



Theory of Estimation-of-Distribution Algorithms

PPSN 2024 Tutorial

1. Estimation-of- (EDAs)
Distribution Algorithms

2. **Theory**

1. Estimation-of- Distribution Algorithms (EDAs)

» **What** are they?

2. Theory

1. Estimation-of- Distribution Algorithms (EDAs)

- » **What** are they?
- » **Focus** on the most commonly theoretically studied ones

2. Theory

1. Estimation-of- Distribution Algorithms (EDAs)

- » **What** are they?
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2. Theory (Mathematical rigor)

1. Estimation-of- Distribution Algorithms (EDAs)

- » **What** are they?
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2. Theory (Mathematical rigor)

- » **What** is studied?

1. Estimation-of- Distribution Algorithms (EDAs)

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2. Theory (Mathematical rigor)

- » **What** is studied?
- » **Selection** of important results
 - ❖ **Understand** the basic idea

1. Estimation-of- Distribution Algorithms (EDAs)

- » **What** are they?
- » **Focus** on the most commonly theoretically studied ones

2. Theory (Mathematical rigor)

- » **What** is studied?
- » **Selection** of important results
 - ❖ **Understand** the basic idea

» **Depth over breadth**



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DOI: [10.1145/3377929.3389888](https://doi.org/10.1145/3377929.3389888)



(more **breadth**)



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(more **breadth**)



1806.05392





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(more **breadth**)

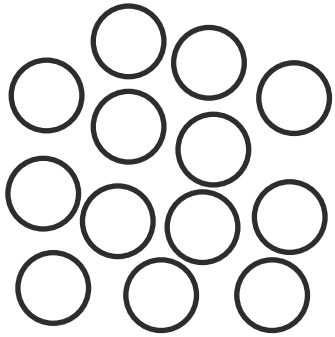


Sunday, Sep. 15

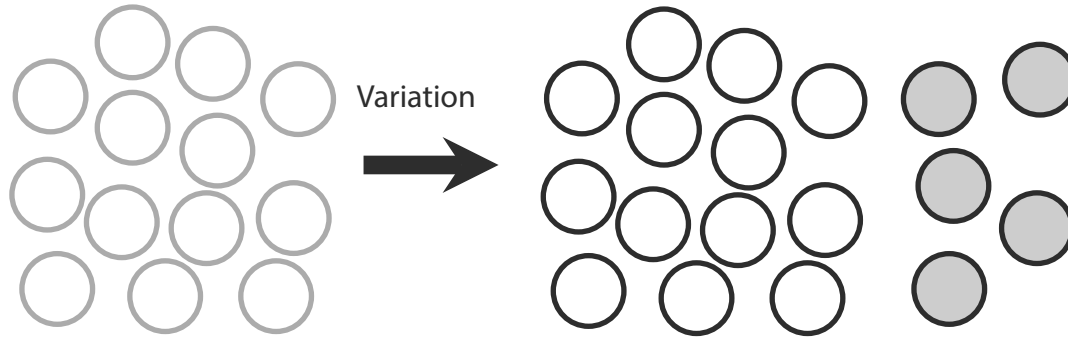
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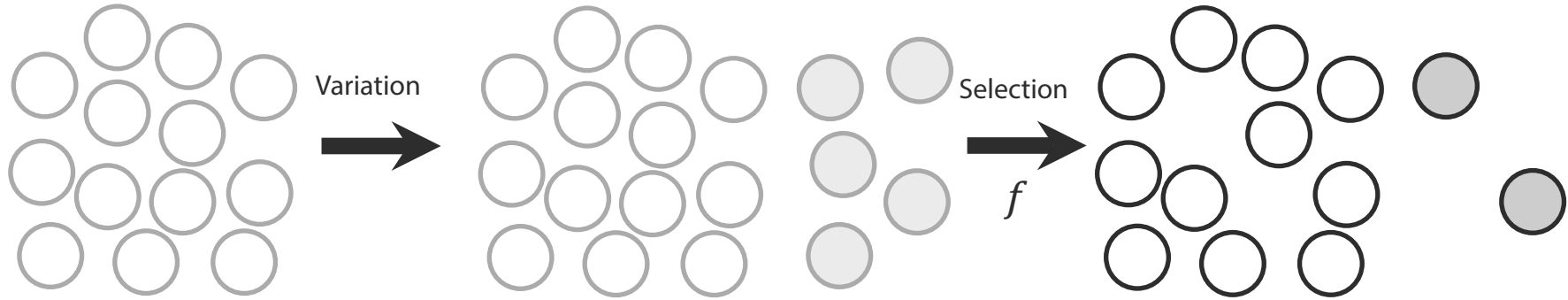
Evolutionary Algorithms



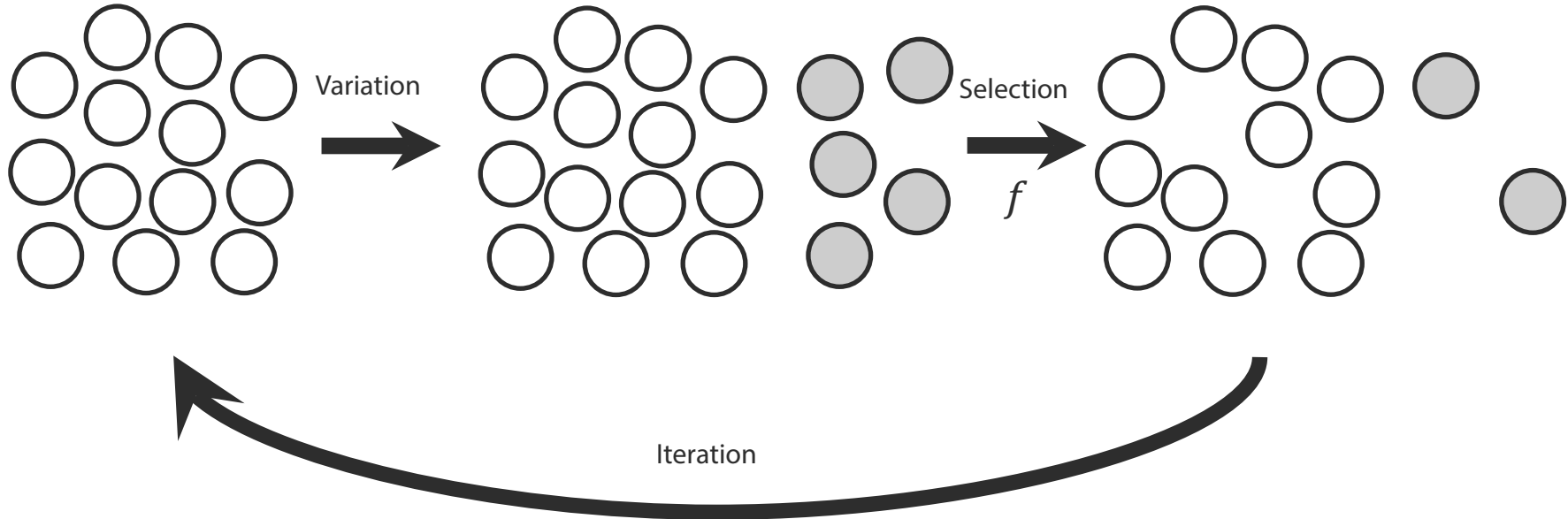
Evolutionary Algorithms



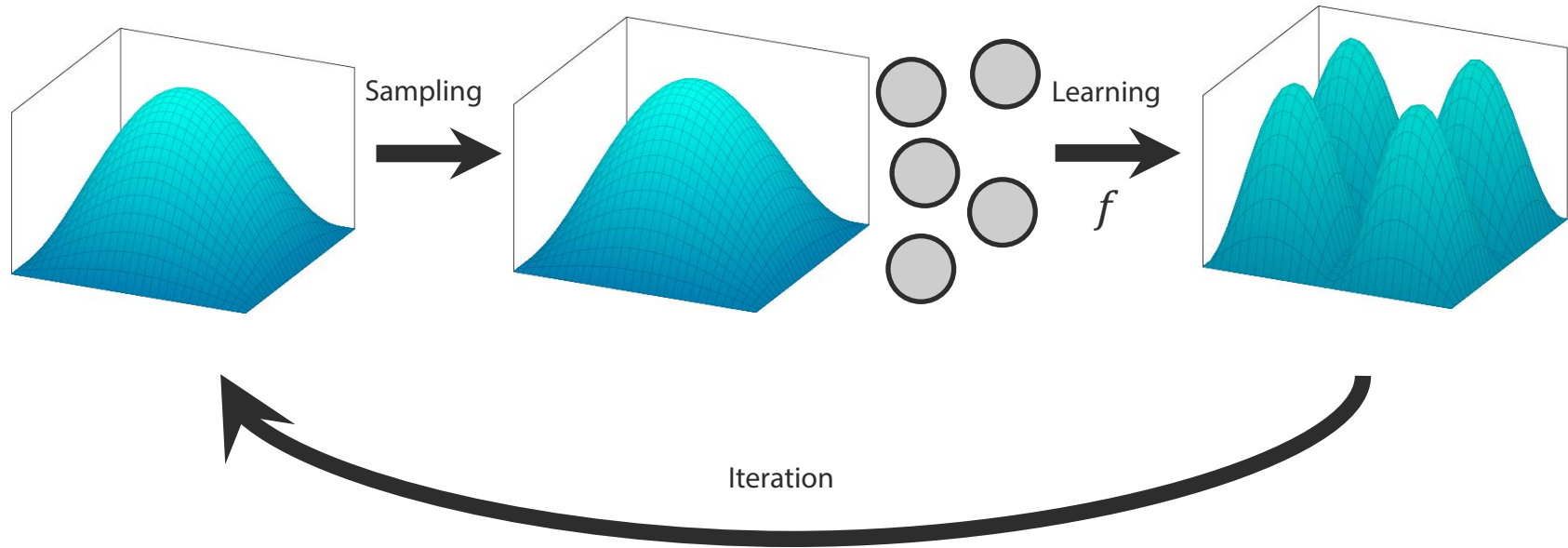
Evolutionary Algorithms



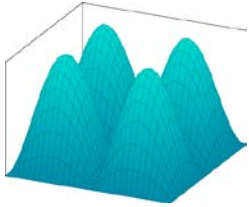
Evolutionary Algorithms



Estimation-of-Distribution Algorithms

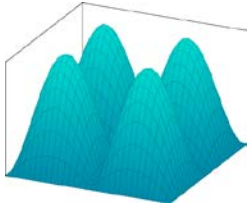


Probabilistic Models



general

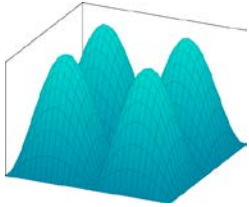
Probabilistic Models



general

x_1	x_2	x_3	x_4	x_5
0	0	0	0	0
0	0	0	0	1
0	0	0	0	2
		\vdots		

Probabilistic Models

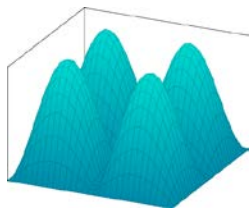


general

x_1	x_2	x_3	x_4	x_5
0	0	0	0	0
0	0	0	0	1
0	0	0	0	2
		\vdots		

$$\prod_{i \in [\#\text{vars}]} (\#\text{vals}(x_i)) - 1$$

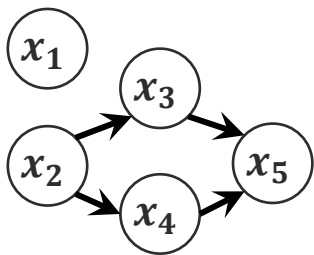
Probabilistic Models



general

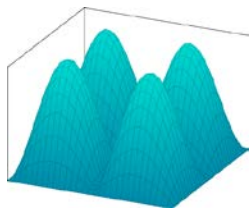
x_1	x_2	x_3	x_4	x_5
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0	0	0	0	2
		\vdots		

$$\prod_{i \in [\text{\#vars}]} (\text{\#vals}(x_i)) - 1$$



compressed, directed

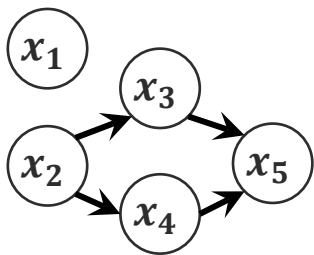
Probabilistic Models



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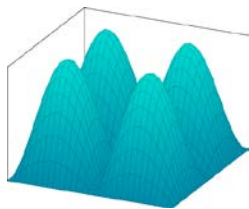
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compressed, directed

x_1	x_2	x_2	x_3	
0	0	0	0	
1	1	0	1	...
2	2	0	2	
\vdots	\vdots	\vdots		

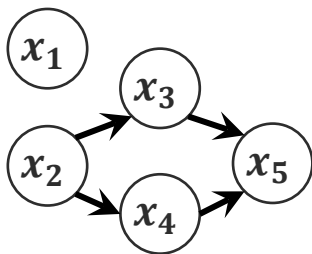
Probabilistic Models



general

x_1	x_2	x_3	x_4	x_5
0	0	0	0	0
0	0	0	0	1
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		⋮		

$$\prod_{i \in [\#vars]} (\#vals(x_i)) - 1$$

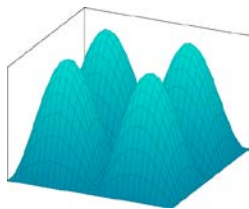


compressed, directed

x_1	x_2	x_3	x_4
0	0	0	0
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⋮	⋮	⋮	⋮

$$\sum_{i \in [\#vars]} (\#vals(x_i) - 1) \prod_{y \in \text{pred}(x_i)} \#vals(y)$$

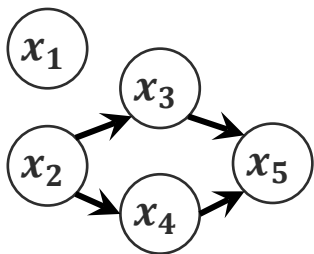
Probabilistic Models



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compressed, directed

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$$\sum_{i \in [\#vars]} (\#vals(x_i) - 1) \prod_{y \in \text{pred}(x_i)} \#vals(y)$$

Theory



univariate

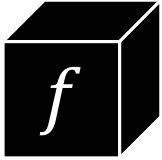
Theory



$$f: \{0,1\}^n \rightarrow \mathbf{R}$$

» Pseudo-Boolean optimization

Theory

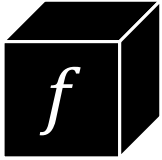


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» Pseudo-Boolean optimization

❖ **Global optimum** often 1^n

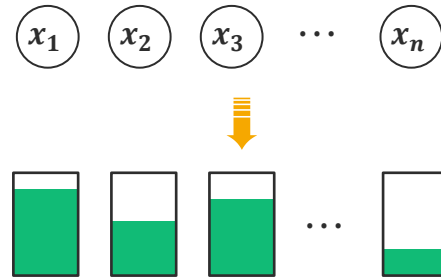
Theory



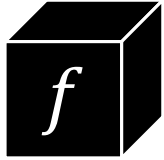
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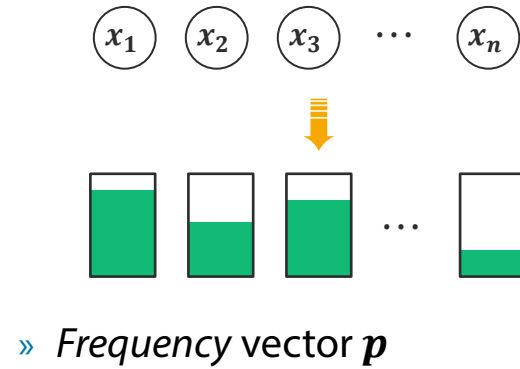
Theory



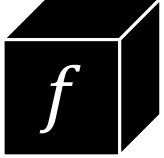
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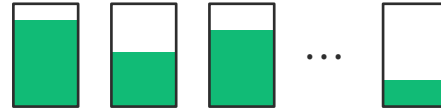
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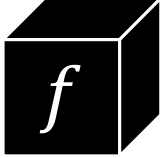
❖ **Global optimum** often 1^n



» *Frequency vector* \mathbf{p}

» \mathbf{p}_i ... probability to sample a 1 at i
(green mass)

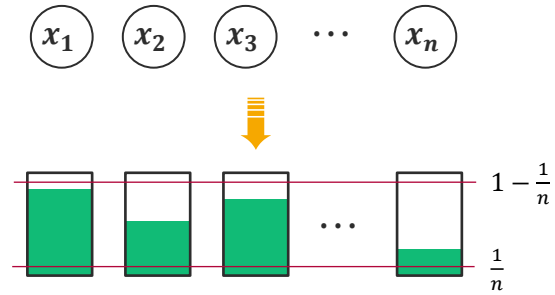
Theory



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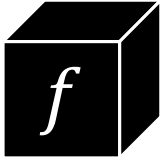
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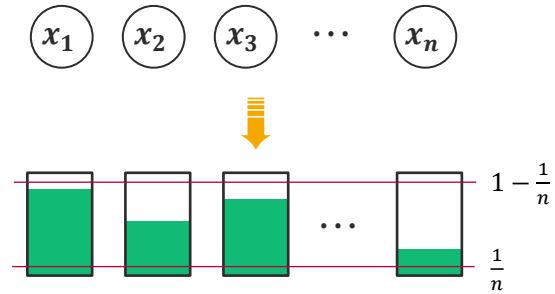
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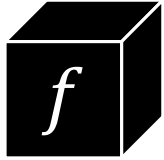
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Multi-valued analyses

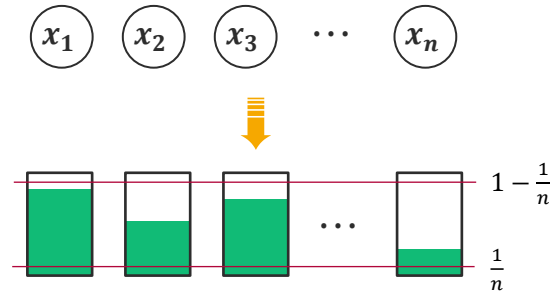
Theory



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(green mass)



Multi-valued analyses

» DOI: [10.1016/j.tcs.2024.114622](https://doi.org/10.1016/j.tcs.2024.114622)



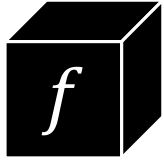
[Ben Jedidia,
Doerr, K.'24]

» *Runtime Analysis of a Multi-Valued Compact Genetic Algorithm on Generalized OneMax*



[Adak, Witt'24]

Theory

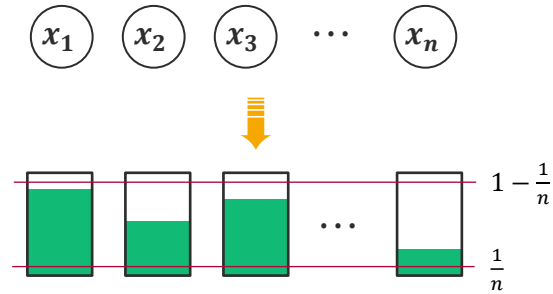


$$f: \{0,1\}^n \rightarrow \mathbf{R}$$

- » Pseudo-Boolean optimization
 - ❖ **Global optimum** often 1^n



- » **Run time** analysis
 - ❖ Number of evaluations of f until a global optimum is sampled



- » *Frequency vector* \mathbf{p}
- » \mathbf{p}_i ... probability to sample a 1 at i (green mass)



Multi-valued analyses

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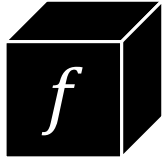


[Ben Jedidia, Doerr, K.'24]



[Adak, Witt'24]

Theory



$$f: \{0,1\}^n \rightarrow \mathbf{R}$$

» Pseudo-Boolean optimization

❖ **Global optimum** often 1^n

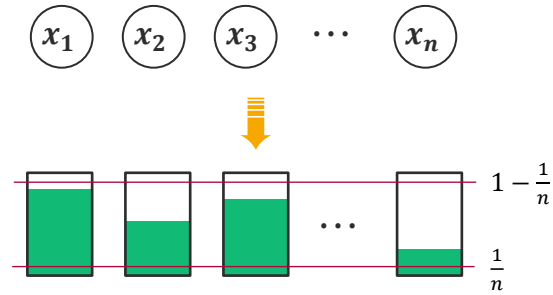


» **Run time** analysis

❖ Number of evaluations of f until a global optimum is sampled



» Favorable albeit not necessary model



» *Frequency vector* \mathbf{p}

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Multi-valued analyses

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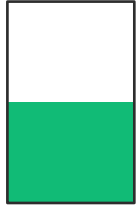


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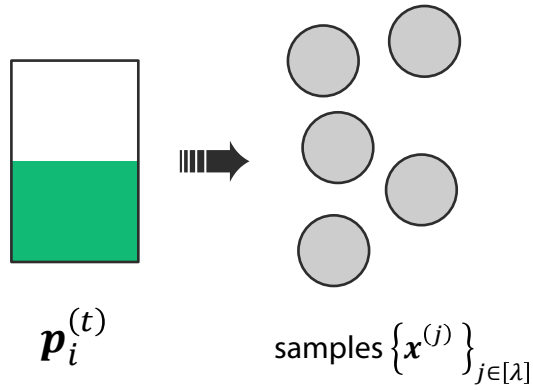
[Adak, Witt'24]

Updating a Frequency (commonly)

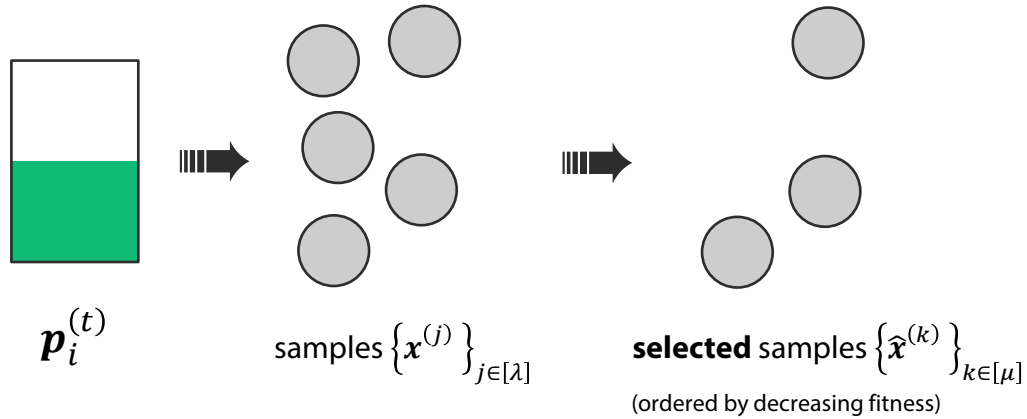


$$p_i^{(t)}$$

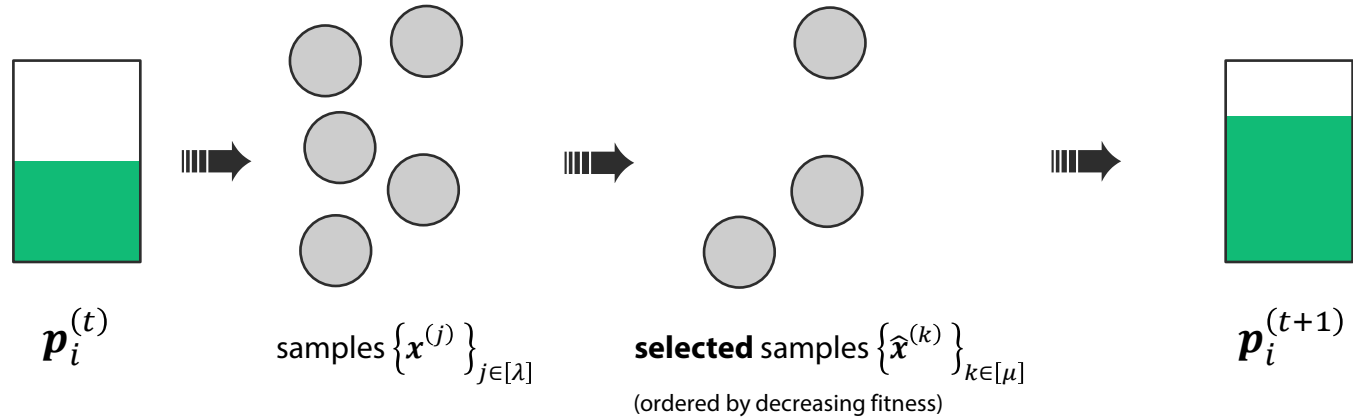
Updating a Frequency (commonly)



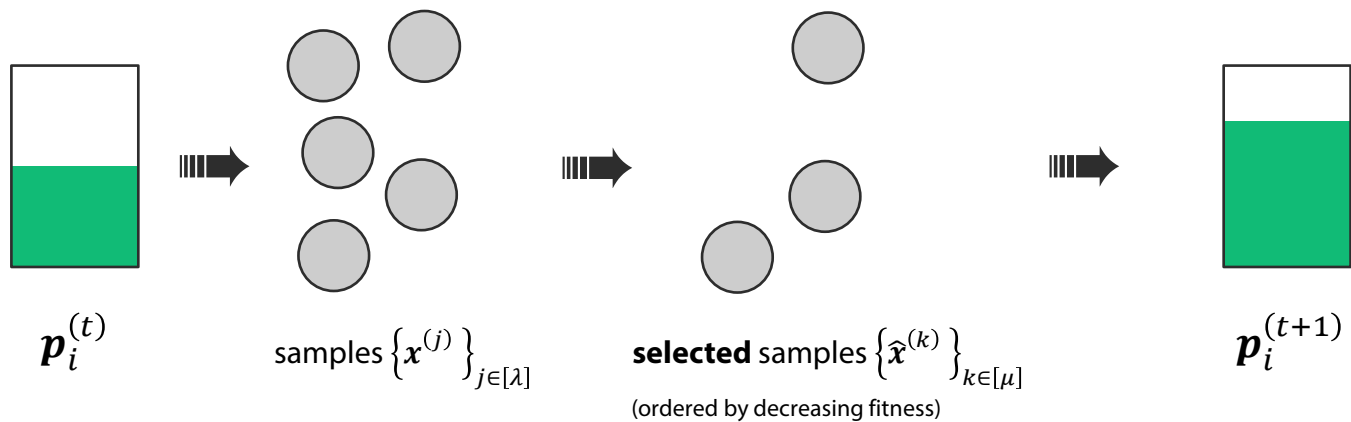
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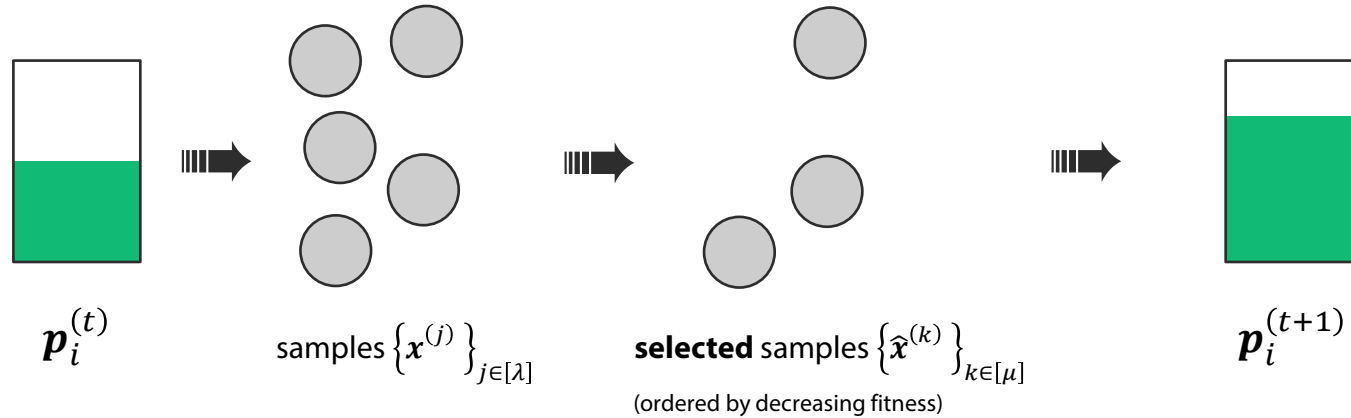
Updating a Frequency (commonly)



$$\mathbb{E} \left[p_i^{(t+1)} - p_i^{(t)} \mid p^{(t)} \right]$$

drift

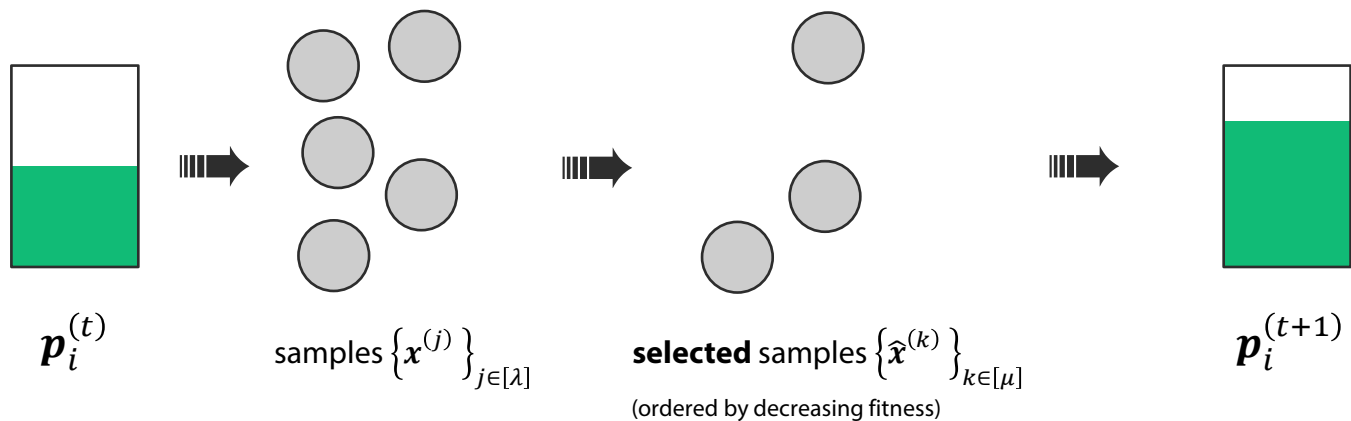
Updating a Frequency (commonly)



$E \left[p_i^{(t+1)} - p_i^{(t)} \mid p^{(t)} \right]$ » Estimating the drift, estimates the expected time that the frequency passes a specific value

drift

Updating a Frequency (commonly)



$$\mathbb{E} \left[p_i^{(t+1)} - p_i^{(t)} \mid p^{(t)} \right]$$

drift

» Estimating the drift, estimates the expected time that the frequency passes a specific value

❖ **Drift analysis**



1712.00964



[Lengler'18]

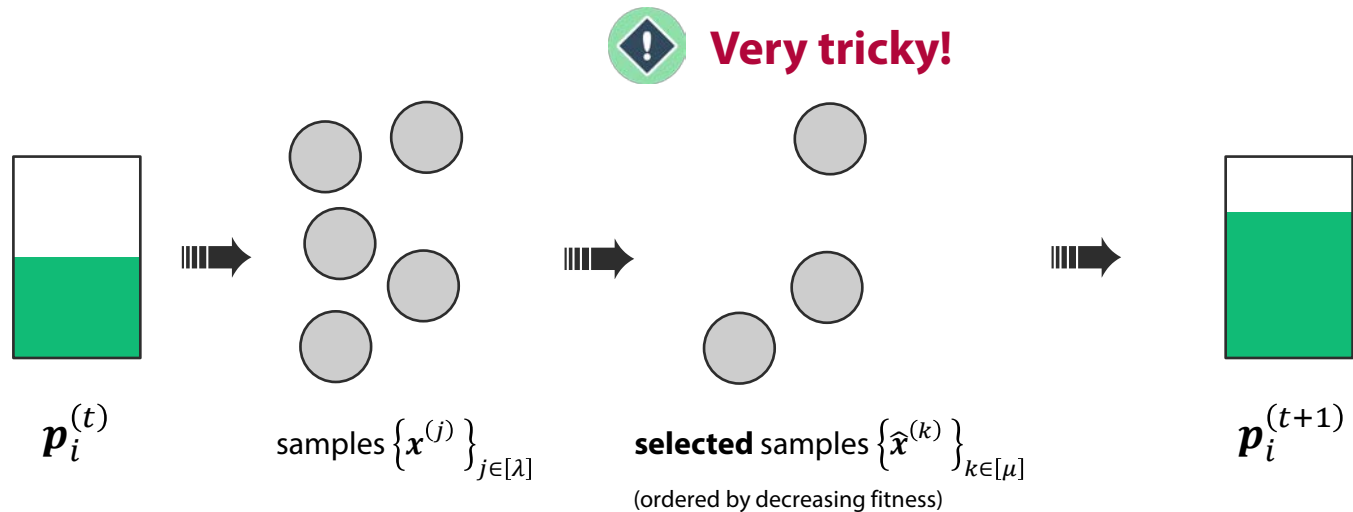


2406.14589



[Kötzing'24]

Updating a Frequency (commonly)



$$\mathbb{E} \left[p_i^{(t+1)} - p_i^{(t)} \mid p^{(t)} \right]$$

drift

» Estimating the drift, estimates the expected time that the frequency passes a specific value

❖ **Drift analysis**



1712.00964



[Lengler'18]



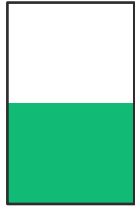
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[Kötzing'24]

Updating a Frequency (Genetic drift)

» **What** happens if the samples are **not** informative?



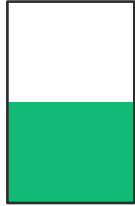
$p_i^{(t)}$

Updating a Frequency (Genetic drift)

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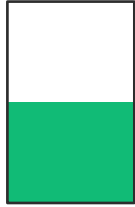
❖ Each $\{\mathbf{x}_i^{(j)}\}_{j \in [\lambda]}$ and $\{\hat{\mathbf{x}}_i^{(k)}\}_{k \in [\mu]}$ follows the same distribution

» Can happen regularly during a run



$p_i^{(t)}$

Updating a Frequency (Genetic drift)



$$p_i^{(t)}$$

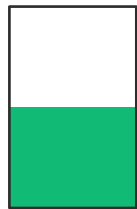
» **What** happens if the samples are **not** informative?

❖ Each $\{x_i^{(j)}\}_{j \in [\lambda]}$ and $\{\hat{x}_i^{(k)}\}_{k \in [\mu]}$ follows the same distribution

➡ **Removes** the tricky situation

» Can happen regularly during a run

Updating a Frequency (Genetic drift)



$p_i^{(t)}$

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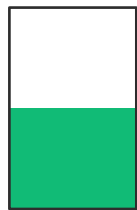
➡ **Removes** the tricky situation

$$E \left[p_i^{(t+1)} \mid p^{(t)} \right] = p_i^{(t)}$$

balanced

» Typical property of univariate EDAs

Updating a Frequency (Genetic drift)



$p_i^{(t)}$

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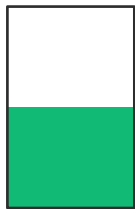
» Typical property of univariate EDAs

balanced

➡ p_i is a **martingale** and **approaches the borders quickly**

» Due to random fluctuations; despite a clear signal

Updating a Frequency (Genetic drift)



$p_i^{(t)}$

» **What** happens if the samples are **not** informative?

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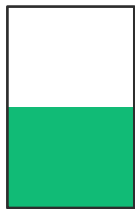
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Updating a Frequency (Genetic drift)



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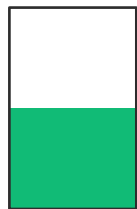
» **How fast** does p_i cover distance $d \in [0,1]$?

Updating a Frequency (Genetic drift)

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❖ Each $\{x_i^{(j)}\}_{j \in [\lambda]}$ and $\{\hat{x}_i^{(k)}\}_{k \in [\mu]}$ follows the same distribution

» Can happen regularly during a run



$p_i^{(t)}$

➡ **Removes** the tricky situation

$$E \left[p_i^{(t+1)} \mid p^{(t)} \right] = p_i^{(t)} \quad \text{» Typical property of univariate EDAs}$$

balanced

➡ p_i is a **martingale** and **approaches the borders quickly**

» Due to random fluctuations; despite a clear signal

❖ **Genetic drift**

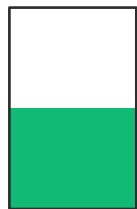


» **How fast** does p_i cover distance $d \in [0,1]$?

drift via
➡
variance

$$\approx \frac{d}{\text{Var} \left[p_i^{(t+1)} \mid p^{(t)} \right]} \quad \text{[K,'19]}$$

Updating a Frequency (Genetic drift)



$p_i^{(t)}$

» **What** happens if the samples are **not** informative?

❖ Each $\{x_i^{(j)}\}_{j \in [\lambda]}$ and $\{\hat{x}_i^{(k)}\}_{k \in [\mu]}$ follows the same distribution

» Can happen regularly during a run

➡ **Removes** the tricky situation

$E[p_i^{(t+1)} | p^{(t)}] = p_i^{(t)}$ » Typical property of univariate EDAs

balanced

➡ p_i is a **martingale** and **approaches the borders quickly**

» Due to random fluctuations; despite a clear signal

❖ **Genetic drift**



» **How fast** does p_i cover distance $d \in [0,1]$?

drift via
➡
variance

$$\approx \frac{d}{\text{Var}[p_i^{(t+1)} | p^{(t)}]} \quad [\text{K}'19]$$

» This hitting time is **concentrated**

[Doerr, Zheng'20]

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration



» Select $\mu = 2$ best samples, ranking them

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]

Commonly Studied EDAs

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[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration



» Select $\mu = 2$ best samples, ranking them



» **Adjust** in favor of victor

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]


Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]

 » $\lambda = 2$ samples each iteration



 » Select $\mu = 2$ best samples, ranking them



 » **Adjust** in favor of victor

$$\mathbf{p}_i^{(t+1)} = \mathbf{p}_i^{(t)} + \frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)})$$

hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



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hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

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Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration



» Select $\mu \leq \lambda$ best samples, ranking them

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration



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hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration



» Select $\mu \leq \lambda$ best samples, ranking them



» **Set** to the relative number of 1s

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration



» Select $\mu = 2$ best samples, ranking them



» **Adjust** in favor of victor

$$\mathbf{p}_i^{(t+1)} = \mathbf{p}_i^{(t)} + \frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)})$$

hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration



» Select $\mu \leq \lambda$ best samples, ranking them



» **Set** to the relative number of 1s

$$\mathbf{p}_i^{(t+1)} = \frac{1}{\mu} \sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)}$$

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



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$$\mathbf{p}_i^{(t+1)} = \mathbf{p}_i^{(t)} + \frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)})$$

hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration



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» **Set** to the relative number of 1s

$$\mathbf{p}_i^{(t+1)} = \frac{1}{\mu} \sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)}$$



» Population-based incremental learning

[Baluja'94]

» 2-Max-Min ant system with iteration-best update

[Neumann, Sudholt, Witt'10]

Commonly Studied EDAs

Compact Genetic Algorithm (cGA)

[Harik, Lobo, Goldberg'99]



» $\lambda = 2$ samples each iteration



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» **Adjust** in favor of victor

$$\mathbf{p}_i^{(t+1)} = \mathbf{p}_i^{(t)} + \frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)})$$

hypothetical population size

Univariate Marginal Distribution Algorithm (UMDA)

[Mühlenbein, Paaß'96]



» λ samples each iteration



» Select $\mu \leq \lambda$ best samples, ranking them



» **Set** to the relative number of 1s

$$\mathbf{p}_i^{(t+1)} = \frac{1}{\mu} \sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)}$$



- » Population-based incremental learning [Baluja'94]
- » 2-Max-Min ant system with iteration-best update [Neumann, Sudholt, Witt'10]

All are **balanced!** (starting with the uniform distribution)

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

Univariate Marginal Distribution Algorithm (UMDA)

$$\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] = \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right]$$

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

Univariate Marginal Distribution Algorithm (UMDA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right]\end{aligned}$$

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

Univariate Marginal Distribution Algorithm (UMDA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \text{Var} \left[\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right]\end{aligned}$$

Commonly Studied EDAs (variances)

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Univariate Marginal Distribution Algorithm (UMDA)

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Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

Univariate Marginal Distribution Algorithm (UMDA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \text{Var} \left[\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \left(\text{Var} \left[\hat{\mathbf{x}}_i^{(1)} \mid \mathbf{p}^{(t)} \right] + \text{Var} \left[\hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \right) \\ &= \frac{2}{K^2} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)})\end{aligned}$$

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \text{Var} \left[\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \left(\text{Var} \left[\hat{\mathbf{x}}_i^{(1)} \mid \mathbf{p}^{(t)} \right] + \text{Var} \left[\hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \right) \\ &= \frac{2}{K^2} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)})\end{aligned}$$

➡ \mathbf{p}_i covers a distance of at most $\frac{1}{4}$ with constant probability within $\Theta(K^2)$ iterations

Univariate Marginal Distribution Algorithm (UMDA)

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} \left(\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \right) \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \text{Var} \left[\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \left(\text{Var} \left[\hat{\mathbf{x}}_i^{(1)} \mid \mathbf{p}^{(t)} \right] + \text{Var} \left[\hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \right) \\ &= \frac{2}{K^2} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)})\end{aligned}$$

➡ \mathbf{p}_i covers a distance of at most $\frac{1}{4}$ with constant probability within $\Theta(K^2)$ iterations

Univariate Marginal Distribution Algorithm (UMDA)

$$\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] = \text{Var} \left[\frac{1}{\mu} \sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)} \mid \mathbf{p}^{(t)} \right]$$

Commonly Studied EDAs (variances)

Compact Genetic Algorithm (cGA)

$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\mathbf{p}_i^{(t)} + \frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)}) \mid \mathbf{p}^{(t)} \right] \\ &= \text{Var} \left[\frac{1}{K} (\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)}) \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \text{Var} \left[\hat{\mathbf{x}}_i^{(1)} - \hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{K^2} \cdot \left(\text{Var} \left[\hat{\mathbf{x}}_i^{(1)} \mid \mathbf{p}^{(t)} \right] + \text{Var} \left[\hat{\mathbf{x}}_i^{(2)} \mid \mathbf{p}^{(t)} \right] \right) \\ &= \frac{2}{K^2} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)})\end{aligned}$$

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$$\begin{aligned}\text{Var} \left[\mathbf{p}_i^{(t+1)} \mid \mathbf{p}^{(t)} \right] &= \text{Var} \left[\frac{1}{\mu} \sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)} \mid \mathbf{p}^{(t)} \right] \\ &= \frac{1}{\mu^2} \text{Var} \left[\sum_{k \in [\mu]} \hat{\mathbf{x}}_i^{(k)} \mid \mathbf{p}^{(t)} \right]\end{aligned}$$

Commonly Studied EDAs (variances)

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➡ \mathbf{p}_i covers a distance of at most $\frac{1}{4}$ with constant probability within $\Theta(\mu)$ iterations

Theoretician's Trick



[Sudholt, Witt'16; Friedrich, Kötzing, K.'16; Doerr, Zheng'20]

» We wish **genetic drift** to be low for at least $\Theta(T)$ iterations

Theoretician's Trick



[Sudholt, Witt'16; Friedrich, Kötzing, K.'16; Doerr, Zheng'20]

» We wish **genetic drift** to be low for at least $\Theta(T)$ iterations

» p_i covers a distance of at most $\frac{1}{4}$ with probability at least $1 - \exp\left(-\Theta\left(\frac{1}{\text{Var}\cdot T}\right)\right)$ [Doerr, Zheng'20]

Theoretician's Trick



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» **Choose** algorithm parameters such that $\text{Var} \in \Theta\left(\frac{1}{T \log n}\right)$ » Guarantee with high probability

Theoretician's Trick



[Sudholt, Witt'16; Friedrich, Kötzing, K.'16; Doerr, Zheng'20]

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Theoretician's Trick



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» **Smart-restart strategy** for EDAs [Doerr, Zheng'20]

❖ Choose some variance and run the EDA for $\Theta\left(\frac{1}{\text{Var}}\right)$ iterations

Theoretician's Trick



[Sudholt, Witt'16; Friedrich, Kötzing, K.'16; Doerr, Zheng'20]

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» **Smart-restart strategy** for EDAs [Doerr, Zheng'20]

❖ Choose some variance and run the EDA for $\Theta\left(\frac{1}{\text{Var}}\right)$ iterations

❖ If not happy, **halve the variance**, repeat the algorithm

Theoretician's Trick



[Sudholt, Witt'16; Friedrich, Kötzing, K.'16; Doerr, Zheng'20]

» We wish **genetic drift** to be low for at least $\Theta(T)$ iterations

» p_i covers a distance of at most $\frac{1}{4}$ with probability at least $1 - \exp\left(-\Theta\left(\frac{1}{\text{Var}\cdot T}\right)\right)$ [Doerr, Zheng'20]

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» Several times derived before, not fully rigorously [Thierens, Goldberg, Pereira'98; Lobo, Goldberg, Pelikan'00; Shapiro'05]

» **Smart-restart strategy** for EDAs [Doerr, Zheng'20]

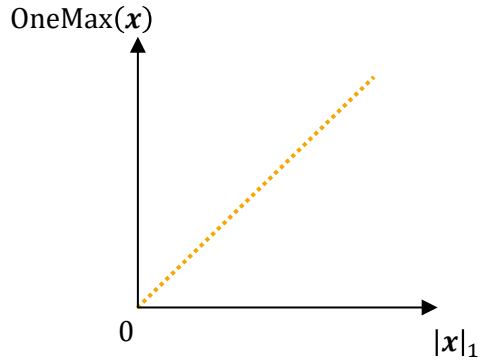
❖ Choose some variance and run the EDA for $\Theta\left(\frac{1}{\text{Var}}\right)$ iterations

❖ If not happy, **halve the variance**, repeat the algorithm

» The trick carries over to weak preferences of bit values [Doerr, Zheng'20]

Commonly Studied Benchmarks

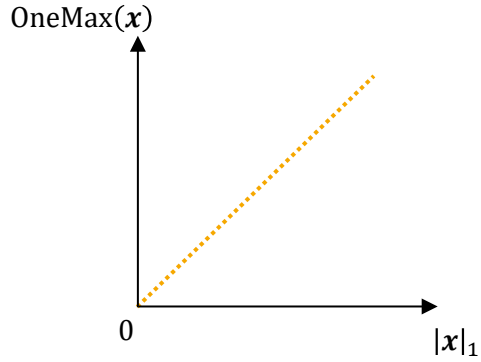
OneMax



$$\mathbf{x} \mapsto \sum_{i \in [n]} x_i =: |\mathbf{x}|_1$$

Commonly Studied Benchmarks

OneMax



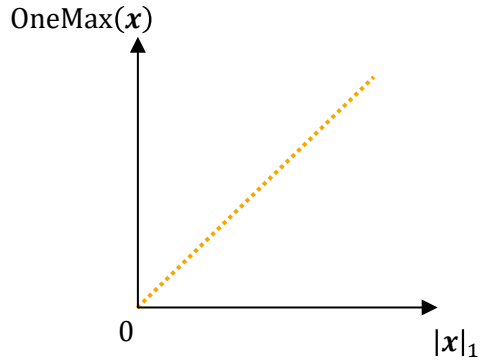
LeadingOnes

110000011010
111110000100

$$x \mapsto \sum_{i \in [n]} x_i =: |x|_1 \quad x \mapsto \max\{i \in [0..n] \mid \forall j \in [i]: x_j = 1\}$$

Commonly Studied Benchmarks

OneMax



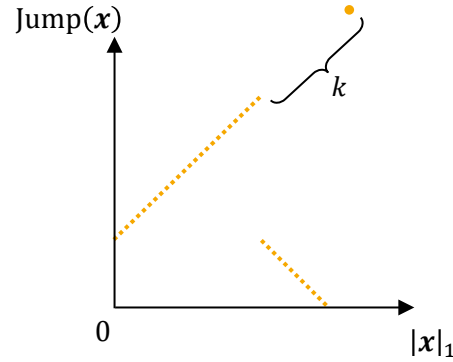
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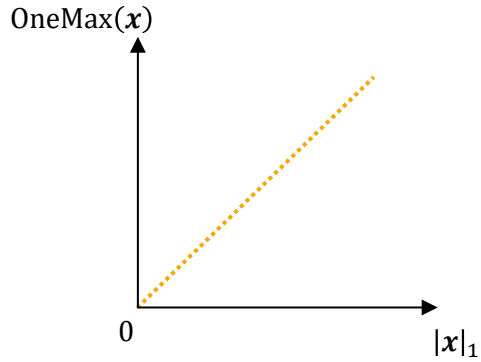
Jump



$$x \mapsto \begin{cases} k + |x|_1 & \text{if } |x|_1 \in [n - k] \cup \{n\} \\ n - |x|_1 & \text{else} \end{cases}$$

Commonly Studied Benchmarks

OneMax



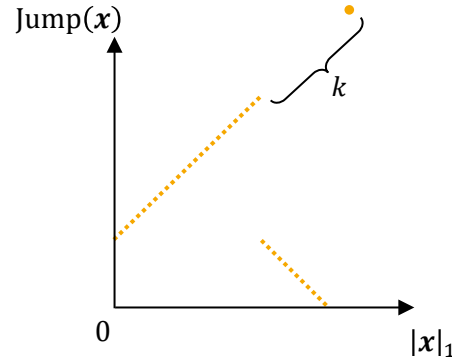
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BinVal









$[110000011010]_2$
 $[111110000100]_2$

$$x \mapsto \sum_{i \in [n]} 2^{n-i} \cdot x_i$$

Commonly Studied Benchmarks

	OneMax	LeadingOnes	Jump	BinVal
cGA	 [Droste'06; Lengler, Sudholt, Witt'18; Sudholt, Witt'19]		 [Hasenöhr, Sutton'18; Doerr'21; Witt'23]	 [Droste'06; Witt'18]
UMDA	 [Dang, Lehre, Nguyen'18; Witt'19; K., Witt'20]	 [Dang, Lehre, Nguyen'18; Doerr, K.'21]		 [Dang, Lehre, Nguyen'18]

Commonly Studied Benchmarks

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Also: (non-exhaustive)

- » **Analyses on noise** [Friedrich, Kötzing, K., Witt'17; Lehre, Nguyen'19; Lehre, Nguyen'21; Kötzing, Radhakrishnan'22]
- » **Analyses on deception** [Lehre, Nguyen'19; Doerr, K.'21]
- » **New EDAs** [Doerr, K.'20; Ajimakin, Devi'23]
- » **Multi-valued EDAs** [Ben Jedidia, Doerr, K.'24; Adak, Witt'24]

cGA on OneMax

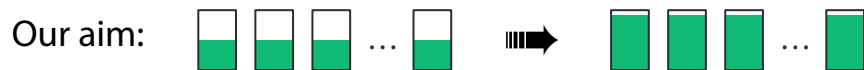


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➡ Since $\underbrace{\left| |\dot{x}^{(1)}|_1 - |\dot{x}^{(2)}|_1 \right|}_{=: D} \geq 2$, position i is not informative

cGA on OneMax (the sampling variance)

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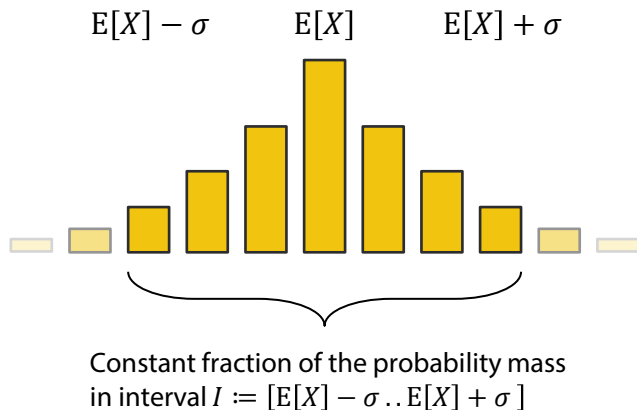
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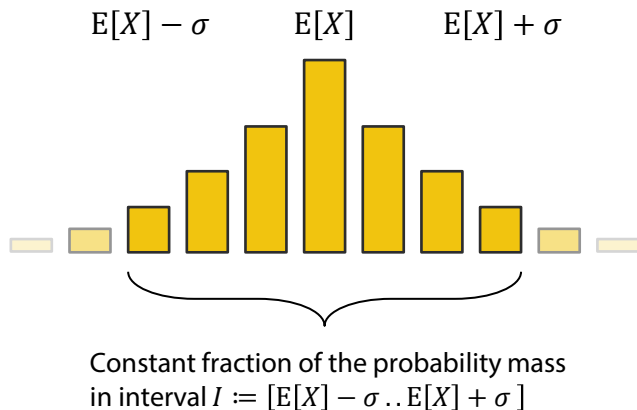
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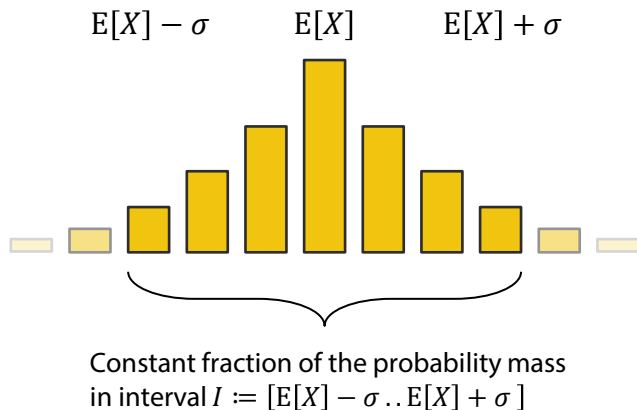
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$$\sqrt{\sum_{j \in [n] \setminus \{i\}} \mathbf{p}_j^{(t)} (1 - \mathbf{p}_j^{(t)})}$$

sampling variance

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» Let $\sigma_i = \sqrt{\sum_{j \in [n] \setminus \{i\}} p_j^{(t)} (1 - p_j^{(t)})}$ and $\sigma = \sqrt{\sum_{i \in [n]} p_i^{(t)} (1 - p_i^{(t)})}$

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» The secret property of φ !
For one-sided bounded frequencies, sampling variance and distance to the optimal model are roughly the same!

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cGA on OneMax

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» We have $\mathbb{E}[\mathbf{p}_i^{(t+1)} - \mathbf{p}_i^{(t)} \mid \mathbf{p}^{(t)}] \gtrsim \frac{2}{K\sigma_i} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)}) \geq \frac{2}{K\sigma} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)})$

$$\begin{aligned} \Rightarrow \mathbb{E}[\varphi^{(t)} - \varphi^{(t+1)} \mid \mathbf{p}^{(t)}] &= \sum_{i \in [n]} \mathbb{E}[\mathbf{p}_i^{(t+1)} - \mathbf{p}_i^{(t)} \mid \mathbf{p}^{(t)}] \\ &\gtrsim \frac{2}{K\sigma} \sum_{i \in [n]} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)}) \\ &= \frac{2}{K} \sigma \end{aligned}$$

» Note that since $\mathbf{p}_i^{(t)} \geq \frac{1}{4}$, we have $\sigma^2 = \sum_{i \in [n]} \mathbf{p}_i^{(t)} (1 - \mathbf{p}_i^{(t)}) \geq \frac{1}{4} \varphi^{(t)}$

» The secret property of φ !
For one-sided bounded frequencies, sampling variance and distance to the optimal model are roughly the same!

$$\Rightarrow \mathbb{E}[\varphi^{(t)} - \varphi^{(t+1)} \mid \mathbf{p}^{(t)}] \gtrsim \frac{1}{K} \sqrt{\varphi^{(t)}} \quad \begin{array}{l} \text{variable} \\ \Rightarrow \\ \text{drift} \end{array} \quad \mathbb{E}[T] \lesssim K\sqrt{n}$$

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- » For $K \in O\left(\frac{\sqrt{n}}{\log(n) \log \log n}\right)$ the run time is in $\Omega(K^{1/3}n + n \log n)$ [Lengler, Sudholt, Witt'18] » Covers **high** genetic drift

UMDA on OneMax (with borders)

[Witt'18] 1. The UMDA with $\mu \in \Omega(\sqrt{n} \log n)$ and $\lambda = (1 + \Theta(1))\mu$ optimizes OneMax in $O(\lambda\sqrt{n})$ function evaluations in expectation. This is $O(n \log n)$ for minimal λ .

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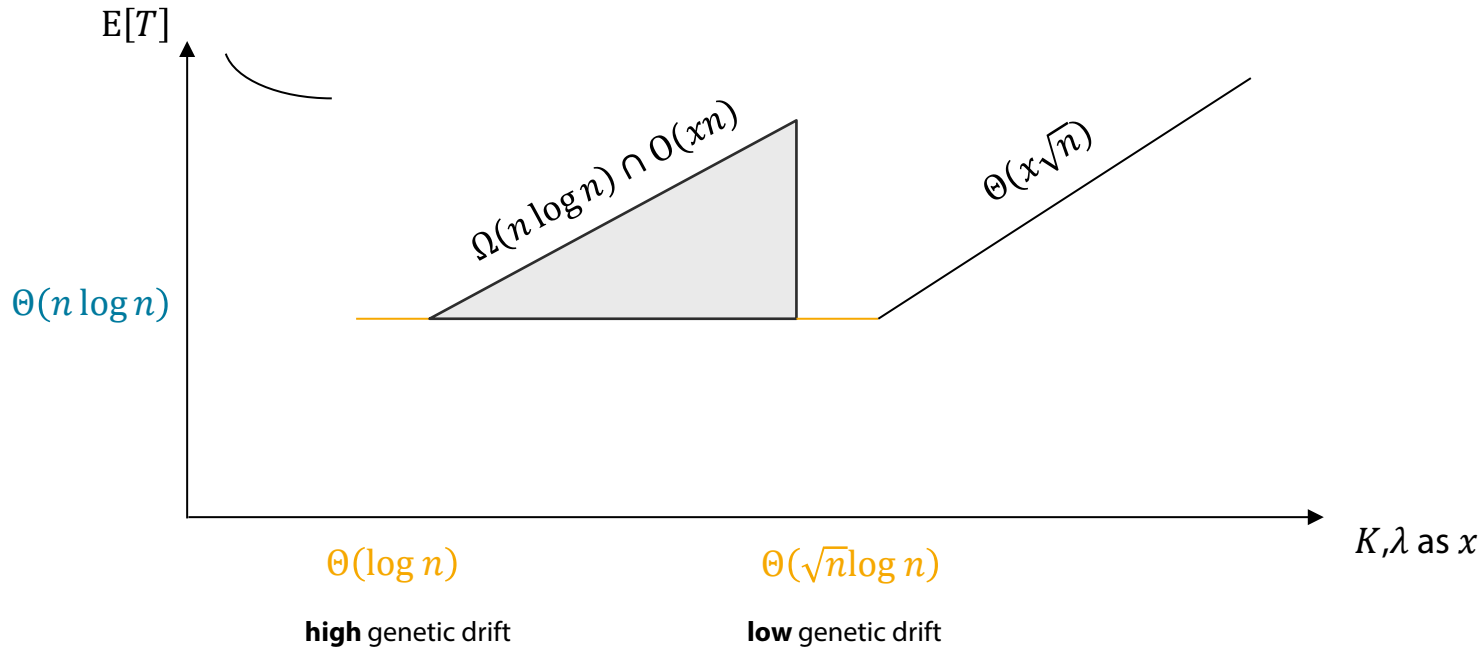
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- » Analysis by Lehre and Nguyen simpler, utilizing a **level-based method**

cGA and UMDA on OneMax



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