

# Genetic Algorithms Parameter Settings

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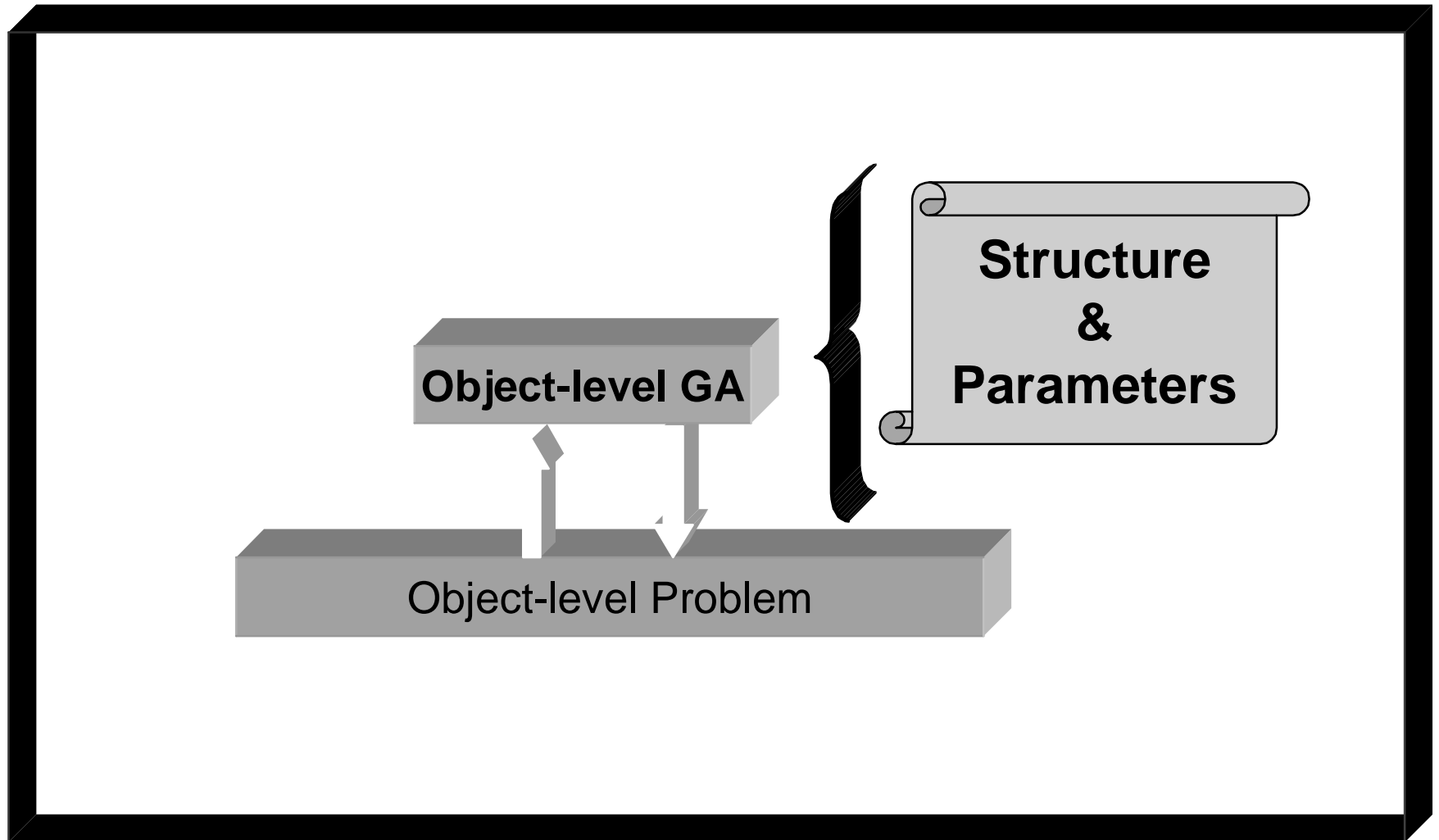
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# GA Parameter Setting - Outline

- **GA Model:** Structure, Parameters
- **GA Parameter Setting**
  - GA Parameter Tuning: Off-line study, Off-line Meta-GA
  - GA Parameter Control: Deterministic Control, Self-Adaptation, Control (FLC)
- **GA Parameter Setting: An Application to Agile Mfg.**
  - Object-level Representation and Complexity
- **First Solution Architecture (U-TGA, TGA, FLC-GA)**
  - FLC-GA: KB, I/O, Control Parameters, Rule base and Scaling Factors
  - Results - Performance Tradeoff
  - Problems with First Solution
  - Summary and Analysis of 30 Experiments
  - Addressing Problems with First Solution
- **Second Solution**
  - FLC: KB, Inputs, Outputs, Termsets, Rule Sets, Control Parameters
  - Statistical Experiments: Set-Up & Measures
  - Summary and Analysis of 1200 Experiments
- **Conclusions and Next Steps**

# GA Model



# GA Structure

## GA Structural Design Selections:

- GA Type:
  - {Simple, Steady-State, Niche,...}
- Chromosome Encoding:
  - {Binary, Integer, Real,...}
- Constraints Representation:
  - {Penalty function, data structure, filters, ...}
- Fitness Function:
  - {Scalar function (weighted aggregation of multiple functions) ...}
- Selection Method:
  - {Proportional Roulette, Tournament, Rank, Uniform, ... }
- Crossover Operator:
  - {Once-cut, Two-cuts, Uniform, BLX, Parent Weighted, ...}
- Mutation Operator:
  - Mutation Rate: {Exponentially Decreasing, Uniform, ..}
  - Value: {Exp. Decreasing, Uniform, Normally Distr., ...}

# GA Parameters

## Adjustable parameters for a GA

- **N** = Population size - large pop. prevent premature convergence
- **P<sub>c</sub>** = Crossover rate:  $P_{cr} * N = \#$  crossovers per generation
- **P<sub>m</sub>** = Mutation rate:  $P_m * N * L = \#$  mutations per generation
- **G** = Generation Gap - percentage of population to be replaced
- **W** = Scaling Window Size = [1, 7]
- **S** = Selection Strategy = {Elitist, Non-Elitist}

## Other possible parameters (that could be adjusted):

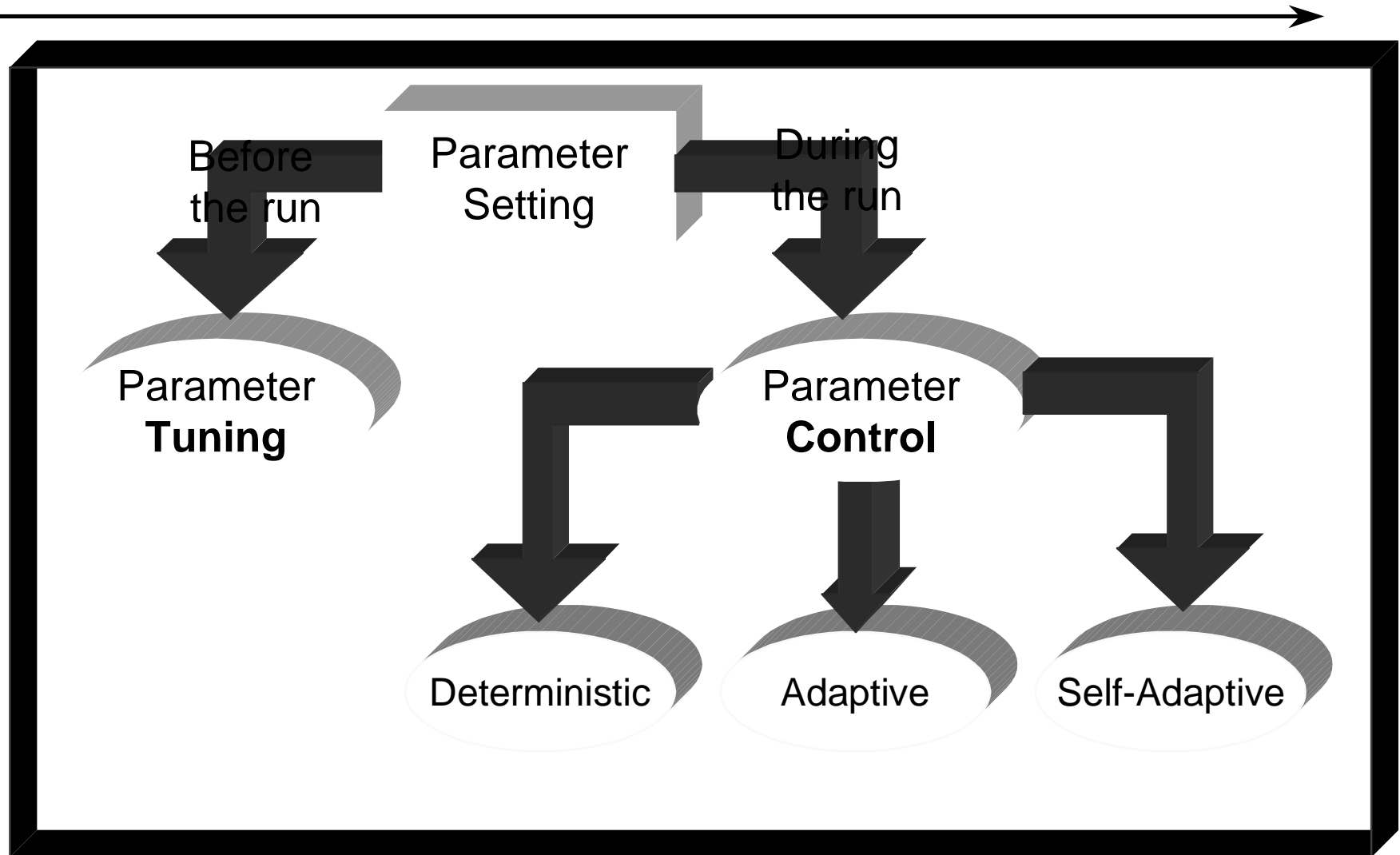
- **T** = Number of Trials =  $\sum N_i$  - where  $i = 1, \text{Max\_Gen}$
- **S<sub>m</sub>** = Mutation step - ( $\sigma$  in Normally distrib. Mutation value)
- **P<sub>S</sub>** = Probability of Selection - (Parametrized slope of prob distrib.)
- **A<sub>S</sub>** = Arity of Parents - number of parents in recombination

# GA Parameter Setting - Outline

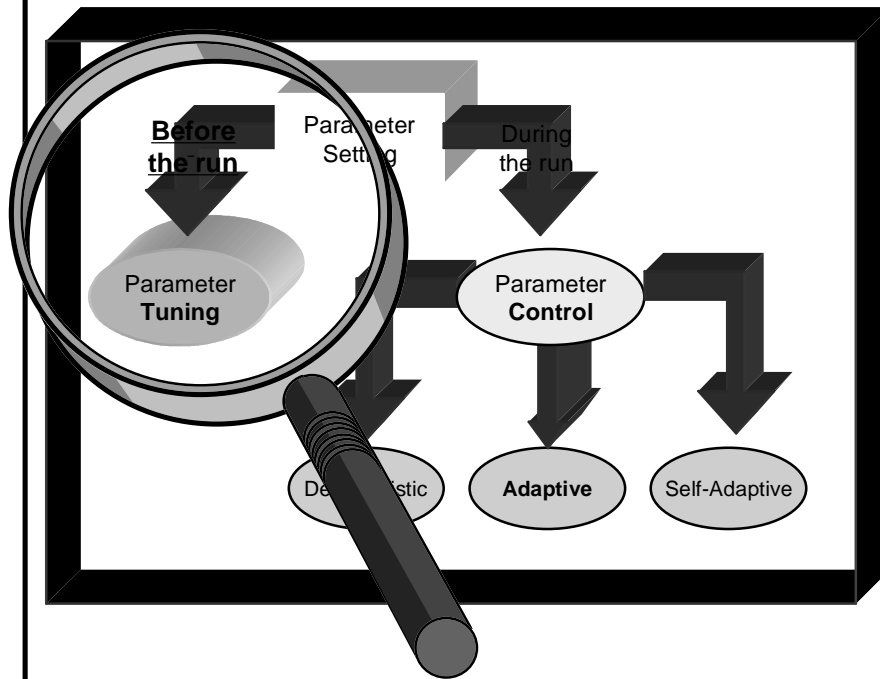
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# GAs Parameter Setting

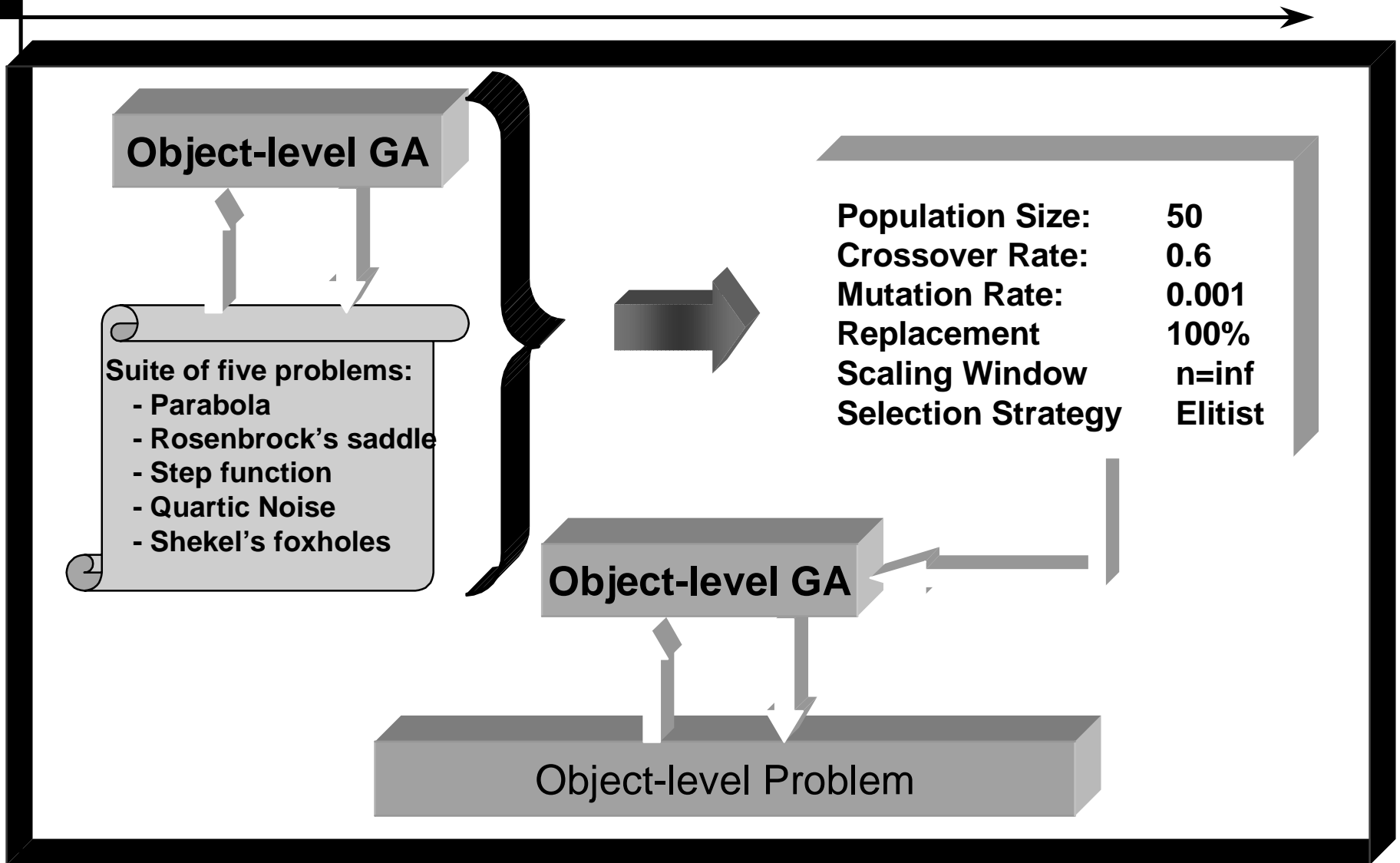


# GAs Parameter Setting: Parameter Tuning

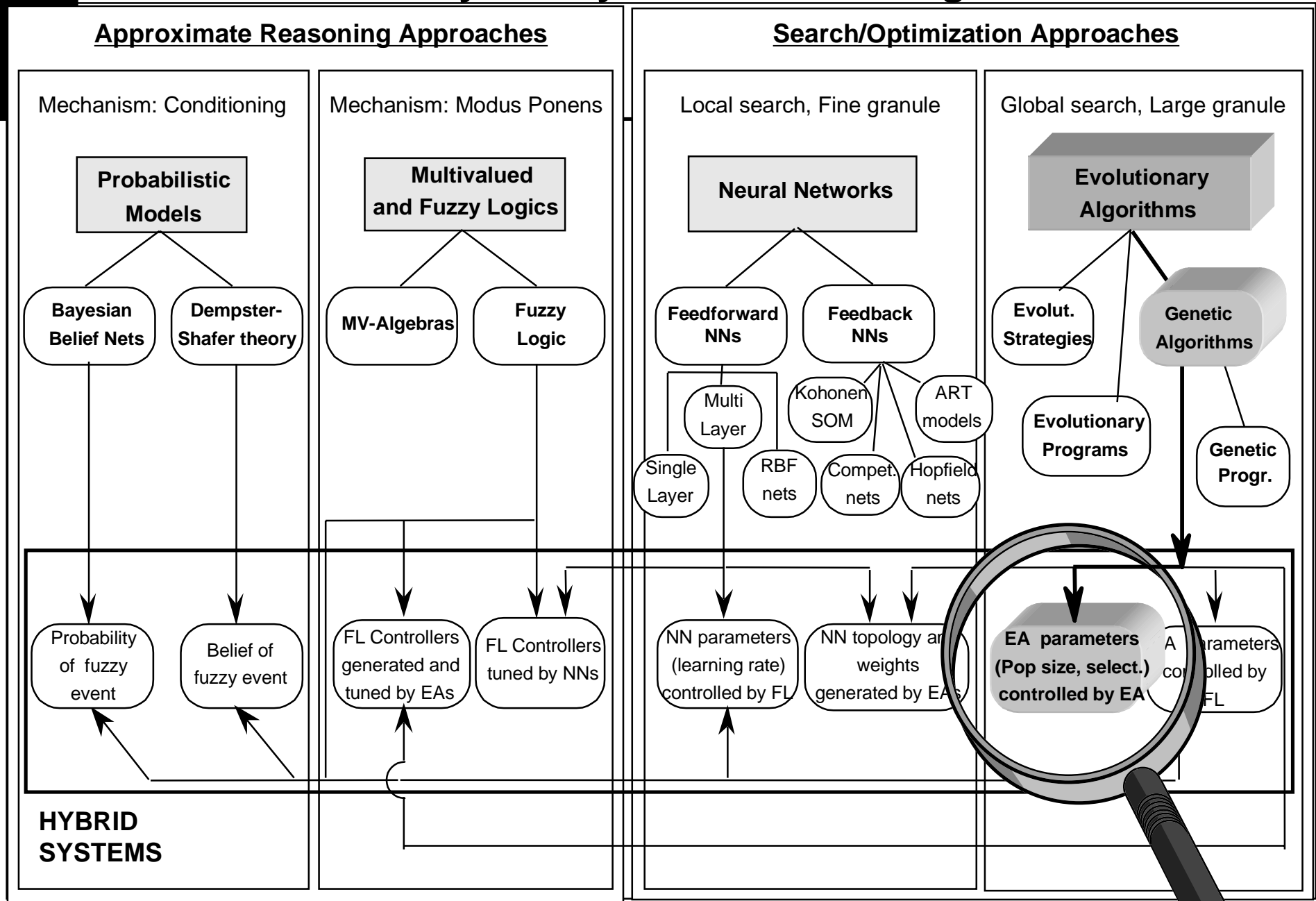


- Off-line Tuning
- Determined before running the GAs on the object-level problem by
  - Studying a subset of five diverse problems (*DeJong, 1975*)
  - Running a Meta-Genetic Algorithm (*Grefenstette, 1986*)

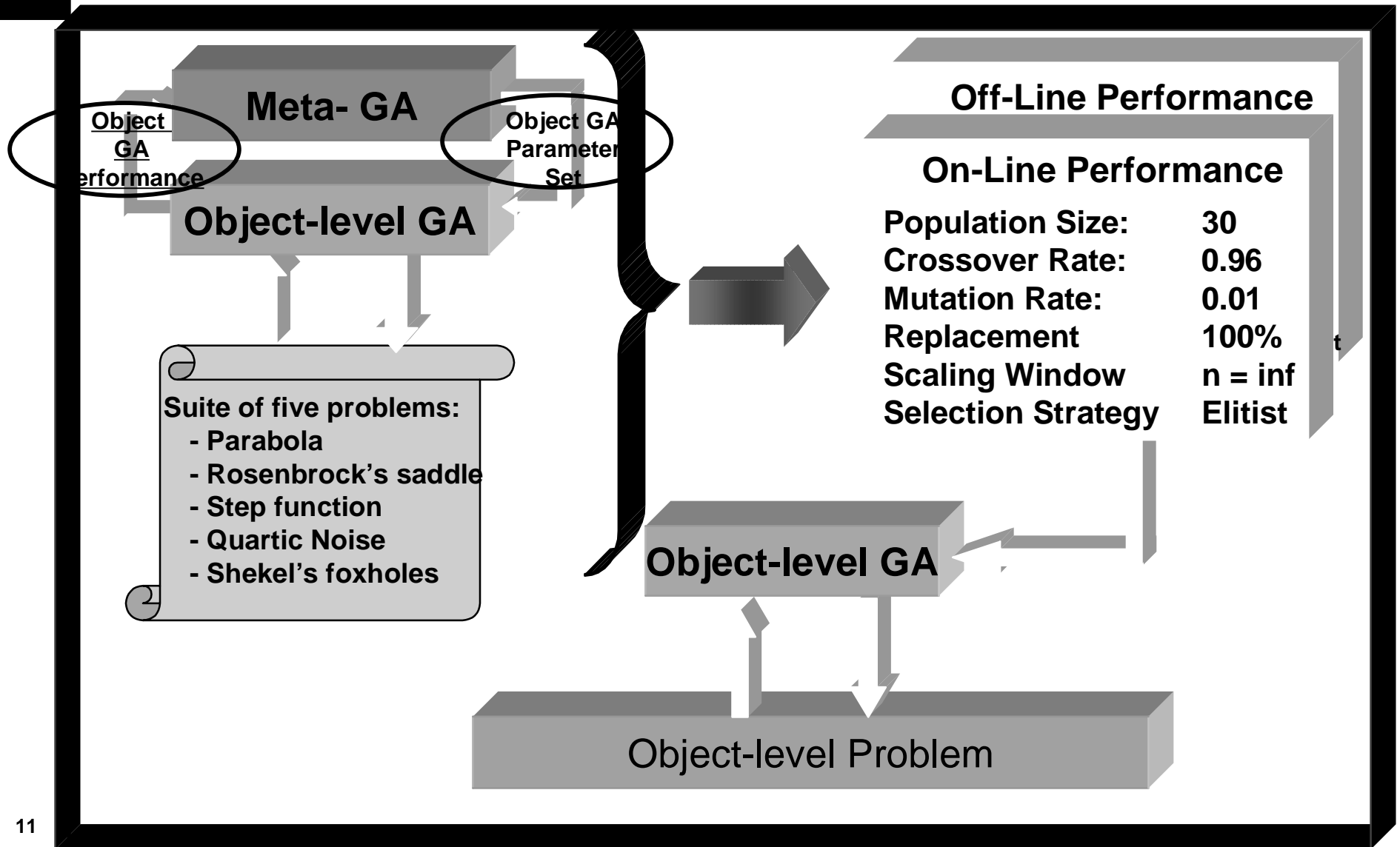
## Off Line Tuning of GA Parameters (*DeJong, 1975*)



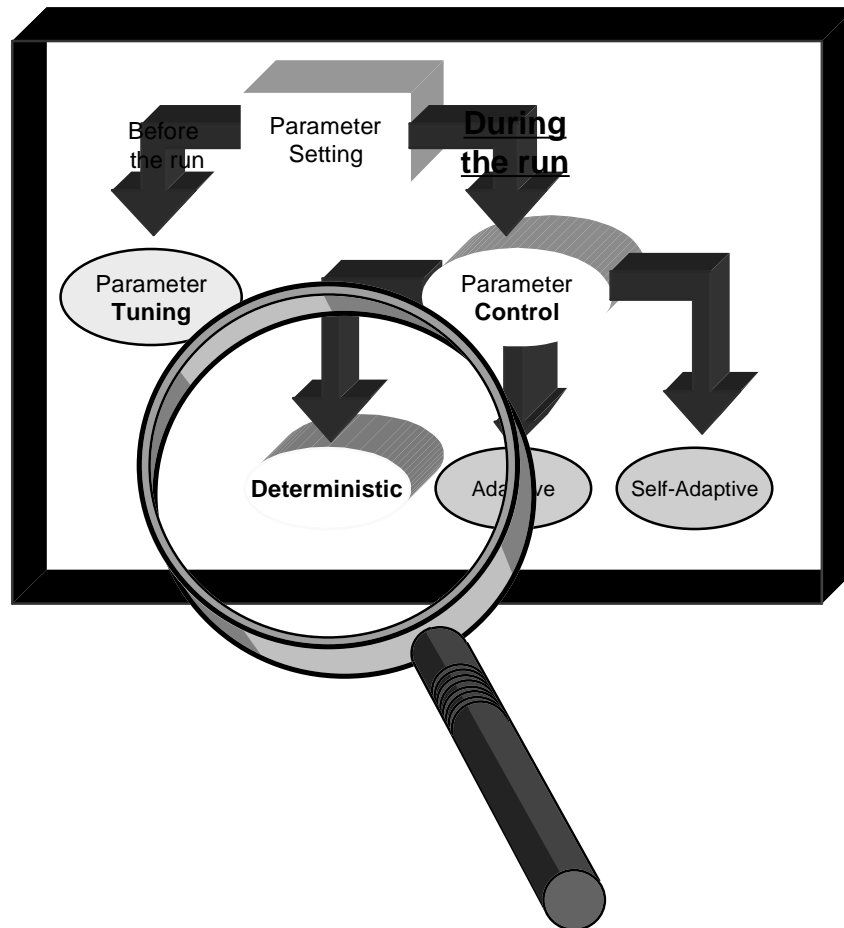
Soft Computing: Evolutionary Comp.: GA parameters control  
**SC Hybrid Systems: EA Tuning EA**



# Off Line Tuning of GA Parameters (*Grefenstette, 1986*)



# GAs Parameter Setting: Deterministic Control

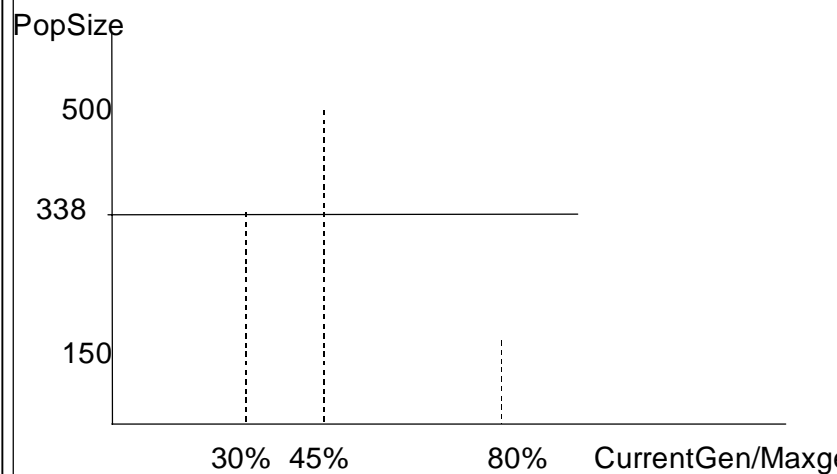


- No feedback information is used.
- A time-varying schedule is used to modify a GA parameter  $p$
- $p$  is replaced by  $p(t)$
- Correct design of  $p(t)$  is very difficult

## GAs Parameter Setting: Deterministic Control - Example

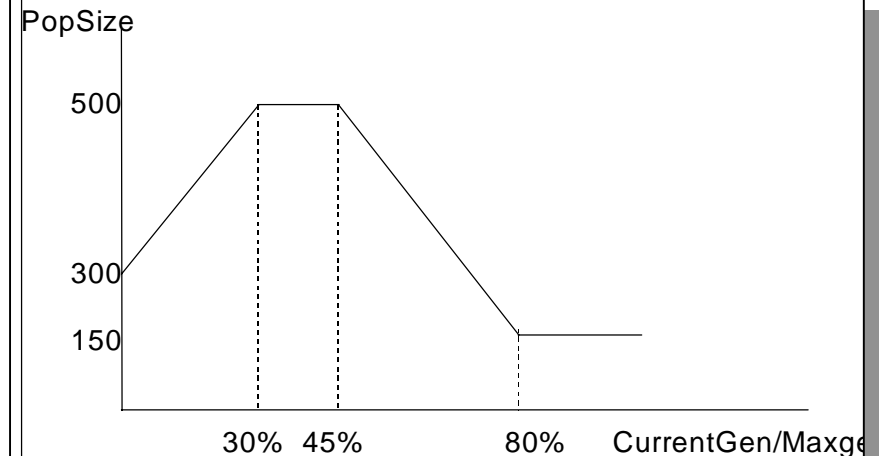
- **Control of Population size**

- By decreasing Population Size toward the last part of the Evolution we are trying to improve the solution refinement (e.g., more generations with same number of trials)

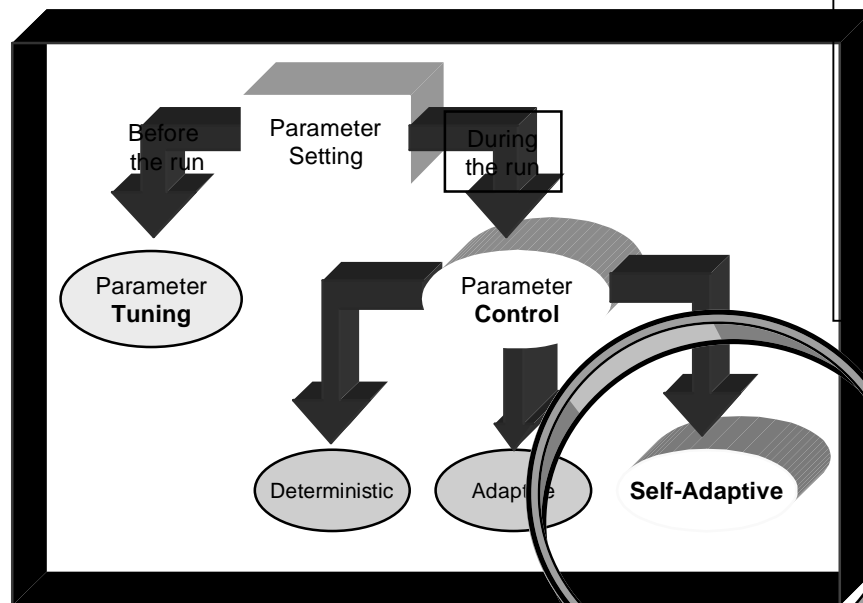


Constant Population size:  $N = 338$   
Number of trials =  $338 * \text{MaxGen}$

Variable Population size:  $N(t)$   
Number of trials =  $338 * \text{MaxGen}$



# GAs Parameter Setting: Self-Adaptive Control



- Incorporate parameters into chromosome making them subject to evolution
- Typically used by ES to determine Mutation Step S:

Mutation Step for Entire Genome

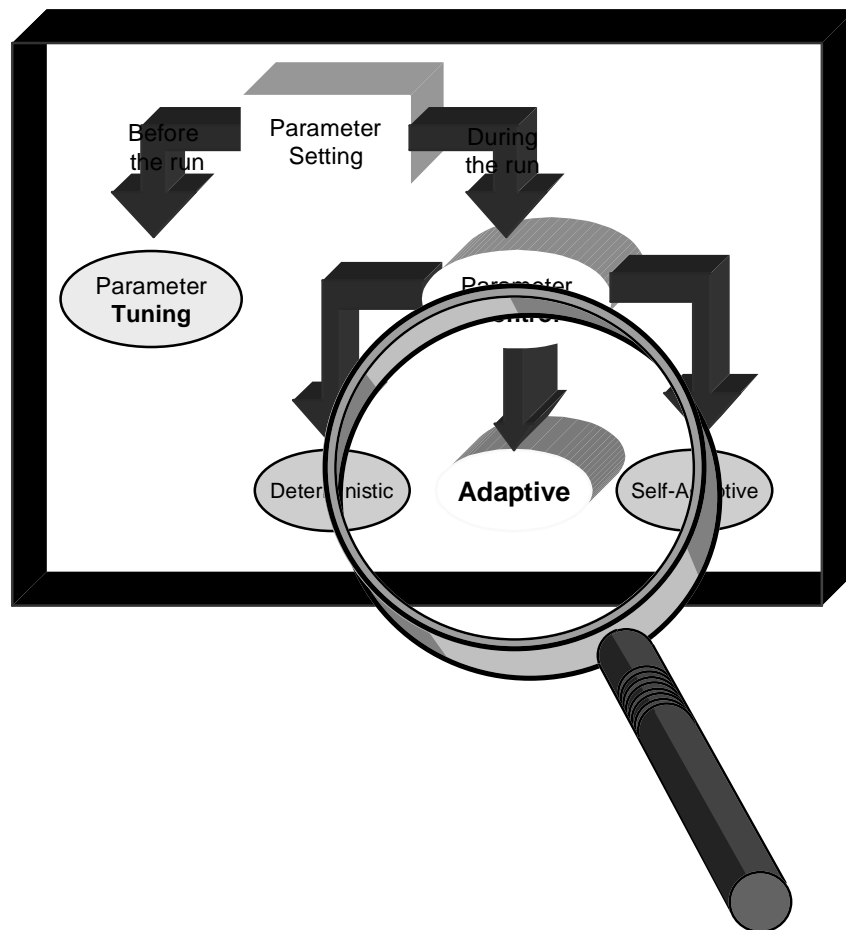
$[g_1 g_2 \dots g_n S]$

or

$[g_1 g_2 \dots g_n S_1 S_2 \dots S_n]$

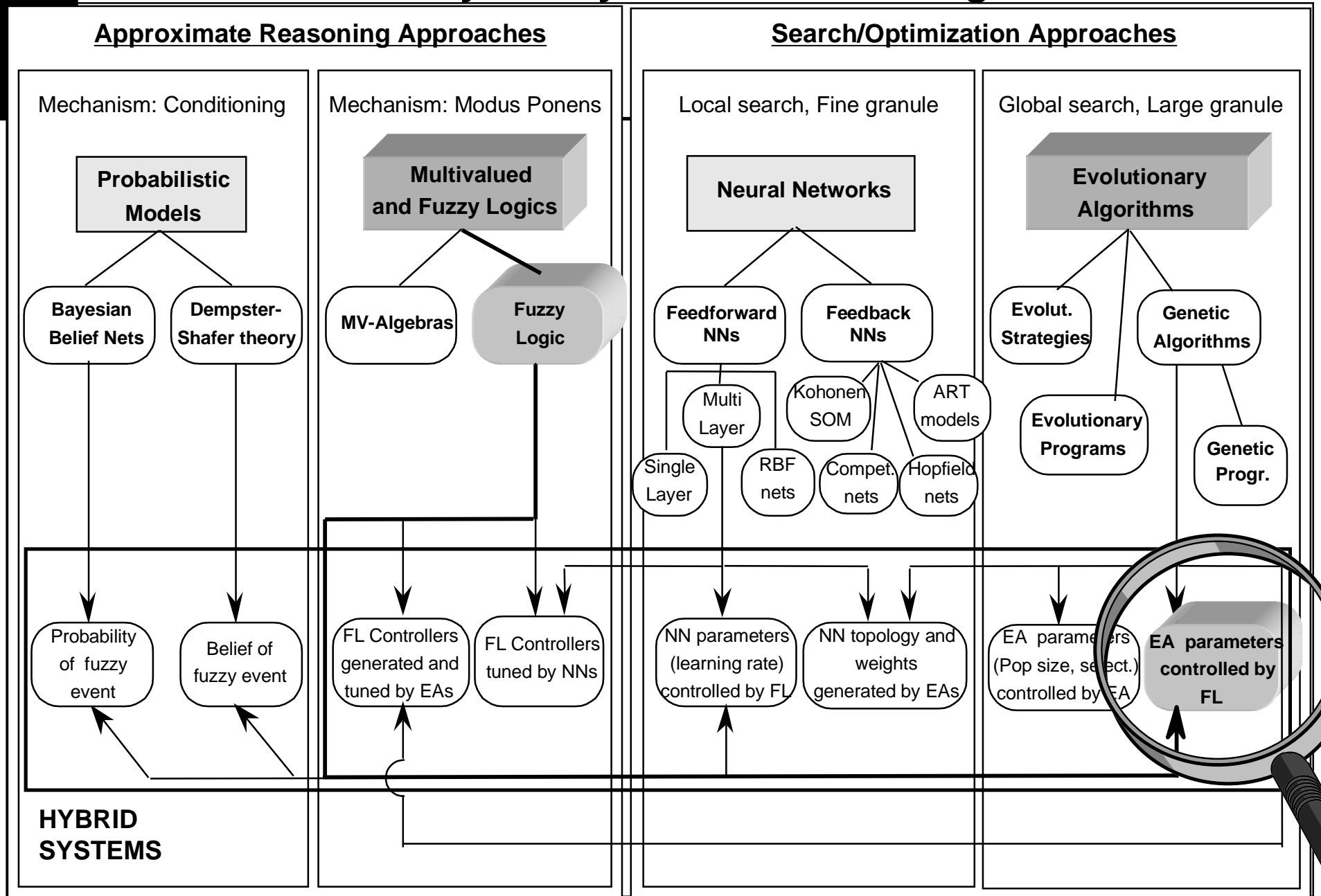
Mutation Steps for Each Genome Value

# GAs Parameter Setting: Adaptive Control

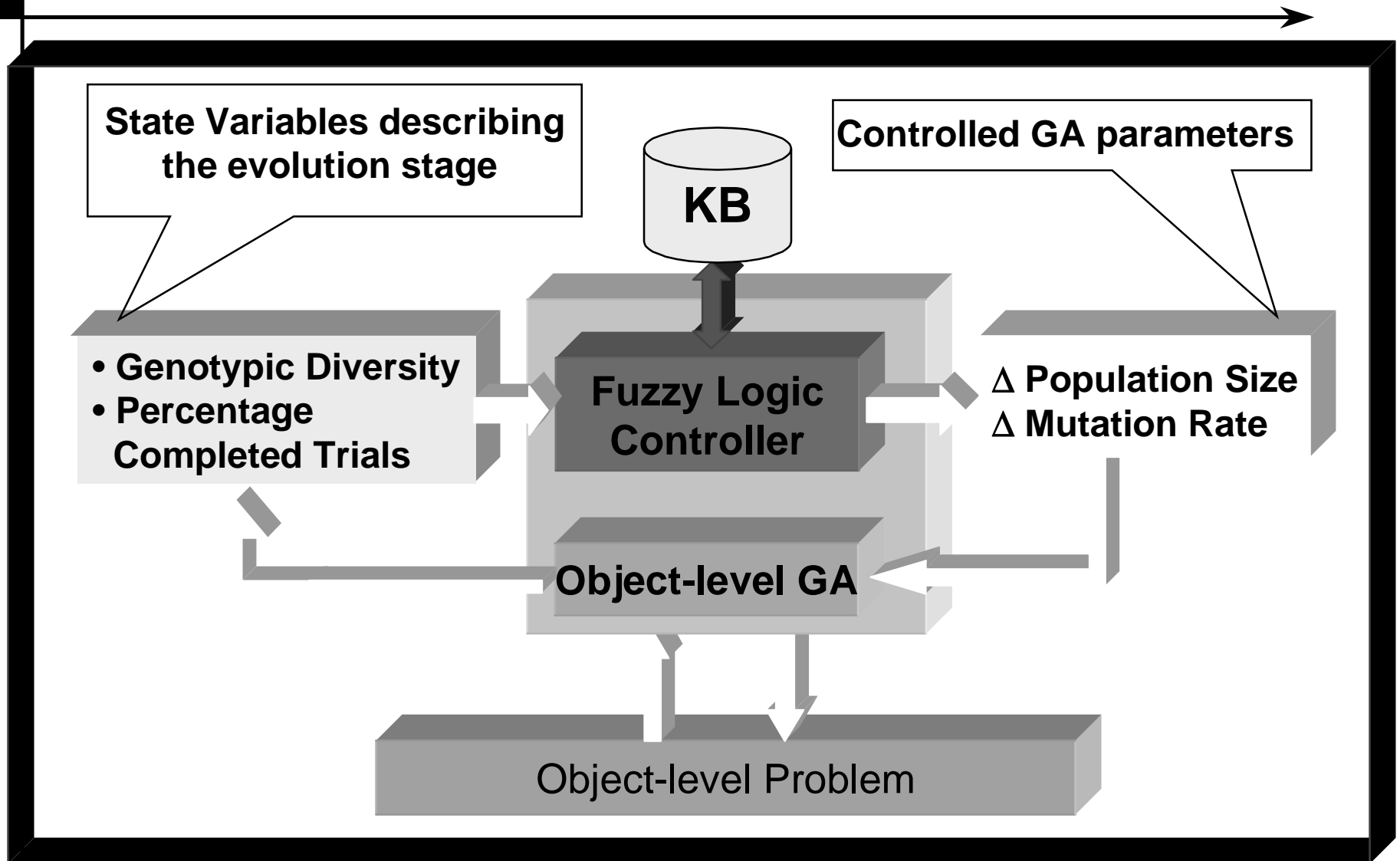


- **Feedback** from the search is used to determine the direction and/or magnitude of the change in the parameter value.
- **A Fuzzy Logic Controller** is used to obtain parameter changes in
  - Population Size
  - Mutation Rate**as a function of**
  - Genotypic Diversity
  - Percentage Completed Trials

Soft Computing: Evolutionary Comp.: GA parameters control  
**SC Hybrid Systems: FLC Tuning EA**



# Fuzzy Logic Controlled GA (FLC-GA)



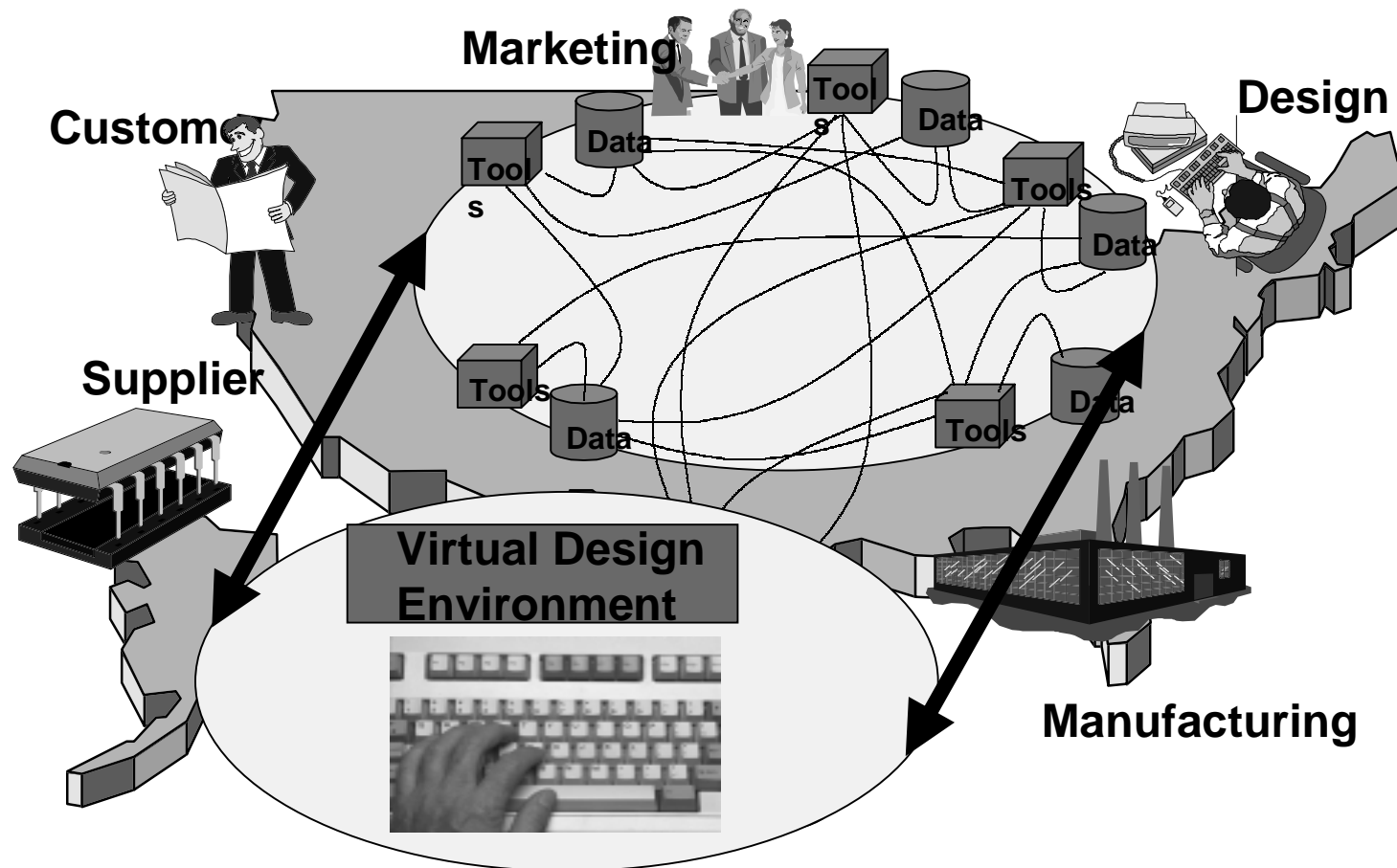
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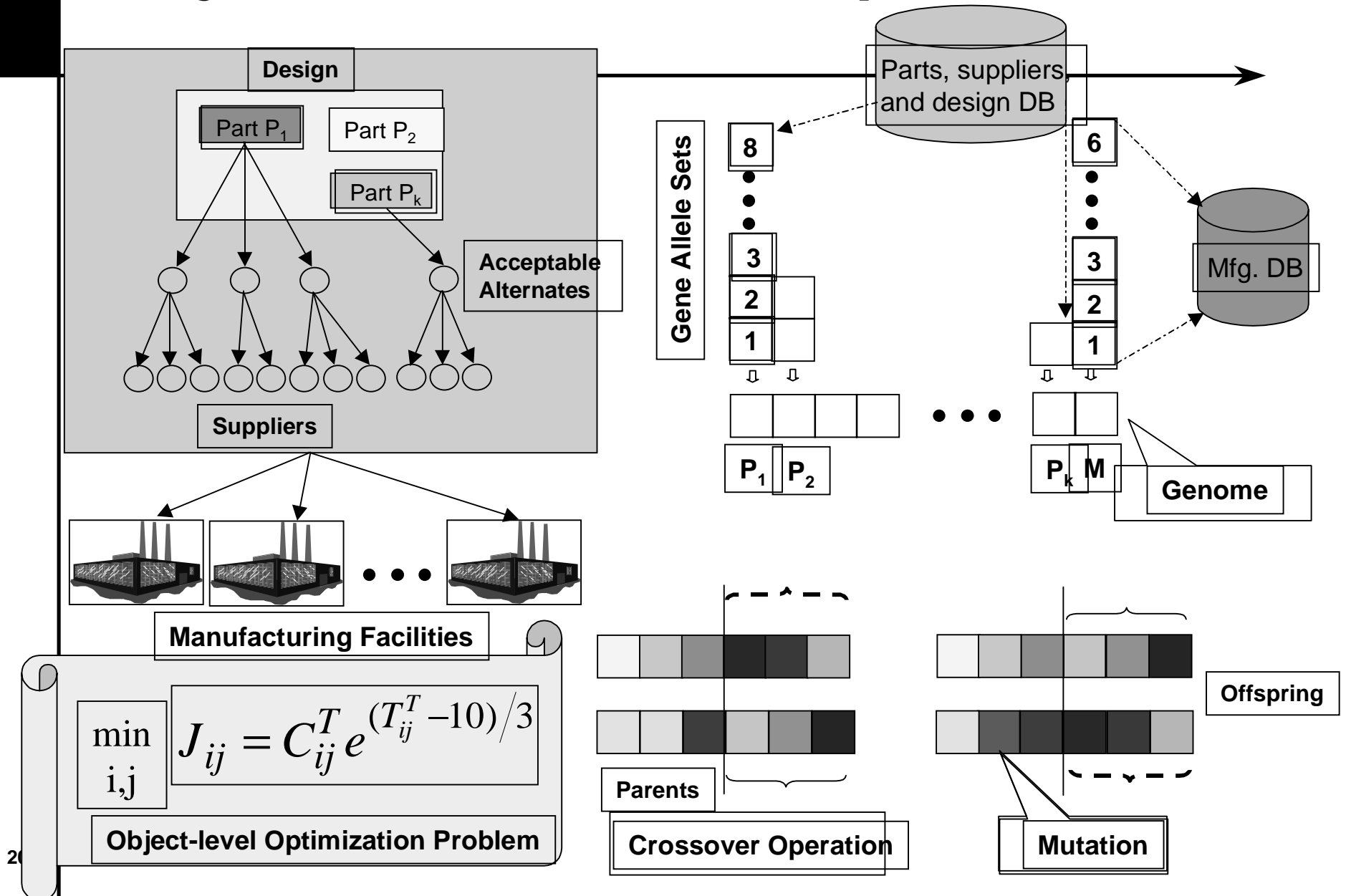


# GA Parameter Setting: An Application

Global optimization of design, manufacturing, supplier planning decisions in a distributed manufacturing environment



# Object-level Problem Representation



# Object-level Problem Complexity

## Search Space Size

- For off-line parameter tuning and system identification (Data Set 1):  $O(10^7)$
- For GA performance validation:  $O(10^{18})$  and  $O(10^{21})$

## Computer (to evaluate run-time performance)

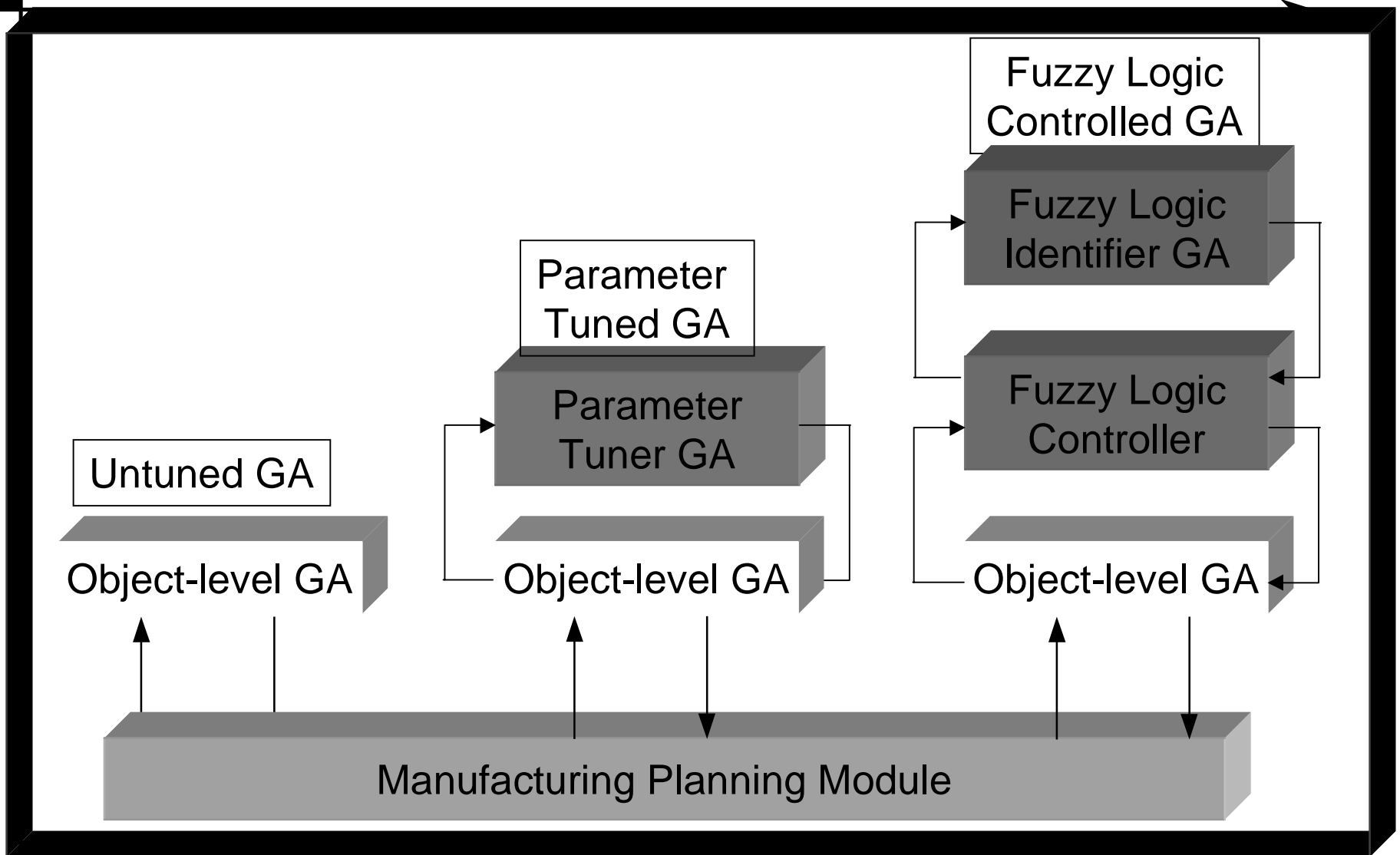
- SUN ULTRA SPARC-1

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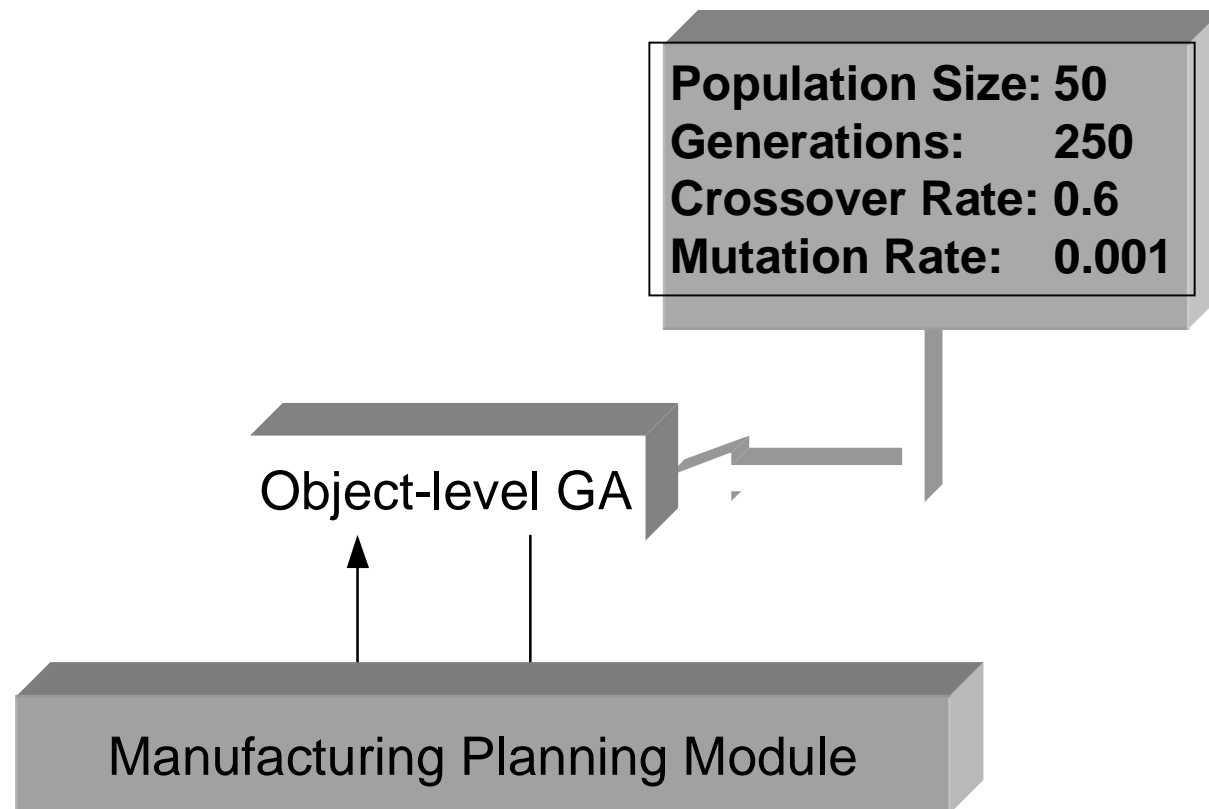
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# First Solution: Architecture



## Untuned GA (U-TGA)



# Parameter Tuned GA (TGA)

## Object GA Performance measure

**M = Overall Best Score +  
Average of Best Scores +  
Standard Deviation of the  
Final Population**

## Object GA Parameters

**Population Size  
Generations  
Crossover Rate  
Mutation Rate**

Parameter  
Tuner GA

Object-level GA

## **Best Results:**

Population Size: 134  
Generations: 250  
Crossover Rate: 0.648  
Mutation Rate: 0.008

Manufacturing Planning Module

# Fuzzy Controller for $\Delta N$ , $\Delta P_c$ , $\Delta P_m$ : Inputs and Outputs

## Inputs

GD = Genotypic Diversity (Normalized Average Hamming Distance)

$$GD = \frac{2}{n(n-1)} \sum_{i=1, j=i+1}^n \frac{d_{ij}}{\text{Genome Length}}$$

where  $d_{ij}$  is the Hamming Distance

GD range is [0, 1] == [Low, High]

PD = Phenotypic Diversity:

$$PD = \frac{f_b}{f_\mu} \quad \begin{array}{l} f_b = \text{Best Fitness} \\ f_\mu = \text{Average Fitness} \end{array}$$

PD range is [0, 1] == [Low, High]

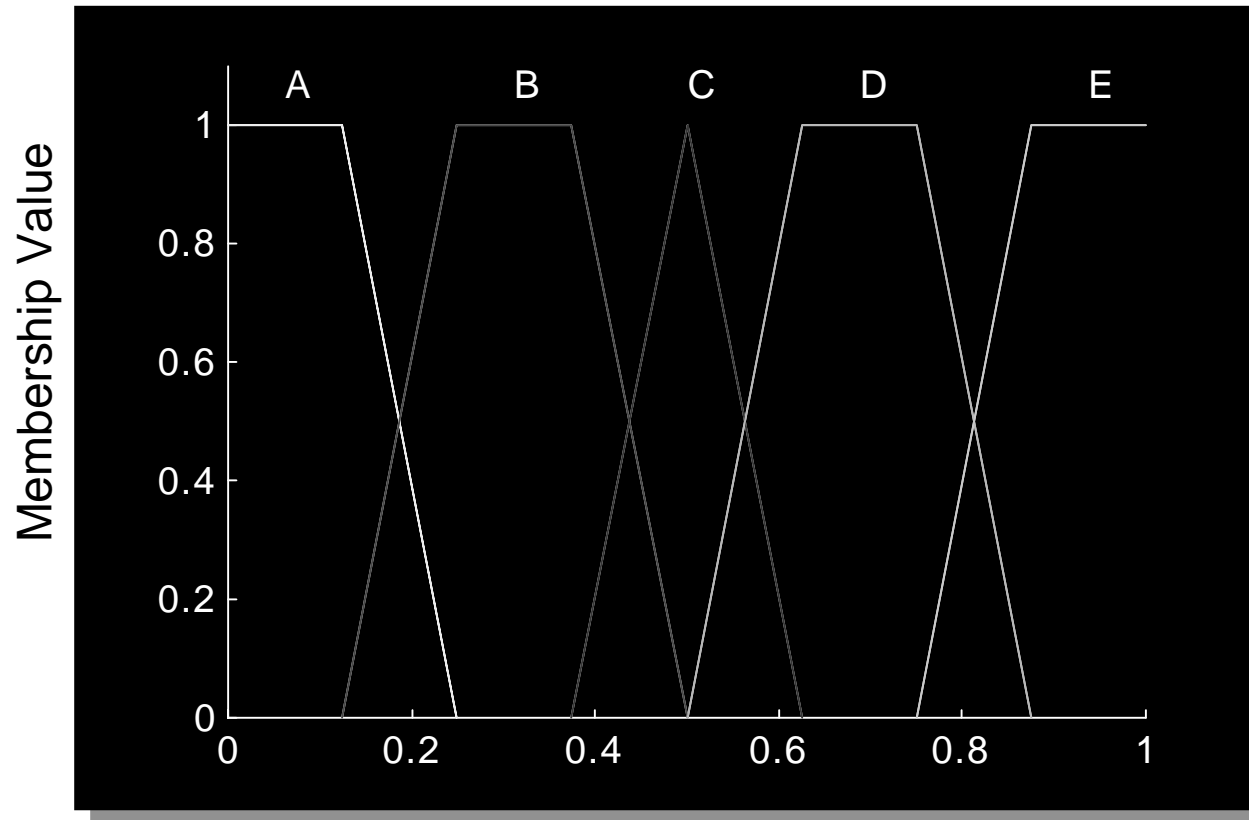
## Outputs

$\Delta$  Population Size

$\Delta$  Crossover Rate

$\Delta$  Mutation Rate

# Termset for State and Control Variables

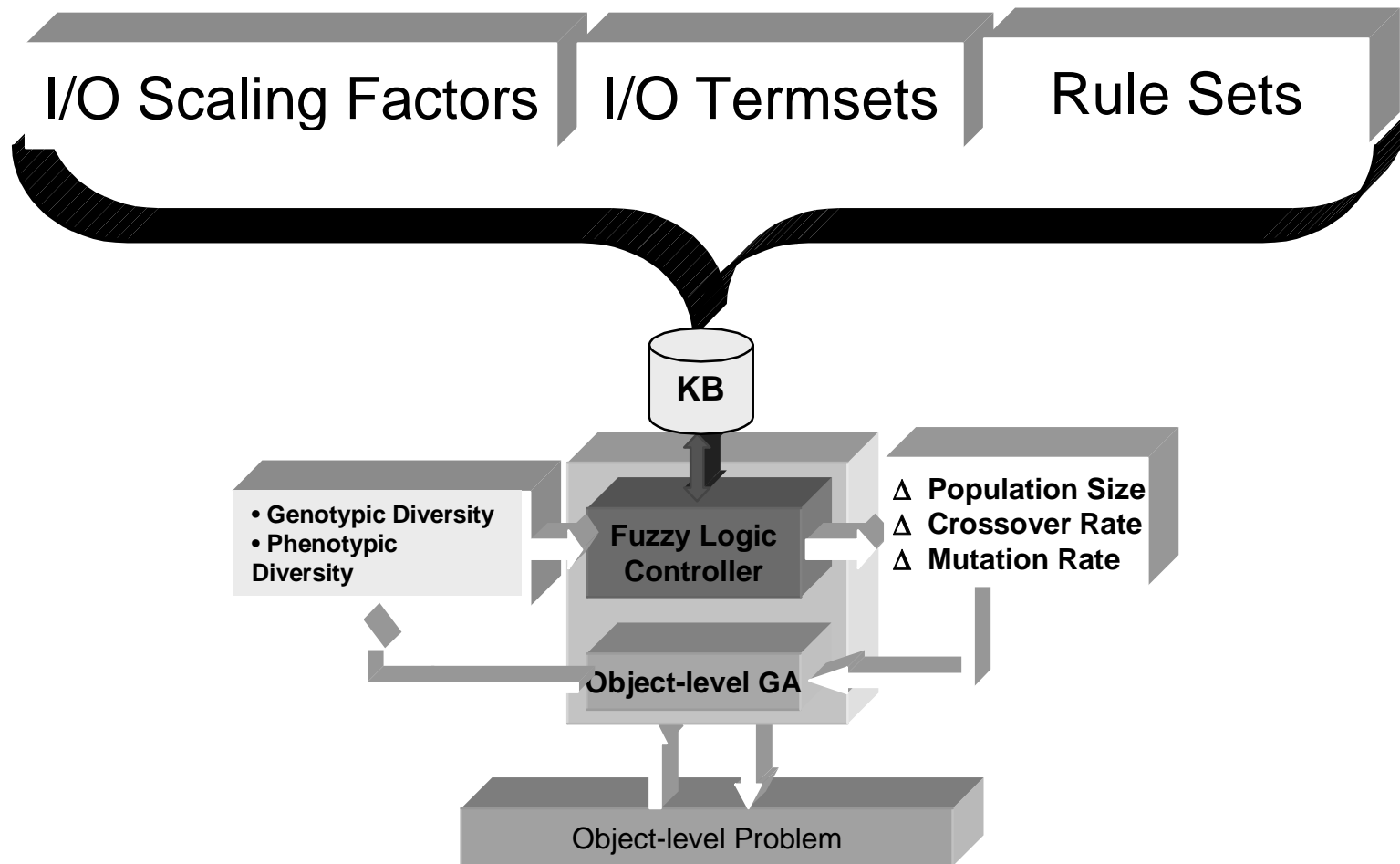


*GD*: A(Very Low), B(Low), C(Medium), D(High), E(Very High)

*PD*: A(Very High), B(High), C(Medium), D(Low), E(Very Low)

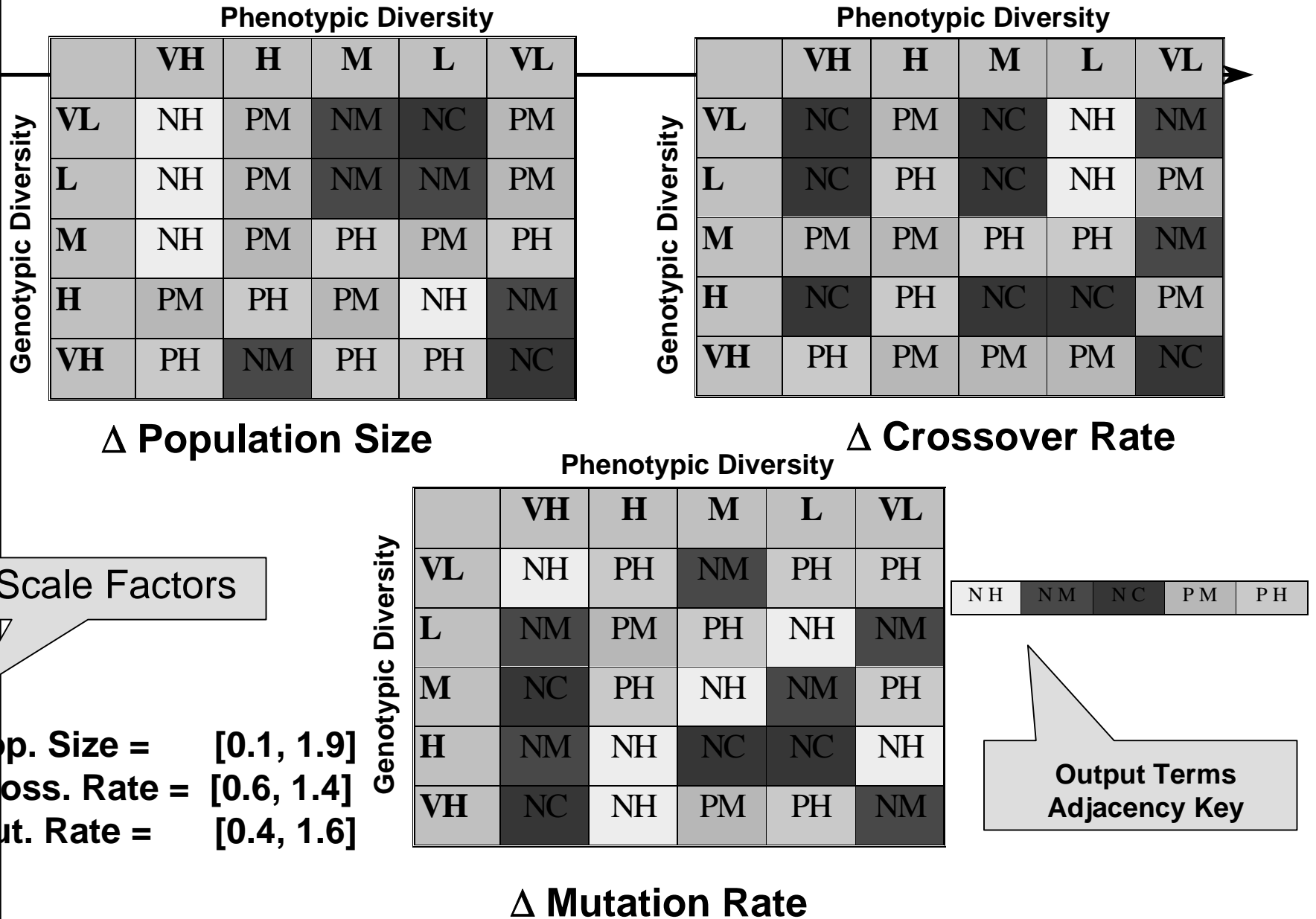
Outputs: A(Neg. High), B(Neg. Medium), C(No Change), D(Pos. Medium), E(Pos. High)

# Fuzzy Logic Controller for GAs: Knowledge Base





# Fuzzy Rule Base and Scaling Factors



# Fuzzy Controller for $\Delta N$ , $\Delta P_c$ , $\Delta P_m$ : Control Parameters

## Frequency of Control Actions

Control Action applied every 5 generations

## Mutation Rate

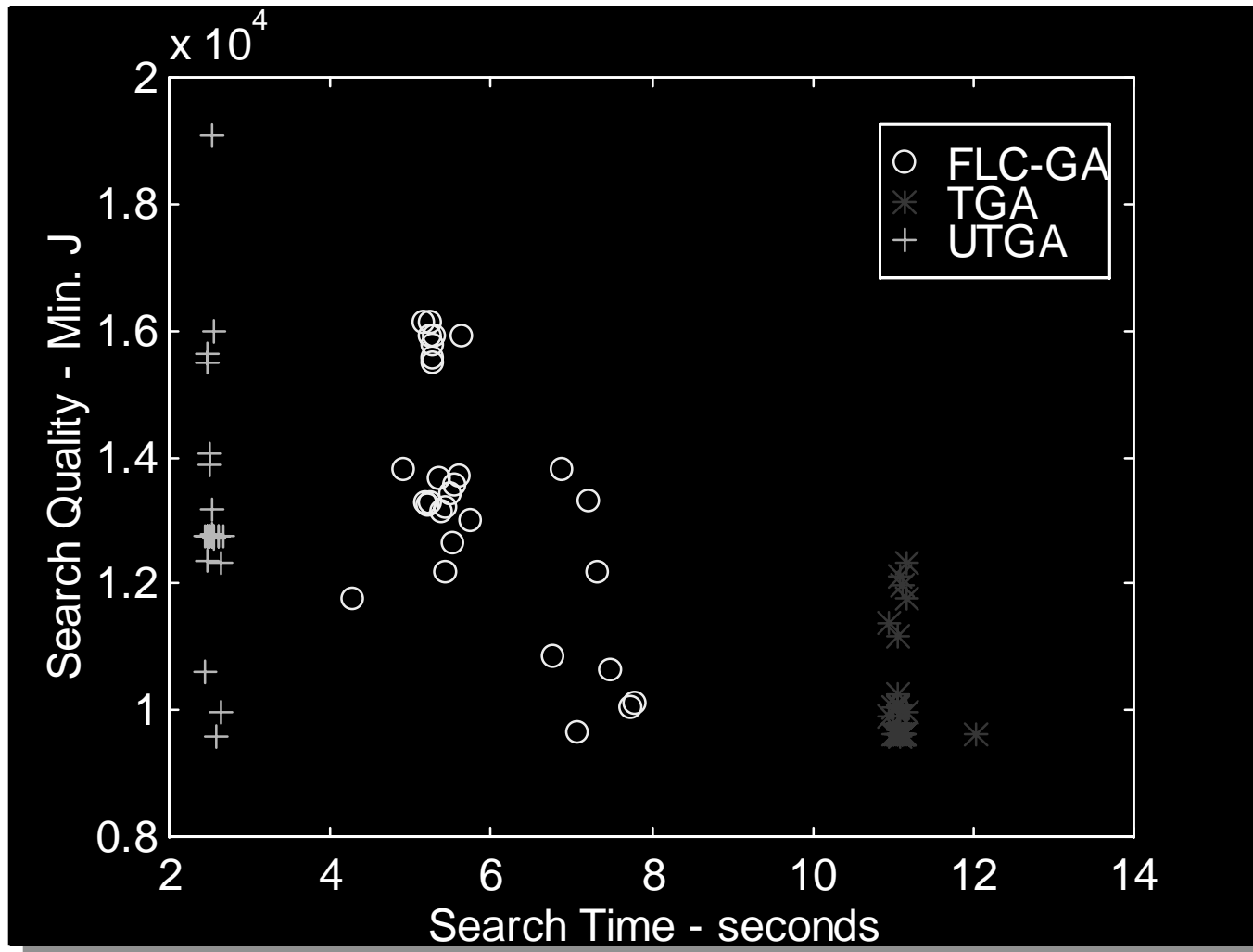
Mutation rates drops exponentially after a control action that increases it

## Inference Engine Parameters

Left Hand Side (LHS) evaluation:	<i>Minimum operator</i>
Rule Firing:	<i>Minimum operator</i>
Rule Output Aggregation:	<i>Maximum operator</i>
Defuzzification:	<i>Center of Gravity (COG)</i>

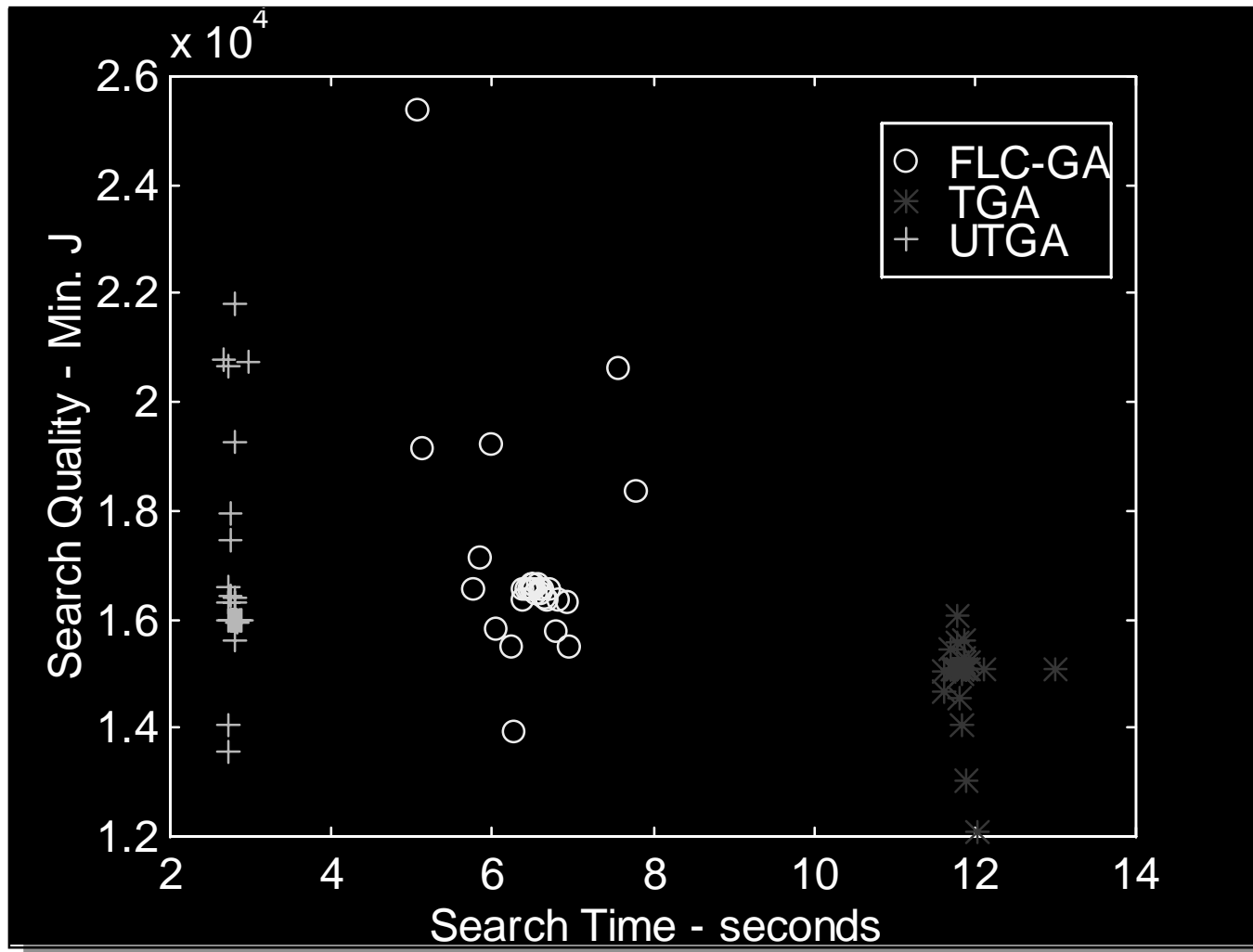


# Results - Performance Tradeoff - $O(10^{18})$



Benchmarked on search space of size  $O(10^{18})$

# Results - Performance Tradeoff - $O(10^{21})$



# Problems with First Solution

## FLC design parameter space:

### - Rule Set:

75 output rules - each rule can have one out of five values

Complexity (possibly different # rules):  $5^{75} \cong 2.64 \times 10^{52}$

### - Scaling Factors:

9 scaling factors - each one can have one out of nine values

Complexity (possibly different # scaling factors):  $9^9 \cong 3.87 \times 10^8$

### - Combining Complexity:

$5^{75} \times 9^9 \cong 1.02 \times 10^{61}$

**Larger than the object-level problem complexity!**

# Summary of 30 Experiments

Data Sets [Complexity]	GA Strategy	Min J			Time	
		$\mu$	$\sigma$	$\sigma/\mu$	$\mu$	$\sigma$
1 $O(10^7)$	FLC-GA	674	74	0.110	4.30	0.39
	TGA	551	23	0.042	6.37	0.08
	UTGA	923	272	0.295	1.45	0.08
2 $O(10^{18})$	FLC-GA	13380	1911	0.143	5.83	0.95
	TGA	10154	871	0.086	11.12	0.18
	UTGA	13048	1761	0.135	2.53	0.06
3 $O(10^{21})$	FLC-GA	17001	2006	0.118	6.46	0.56
	TGA	14939	738	0.049	11.86	0.24
	UTGA	16793	1946	0.116	2.78	0.06

## Analysis of Results

- **Offline computation time - using Data set 1 - for:**
  - **FLC rule set and scaling factors: 14 hrs**
  - **Meta-GA: 2 hrs**
- **Statically tuned Meta-GA (TGA):**
  - **slowest one but has the best search quality performance (J)**
- **Untuned GA (UTGA):**
  - **is the fastest but has the lowest search quality**
- **FLC controlled GA (FLC-GA):**
  - **has large spread of search quality but it is faster than TGA**

**Identified problems with FLC-GA approach that need to be solved**

## Addressing the Problems with First Solution

- ***Minimize high-level search space size for FLC-GA by***
  - ***Identify primary drivers (influences) of GA search***  
(use DOE to identify main effects and control only those drivers, minimizing number of controllers, hence number of parameters)
  - ***Control primary drivers by few simple heuristic rules***  
(eliminating learning of rule set and scaling factors)
- ***Optimal FLC firing rate?***
  - ***Reduction of firing frequency to reduce overhead***
- ***Extensive & statistically significant empirical evidence***
  - ***Let's use something we learned in  $6\sigma$  !***

# GA Parameter Setting - Outline

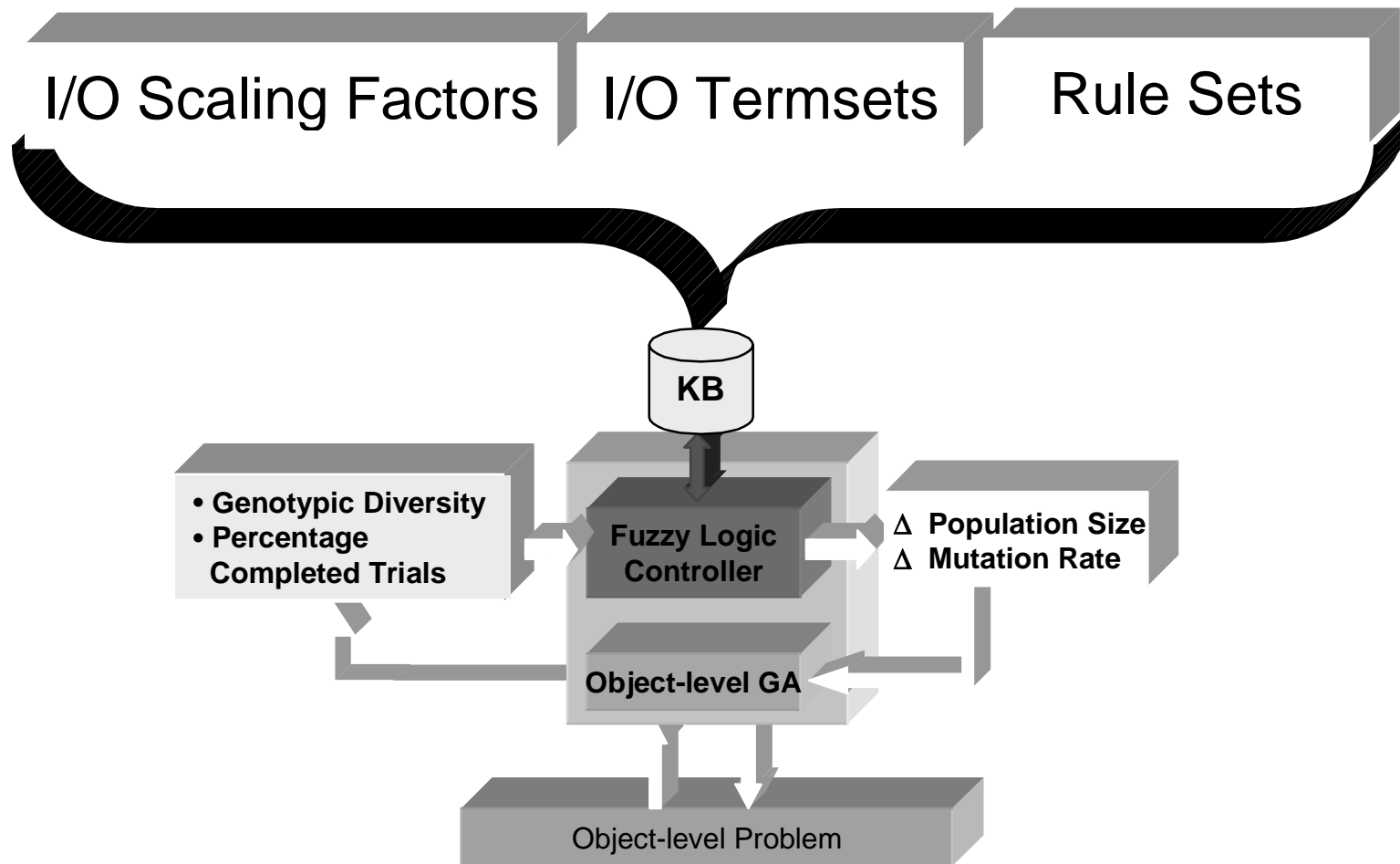
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## Second Solution

- **Minimize high-level search space size for FLC-GA by**
  - **Identify primary drivers (influences) of GA search**  
DOE determined that the two main drivers were:  
**Population Size ( $N$ ) and Mutation Rate ( $P_m$ )**
  - **Control primary drivers by few simple heuristic rules**  
Built two FLC controllers with heuristic rule sets and SF  
Changed on input (state variable) to capture evolution stage
- **Optimal FLC firing rate?**
  - **Reduction of firing frequency**  
From an action every 5 to one every 10 generation
- **Extensive & statistically significant empirical evidence**
  - **Use  $t$ -test and  $F$ -tests to analyze  $\mu$  and  $\sigma$  improvements**

# Fuzzy Logic Controller for GAs: Knowledge Base



# Fuzzy Controller for $\Delta N$ and $\Delta P_m$ : Inputs

## Inputs

GD = Genotypic Diversity:

Normalized Average Hamming Distance

$$GD = \frac{2}{n(n-1)} \sum_{i=1, j=i+1}^n \frac{d_{ij}}{\text{Genome Length}}$$

where  $d_{ij}$  is the Hamming Distance  
 GD range is [0, 1] == [Low, High]

PFE = Percentage Fitness Evaluations:

(Completed # Trials) / (Max Allocated # Trials)

PFE range is [0, 1] == [Low, High]

## Fuzzy Controller for $\Delta N$ and $\Delta P_m$ : Outputs

### Outputs

$\Delta N$  = Change in Population Size (Mult. Factor)

$\Delta N$  range is [0.5, 1.5] == [Neg High, Pos High]  
so that NC corresponds to 100% of previous Pop Size

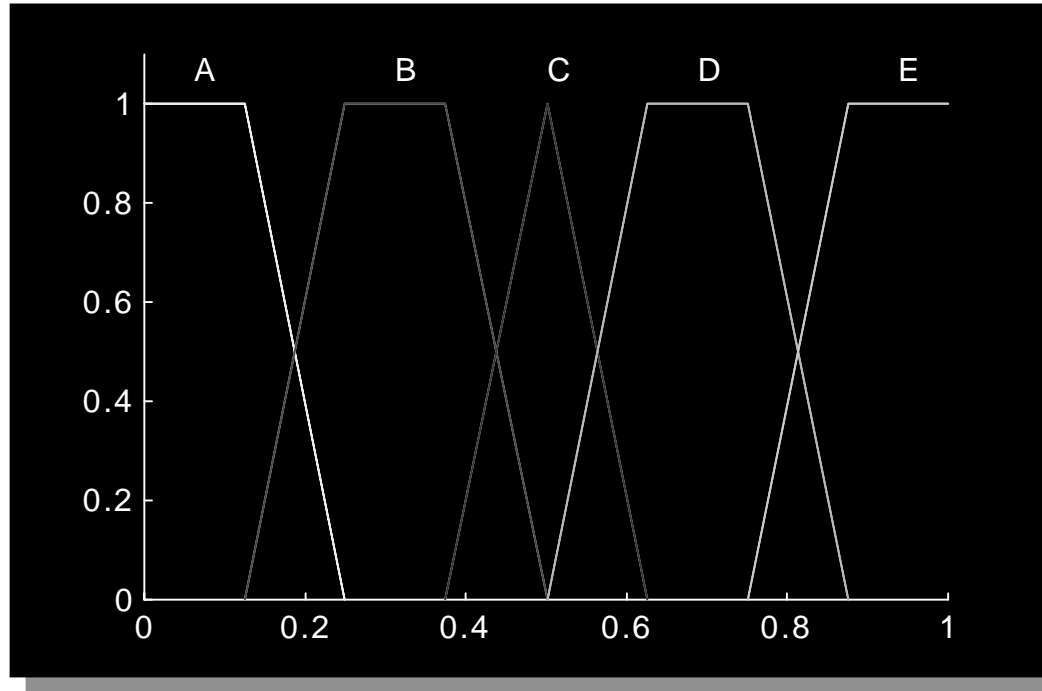
Population Size is *clamped* within [25, 150]

$\Delta P_m$  = Change in Mutation Rate (Mult. Factor)

$\Delta P_m$  range is [0.5, 1.5] == [Neg High, Pos High]  
so that NC corresponds to 100% of previous  $P_m$

Mutation Rate is *clamped* within [0.005, 0.10]

# Fuzzy Controller for $\Delta N$ and $\Delta P_m$ : Termsets



**Inputs:**

**GD:** A(Very Low), B(Low), C(Medium), D(High), E(Very High)

**PFE:** A(Very Low), B(Low), C(Medium), D(High), E(Very High)

**Outputs (for both  $\Delta N$  and  $\Delta P_m$ ):**

A(Neg. High), B(Neg. Medium), C(No Change), D(Pos. Medium),  
E(Pos. High)

# Fuzzy Controller for Population Size: Rule Set

GD, PFE  $\rightarrow$   $\Delta N$

GD = Genotypic Diversity:  
Normalized Average Hamming Distance

PFE = Percentage Fitness Evaluations:  
(Completed # Trials) / (Max Allocated # Trials)

$\Delta N$  = Change in Population Size

Genotypic Diversity (GD)	Percentage Fitness Evaluation (PFE)				
	<i>Very Low</i>	<i>Low</i>	<i>Medium</i>	<i>High</i>	<i>Very High</i>
<i>Very Low</i>	Pos High	Pos High	Pos High	Pos Medium	No Change
<i>Low</i>	Pos High	Pos High	Pos Medium	No Change	Neg Medium
<i>Medium</i>	Pos High	Pos Medium	No Change	Neg Medium	Neg High
<i>High</i>	Pos Medium	No Change	Neg Medium	Neg High	Neg High
<i>Very High</i>	No Change	Neg Medium	Neg High	Neg High	Neg High

**Exploration Stage**  
Increase population/ broaden search

**Exploitation Stage**  
Reduce population/ Refine best

# Fuzzy Controller for Mutation Rate: Rule Set

GD, PFE  $\rightarrow$   $\Delta P_m$

- GD = Genotypic Diversity:  
Normalized Average Hamming Distance
- PFE = Percentage Fitness Evaluations:  
(Completed # Trials) / (Max Allocated # Trials)
- $\Delta P_m$  = Change in Mutation Rate

Genotypic Diversity (GD)	Percentage Fitness Evaluation (PFE)				
	<i>Very Low</i>	<i>Low</i>	<i>Medium</i>	<i>High</i>	<i>Very High</i>
<i>Very Low</i>	Pos High	Pos High	Pos Medium	Pos Medium	No Change
<i>Low</i>	Pos High	Pos Medium	Pos Medium	No Change	No Change
<i>Medium</i>	Pos Medium	Pos Medium	No Change	No Change	No Change
<i>High</i>	Pos Medium	No Change	No Change	No Change	No Change
<i>Very High</i>	No Change	No Change	No Change	No Change	No Change

# Fuzzy Controller for $\Delta N$ and $\Delta P_m$ : Control Parameters

## Frequency of Control Actions

Control Action:

- mutation rate changed every 10 generations
- population size change every generation

## Mutation Rate

Mutation rates drops exponentially after a control action that increases it

## Inference Engine Parameters

Left Hand Side (LHS) evaluation:	<i>Minimum operator</i>
Rule Firing:	<i>Minimum operator</i>
Rule Output Aggregation:	<i>Maximum operator</i>
Defuzzification:	<i>Center of Gravity (COG)</i>

# Statistical Experiments: Set-Up

## Data Set for Experiments

- Seven part classes corresponding to a complexity of  $O(10^7)$

## GA Structure:

- Type: {Simple, Steady-State}
- Chromosome Encoding: Integer
- Fitness Function: Three type of cost functions
- Selection Method: Proportional Roulette
- Crossover Operator: Uniform
- Mutation Operator: Exponentially Decreasing

## GA Parameters

- $N$  = Base Population size: 50
- $P_c$  = Crossover rate: 0.6
- $P_m$  = Mutation rate: 0.005
- $G$  = Generation Gap  
100% replac. - Simple GA (SGA)  
25% replac. - Steady State GA (SSGA)
- $S$  = Selection Strategy: Elitist

# Statistical Experiments: Set-Up

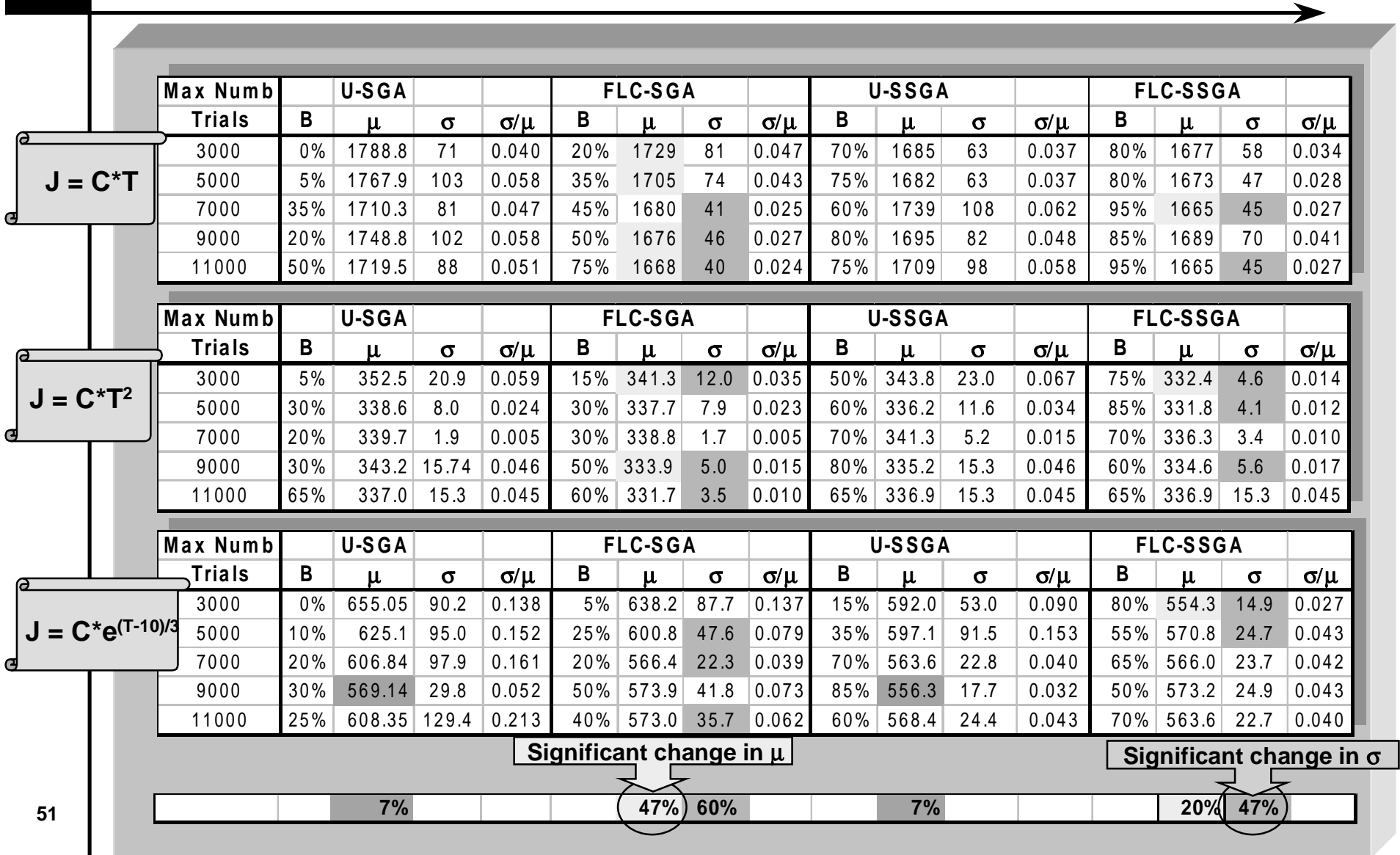
## Set-Up for 1200 experiments:

- We defined 4 GA configurations
  - (a) Untuned Simple GA (U-SGA)
  - (b) FL Controlled Simple GA (FLC-SGA)
  - (c) Untuned Steady State GA (U-SSGA)
  - (d) FL Controlled Steady State GA (FLC-SSGA)
- For each configuration we performed 300 experiments:
  - 20 runs for each pair of (Cost function, Max number of Trials)
  - 15 different pairs of (Cost function, Max number of Trials)
    - Three types of *cost functions*:  
**(1)  $J = C \cdot T$ ; (2)  $J = C \cdot T^2$ ; (3)  $J = C \cdot e^{(T-10)/3}$**
    - Five values of maximum *number of Trials* (to evaluate effect of control for different evolution lengths):  
**(i) 3,000; (ii) 5,000 (iii) 7,000 (iv) 9,000 (v) 11,000**

# Statistical Experiments: Measures

- For each of the four configuration (a-d) we ran 20 experiments with the same parameters
- Then we considered the following measures:
  - $\hat{B}$  = sample average *over 20 experiments* of Best score frequency (number of time cost function J reached its minimal value - known a priori for small size experiment)
  - $\hat{\mu}$  = average of population best
  - $\hat{\sigma}$  = standard deviation of population best
- **We performed an ANOVA test (using both t and F test -using the usual value of  $p < 0.05$  to identify statistical significance) to see how many times the following statements were true:**
  - Cost (U-SGA) significantly higher than cost ( FLC-SGA)
  - Cost (U-SGA) significantly higher than cost ( U-SSGA)
  - Cost (U-SSGA) significantly higher than cost ( FLC-SSGA)
- **Finally we verified if the FLC caused the GA to perform any worse than its corresponding untuned GA, i.e.:**
  - Cost (U-SGA) significantly lower than cost ( FLC-SGA)
  - Cost (U-SSGA) significantly lower than cost ( FLC-SSGA)

# Summary of 1200 Experiments



# Statistical Experiments: Summary

Analysis of  $\hat{\mu}$  for controlled GAs with Untuned GAs:

- FLC-SGA significantly better than U-SGA: **47%** of the times (7/15)
- FLC-SSGA significantly better than U-SSGA: **20%** of the times (3/15)
- FLC-SGA significantly worse than U-SGA: **7%** of the times (1/15)
- FLC-SSGA significantly worse than U-SSGA: **7%** of the times (1/15)

Analysis of  $\hat{\sigma}$  for controlled GAs with Untuned GAs:

- FLC-SGA significantly better than U-SGA : **60%** of the times (9/15)
- FLC-SSGA significantly better than U-SSGA: **47%** of the times (7/15)
- FLC-SGA significantly worse than U-SGA: **Never**
- FLC-SSGA significantly worse than U-SSGA : **Never**

Comparison of Untuned GAs (Simple vs SS):

- $\hat{\mu}$  U-SSGA significantly better than U-SGA: **15%** of the times
- $\hat{\sigma}$  U-SSGA significantly better than U-SSGA: **30%** of the times

# GA Parameter Setting - Outline

- GA Model: Structure, Parameters
- GA Parameter Setting
  - GA Parameter Tuning: Off-line study, Off-line Meta-GA
  - GA Parameter Control: Deterministic Control, Self-Adaptation, Control (FLC)
- GA Parameter Setting: An Application
  - Object-level Representation and Complexity
- First Solution Architecture (U-TGA, TGA, FLC-GA)
  - FLC-GA: KB, I/O, Control Parameters, Rule base and Scaling Factors
  - Results - Performance Tradeoff
  - Problems with First Solution
  - Summary and Analysis of 30 Experiments
  - Addressing Problems with First Solution
- Second Solution
  - FLC: KB, Inputs, Outputs, Termsets, Rule Sets, Control Parameters
  - Statistical Experiments: Set-Up & Measures
  - Summary and Analysis of 1200 Experiments
- Conclusions and Next Steps



# Conclusions

## FLC State Representation: [Evolution Stage]

- Since the objectives of the GA are different in the earliest and latest stages, we need time to be an explicit state variable
- Diversity is an essential measure of the evolutionary stage:
  - Percentage Fitness Evaluations (PFE)
  - Genotypic Diversity (GD)

## FLC Control Variables: [GA Adaptable Parameters (Subset )]

- $\Delta N$  = Change in Population Size
- $\Delta P_m$  = Change in Mutation Rate

## Main Result

- By using the FLC with the above State and Control variables, we achieved a good improvement of the population average and an even better improvement of the population variance.
- No major negative effects on GA performance using FLC

# Next Steps: Multi-Criteria Fitness Functions

- **Scalar-Valued Fitness Functions**
  - Aggregated fitness function focuses on one tradeoff point in frontier
    - Use of *variable weights* in multi-criteria aggregating function
    - **Dynamic Control of weights over time**
- **GA Generation of Pareto Frontier**
  - **VE-GA:** Vector Evaluated GA (*Shaffer & Grefenstette 1985*)
    - **Multi-sexual GA** (*Lis & Elben 1997*)
  - **VOW-GA:** Variable Objective Weighting GA (*Hajela & Lin 1992*)
  - **MO-GA:** Multiobjective Optimization GA (*Fonseca & Fleming 1994*)
  - **NP-GA:** Niche Pareto GA (*Horn, Nafpilotis, Goldberg, 1994*)
  - **NS-GA:** Non Dominated Sorting GA (*Srinivas & Deb, 1995*)
  - **RW-GA:** Random Weights GA (*Ishibushi & Murata, 1998*)
- **Great parameter control opportunity!**

# Next Steps: Multi-Criteria Fitness Functions

## GA Generation of Pareto Frontier (Past Work)

### - VE-GA: Vector Evaluated GA (*Shaffer & Grefenstette 1985*)

- $N/k$  sub-populations created (where  $k$  is the number of criteria;  $N = \text{Pop-Size}$ )
- Modified selection operator performs proportional selection for each sub-population according to each objective function
- Sub-population shuffled to generate population of size  $N$
- Crossover and Mutation applied to population

Problems with “speciation” (sub-population excelling in only one aspect of fitness)

Problems with maintaining “middlings” (well-round individual not outstanding in any specific aspect of fitness)

Suggested: Heuristic selection to protect middling + heuristic crossbreeding of species

Equivalent to linear combination of objectives - Cannot generate Pareto-optimal solutions in the presence of non-convex search spaces.

# Next Steps: Multi-Criteria Fitness Functions

## GA Generation of Pareto Frontier (Past Work)

- **VE-GA:** Vector Evaluated GA (*Shaffer & Grefenstette 1985*)
  - **Multi-sexual GA** (*Lis & Elben 1997*)

A variation of VE-GA:

- A fitness function for each criterion
- Gender-tagged genotypes - a gender for each criterion
- Multiple-parents recombination operator
  - Replace aggregation of multiple fitness functions with aggregation of multiple genetic material

NEEDS

- **Dynamic determination of offspring gender**
- **Dynamic control of genders distribution in population**

# Next Steps: Multi-Criteria Fitness Functions

## GA Generation of Pareto Frontier (Past Work)

### - **VOW-GA:** Variable Objective Weighting GA (*Hajela & Lin 1992*)

- Each objective is assigned a (non-negative, normalized) weight
- Scalar fitness function is computed with weighted average of objectives
- Weights are encoded in genotype
- Diversity of weight combinations promoted by phenotypic fitness “sharing”
- EA evolves solutions and weights simultaneously

### - **RW-GA:** Random Weights GA (*Ishibushi & Murata, 1998*)

- Each pair of parents is randomly assigned a weight vector used in their offspring evaluation
- The (non-negative, normalized) weights represent a direction in the search
- Each individual 's fitness is computed by a weighted average of objectives
- Each offspring is locally improved by a k-neighborhood search around it
- At each generation, a provisional set of non-dominated solutions is identified, stored, and updated.
- A subset of these non-dominated solutions are re-introduced in population (elitism)

# Next Steps: Multi-Criteria Fitness Functions

## GA Generation of Pareto Frontier (Past Work)

- **MO-GA:** Multiobjective Optimization GA (*Fonseca & Fleming 1994*)  
{description - to be continued}
- **NP-GA:** Niche Pareto GA (*Horn, Nafpitolis, Goldberg, 1994*)  
{description - to be continued}
- **NS-GA:** Non Dominated Sorting GA (*Srinivas & Deb, 1995*)  
{description - to be continued}

## More Next Steps: Controlling Other Parameters

- **Run-time Controlled GAs Parameters:**

DONE

- Population size:

- larger size: increase parallel search in solution space
- smaller size: focus on current existing regions

DONE

- Probability mutation:

- Higher prob. of mutation disrupts current solutions - exploration
- Lower probability of mutation favors current solutions - exploitation

- **Other Possible Run-time Controllable GAs Parameters:**

- Customized mutation operators:

- Variable amount of changes
  - smaller for good solutions, larger for bad ones

- Fitness function:

- Evolving fitness function (variable weights in multi-criteria aggregating function)

## GAs controlled by FL (cont.)

- **Probability of Selection:**
  - Parametrized slope distribution ranging from:
    - **Uniform** probability: ignore fitness function and perform random selection of parents - extreme case of exploration, to
    - **Proportional** selection with rescaling and other intermediate strategies - compromise between exploration and exploitation cases, and
    - **Ranking**: always select the best N and ignore the rest - extreme case of exploitation
  - Probability as function of fitness and genotypical distance with other solutions - enforcing diversity and favoring exploration
- **Probability of crossover:**
  - Constraints applicability to mostly good solutions
- **Customized-crossover operators** (for real-coded GAs):
  - Selection of crossovers based on T-norms and T-conorms causes offsprings to take more extreme values (exploration)
  - Selection of crossovers based on aggregating operators causes offsprings to take average values (exploitation)