation ratio (SDR) triggered variance scaling

Real-Valued PMBGAs: Discretization

- Idea: Transform into discrete domain.
- Fixed models
 - \Box 2^k equal-width bins with k-bit binary string.
 - □ Goldberg (1989).
 - □ Bosman & Thierens (2000); Pelikan et al. (2003).
- Adaptive models
 - □ Equal-height histograms of 2k bins.
 - $\hfill\square$ k-means clustering on each variable.
 - □ Pelikan, Goldberg, & Tsutsui (2003); Cantu-Paz (2001).

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Real-Valued PMBGAs: Summary

- Discretization
 - □ Fixed
 - Adaptive
- Real-valued models
 - \Box Single or multiple peaks?
 - □ Same variance or different variance?
 - □ Covariance or no covariance?
 - □ Mixtures?
 - Treat entire vectors, subsets of variables, or single variables?

Real-Valued PMBGAs: Recommendations

- Multimodality?
 - □ Use multiple peaks.
- Decomposability?
 - $\hfill \Box$ All variables, subsets, or single variables.
- Strong linear dependencies?
 Covariance.
- Partial differentiability?
 - $\hfill\square$ Combine with gradient search.

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PMBGP (Genetic Programming)

New challenge

- □ Structured, variable length representation.
- □ Possibly infinitely many values.
- \Box Position independence (or not).
- Low correlation between solution quality and solution structure (Looks, 2006).
- Approaches
 - □ Use explicit probabilistic models for trees.
 - □ Use models based on grammars.

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PIPE

	Probabilistic incremental program evolution		
	(Salustowicz &	X	P(X)
	Schmidhuber, 1997)	sin	0.15
	Store frequencies of	+	0.35
	operators/terminals in nodes of a <i>maximum</i> tree	-	0.35
	Sampling generates tree	Х	0.15
	from top to bottom		

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eCGP

- Sastry & Goldberg (2003)
- ECGA adapted to program trees.
- Maximum tree as in PIPE.
- But nodes partitioned into groups.



BOA for GP

- Looks, Goertzel, & Pennachin (2004)
- Combinatory logic + BOA
 - $\hfill\square$ Trees translated into uniform structures.
 - □ Labels only in leaves.
 - □ BOA builds model over symbols in different nodes.
- Complexity build-up
 - $\hfill\square$ Modeling limited to max. sized structure seen.
 - $\hfill\square$ Complexity builds up by special operator.

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MOSES

- Looks (2006).
- Evolve demes of programs.
- Each deme represents similar structures.
- Apply PMBGA to each deme (e.g. hBOA).
- Introduce new demes/delete old ones.
- Use normal forms to reduce complexity.

PMBGP with Grammars

- Use grammars/stochastic grammars as models.
- Grammars restrict the class of programs.
- Some representatives
 - □ Program evolution with explicit learning (Shan et al., 2003)

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- □ Grammar-based EDA for GP (Bosman, de Jong, 2004)
- □ Stochastic grammar GP (Tanev, 2004)
- □ Adaptive constrained GP (Janikow, 2004)

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PMBGP: Summary

- Interesting starting points available.
- But still lot of work to be done.
- Much to learn from discrete domain, but some completely new challenges.
- Research in progress

PMBGAs for Permutations

- New challenges
 - \Box Relative order
 - □ Absolute order
 - Permutation constraints
- Two basic approaches
 - □ Random-key and real-valued PMBGAs
 - $\hfill\square$ Explicit probabilistic models for permutations

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Random Keys and PMBGAs

- Bengoetxea et al. (2000); Bosman et al. (2001)
- Random keys (Bean, 1997)
 - □ Candidate solution = vector of real values
 - □ Ascending ordering gives a permutation
- Can use any real-valued PMBGA (or GEA)
 - □ IDEAs (Bosman, Thierens, 2002)
 - □ EGNA (Larranaga et al., 2001)
- Strengths and weaknesses
 - □ Good: Can use any real-valued PMBGA.
 - $\hfill\square$ Bad: Redundancy of the encoding.

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Direct Modeling of Permutations

- Edge-histogram based sampling algorithm (EHBSA) (Tsutsui, Pelikan, Goldberg, 2003)
 - \Box Permutations of *n* elements
 - \square Model is a matrix $A = (a_{i,j})_{i,j=1,2,...,n}$
 - \Box a_{i,i} represents the probability of edge (i, j)
 - $\hfill\square$ Uses template to reduce exploration
 - □ Applicable also to scheduling

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ICE: Modify Crossover from Model

ICE

- □ Bosman, Thierens (2001).
- □ Represent permutations with random keys.
- □ Learn multivariate model to factorize the problem.
- □ Use the learned model to modify crossover.

Performance

□ Typically outperforms IDEAs and other PMBGAs that learn and sample random keys.

Multivariate Permutation Models

- Basic approach
 - □ Use any standard multivariate discrete model.
 - $\hfill\square$ Restrict sampling to permutations in some way.
 - □ Bengoetxea et al. (2000), Pelikan et al. (2007).
- Strengths and weaknesses
 - $\hfill\square$ Use explicit multivariate models to find regularities.
 - High-order alphabet requires big samples for good models.
 - □ Sampling can introduce unwanted bias.
 - Inefficient encoding for only relative ordering constraints, which can be encoded simpler.

Conclusions

- Competent PMBGAs exist
 - $\hfill\square$ Scalable solution to broad classes of problems.
 - □ Solution to previously intractable problems.
 - □ Algorithms ready for new applications.
- PMBGAs do more than just solve the problem
 - □ They provide us with sequences of probabilistic models.
 - $\hfill\square$ The probabilistic models tell us a lot about the problem.
- Consequences for practitioners
 - □ Robust methods with few or no parameters.
 - □ Capable of learning how to solve problem.
 - □ But can incorporate prior knowledge as well.
 - □ Can solve previously intractable problems.

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Starting Points

- World wide web
- Books and surveys
 - □ Larrañaga & Lozano (eds.) (2001). Estimation of distribution algorithms: A new tool for evolutionary computation. Kluwer.
 - Pelikan et al. (2002). A survey to optimization by building and using probabilistic models. Computational optimization and applications, 21(1), pp. 5-20.
 - Pelikan (2005). Hierarchical BOA: Towards a New Generation of Evolutionary Algorithms. Springer.
 - □ Lozano, Larrañaga, Inza, Bengoetxea (2006). Towards a New Evolutionary Computation: Advances on Estimation of Distribution Algorithms, Springer.
 - Pelikan, Sastry, Cantu-Paz (eds.) (2006). Scalable Optimization via Probabilistic Modeling: From Algorithms to Applications, Springer.

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Online Code (1/2)

- BOA, BOA with decision graphs, dependency-tree EDA <u>http://medal-lab.org/</u>
- ECGA, xi-ary ECGA, BOA, and BOA with decision trees/graphs <u>http://www.illigal.org/</u>
- mBOA http://jiri.ocenasek.com/
- PIPE <u>http://www.idsia.ch/~rafal/</u>
- Real-coded BOA <u>http://www.evolution.re.kr/</u>

Online Code (2/2)

- Demos of APS and EHBSA <u>http://www.hannan-u.ac.jp/~tsutsui/research-e.html</u>
- RM-MEDA: A Regularity Model Based Multiobjective EDA Differential Evolution + EDA hybrid http://cswww.essex.ac.uk/staff/gzhang/mypublication.htm
- Naive Multi-objective Mixture-based IDEA (MIDEA) Normal IDEA-Induced Chromosome Elements Exchanger (ICE) Normal Iterated Density-Estimation Evolutionary Algorithm (IDEA) <u>http://homepages.cwi.nl/~bosman/code.html</u>