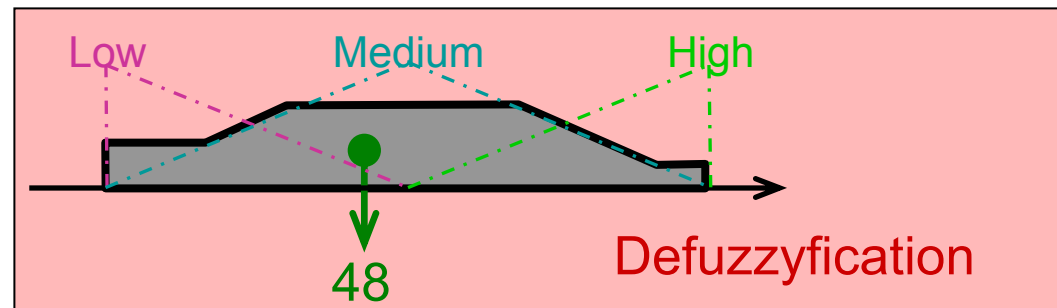
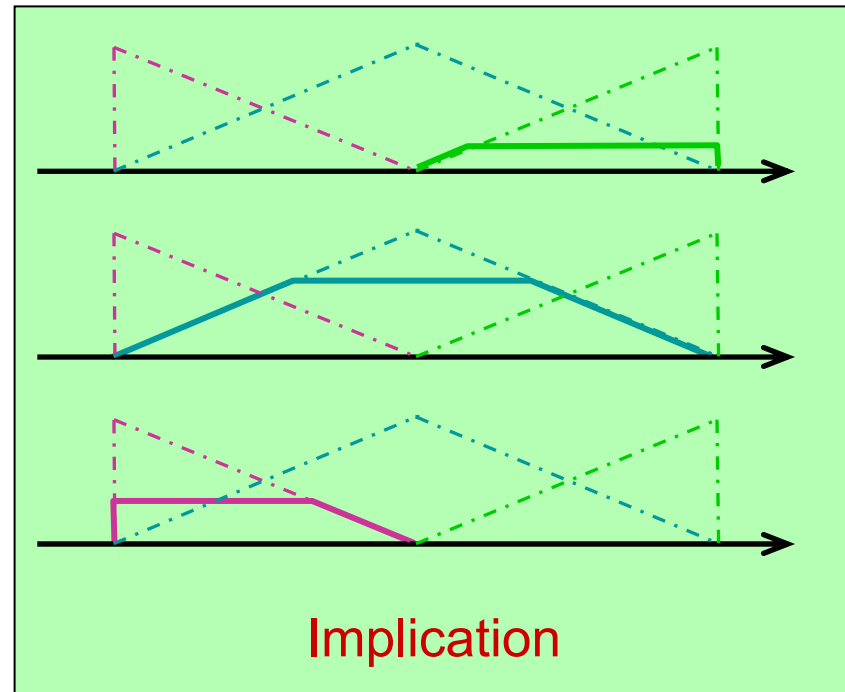
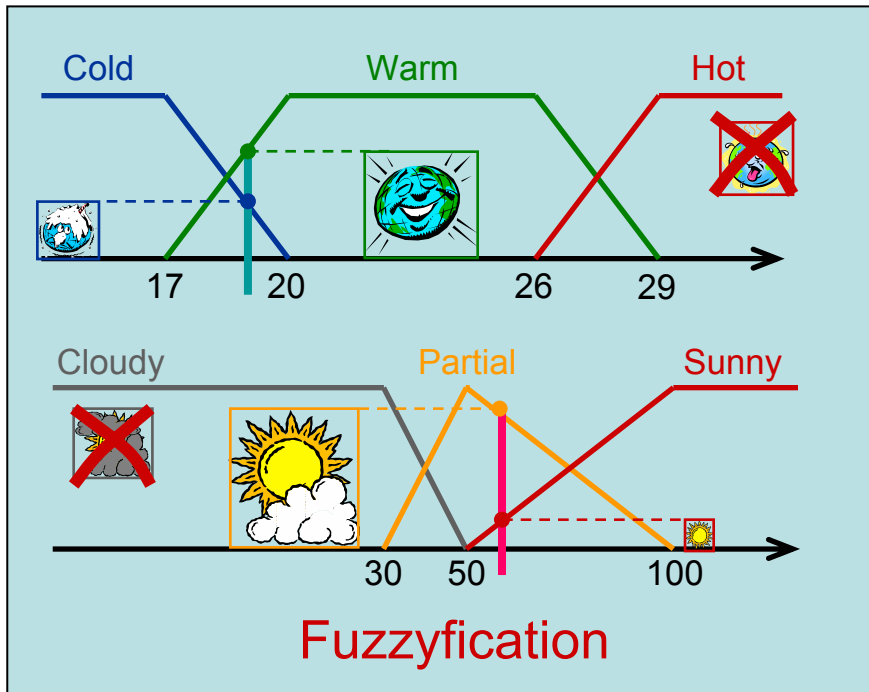
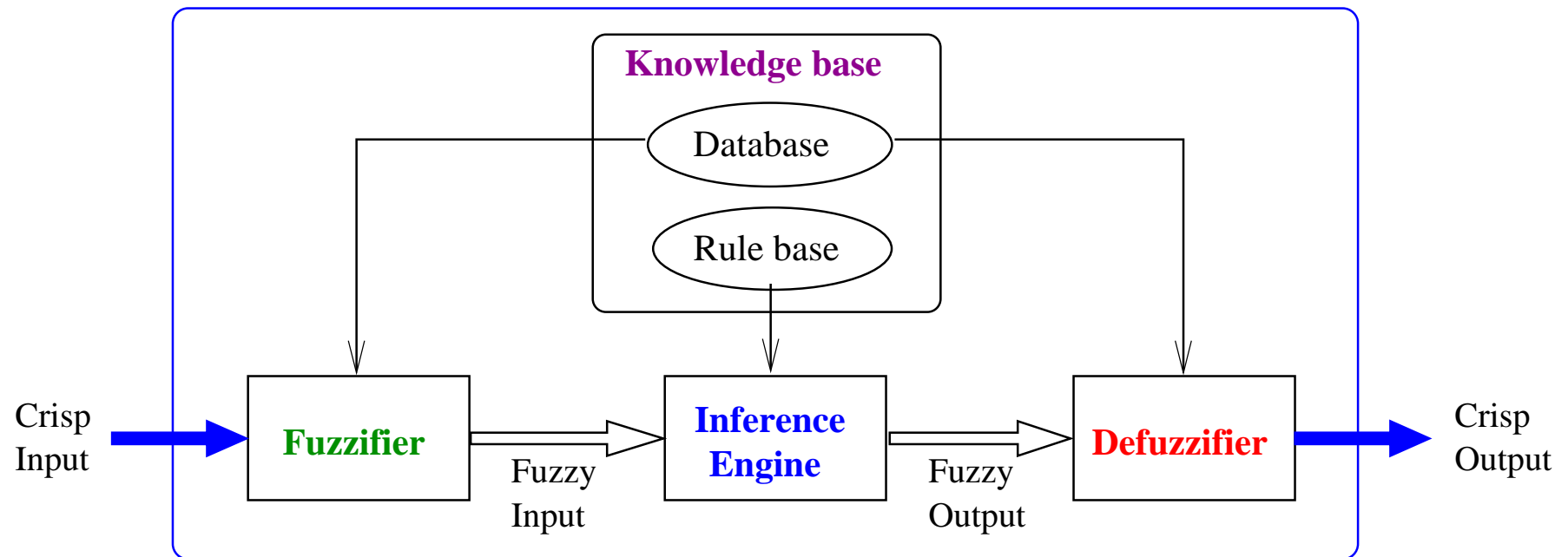


Fuzzy inference



Fuzzy inference systems



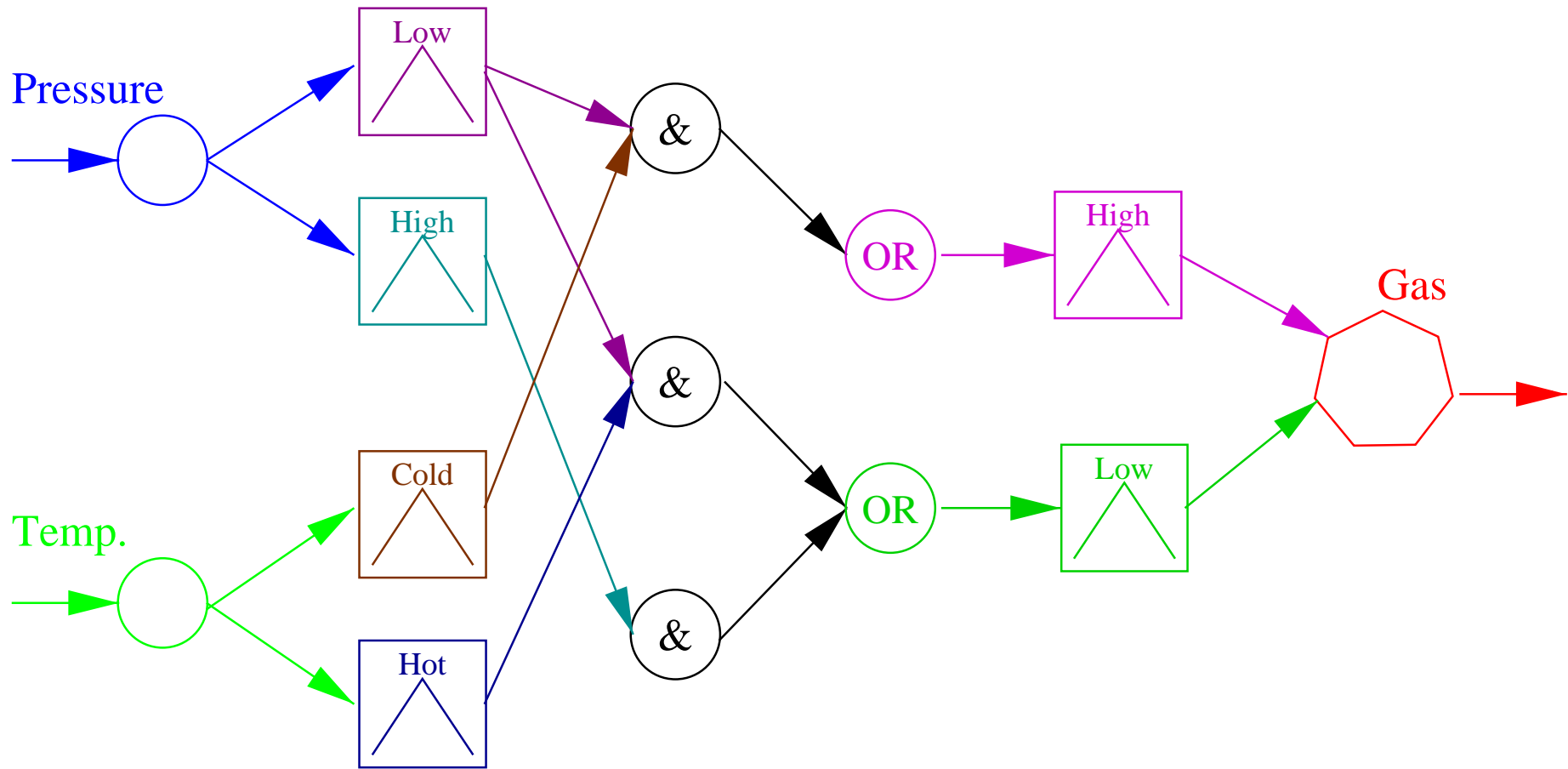
Fuzzifier: translates crisp inputs into fuzzy values

Inference engine: applies reasoning to compute fuzzy outputs

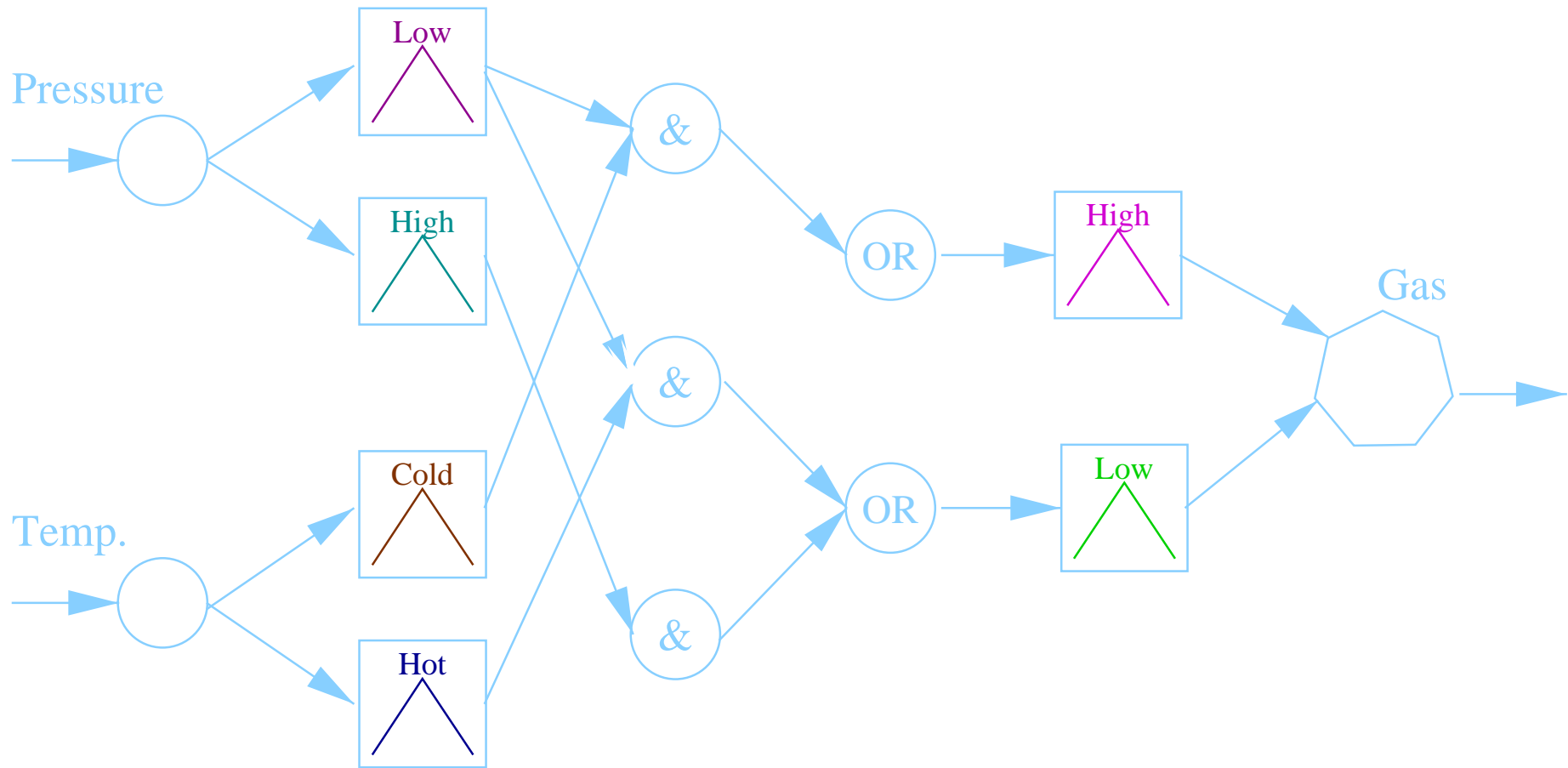
Defuzzifier: translates fuzzy outputs into crisp values

Knowledge base: defines rules and membership functions

Network-like view of a fuzzy system

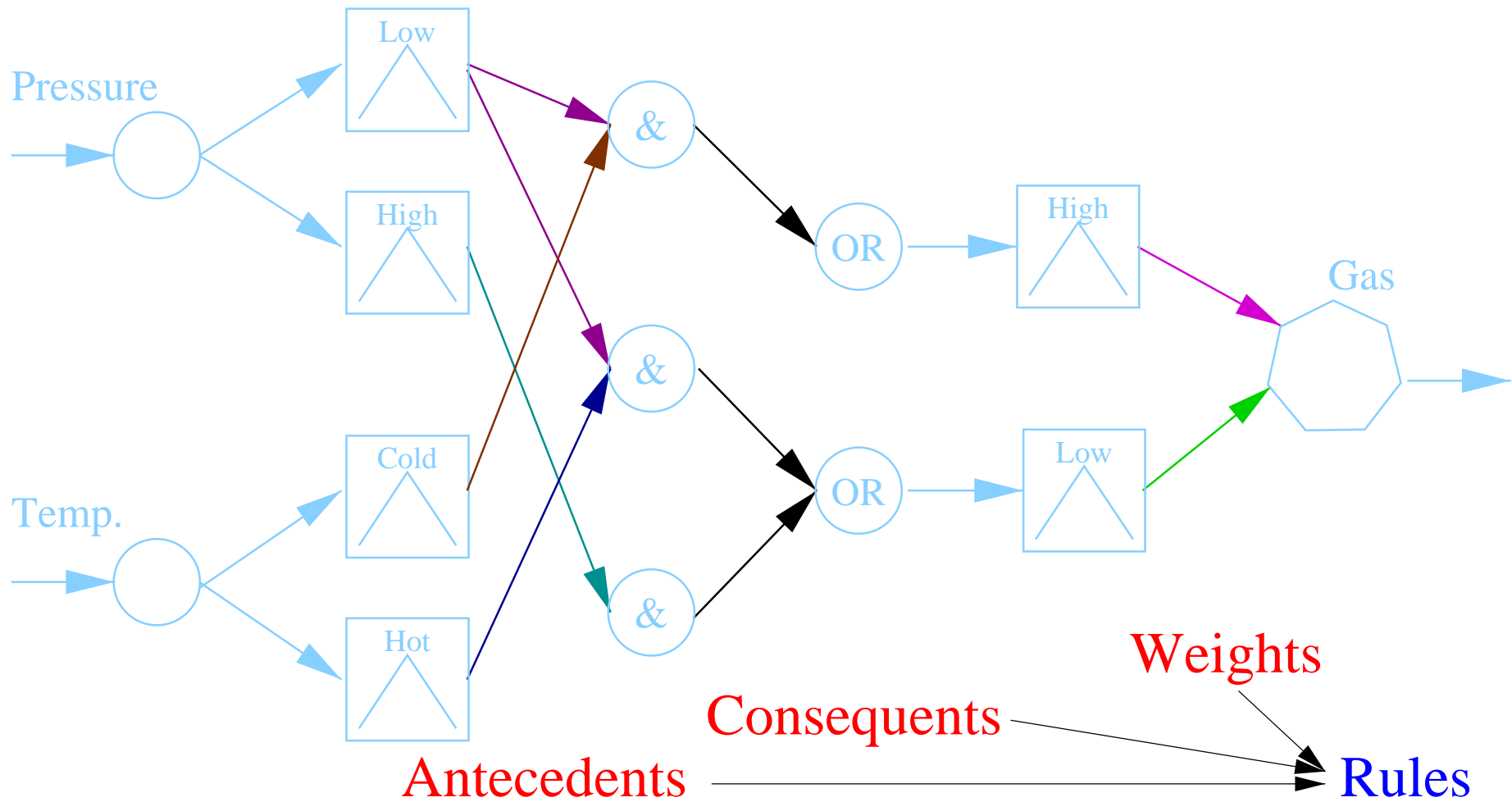


Operational parameters

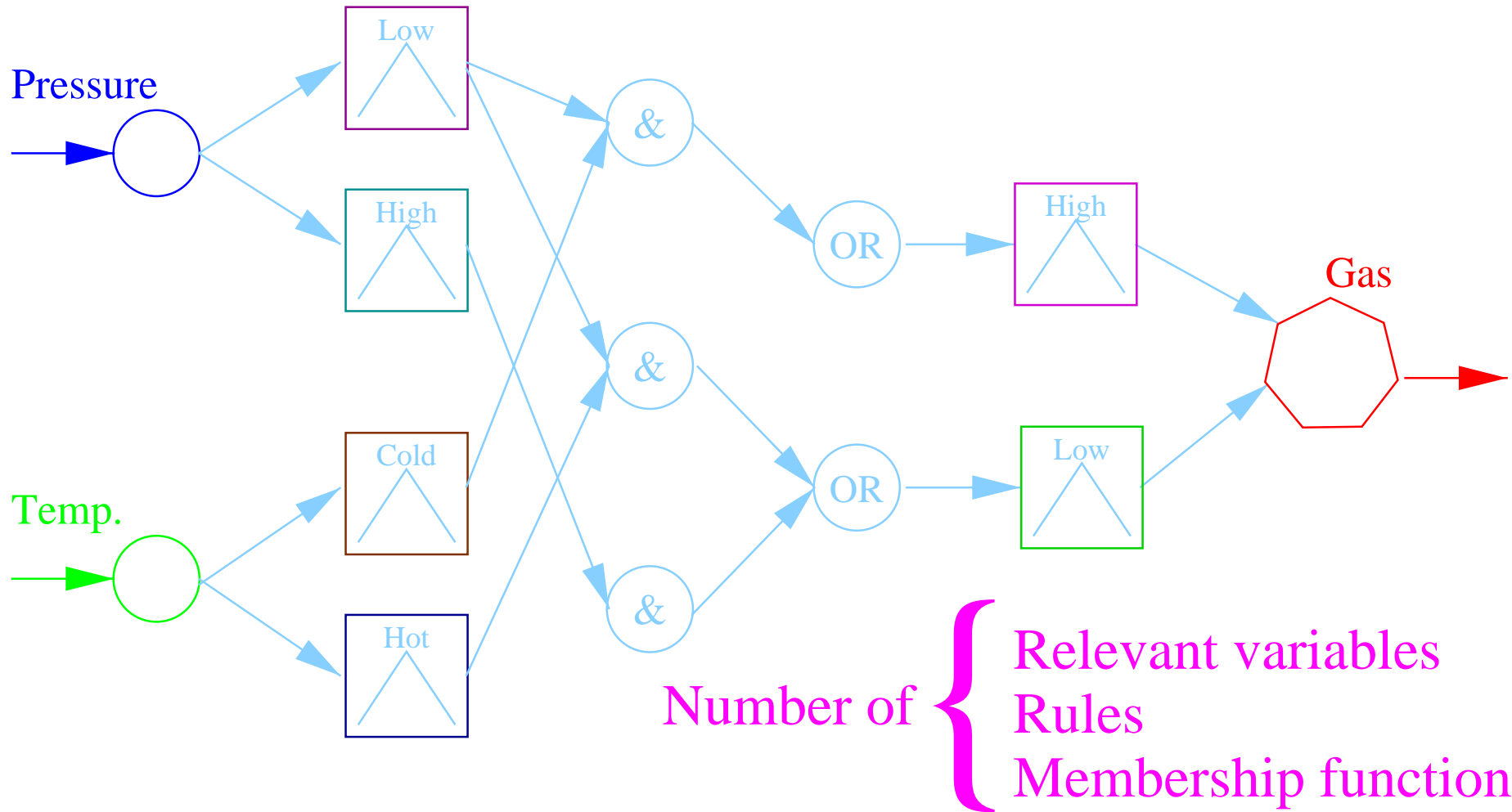


Membership function values

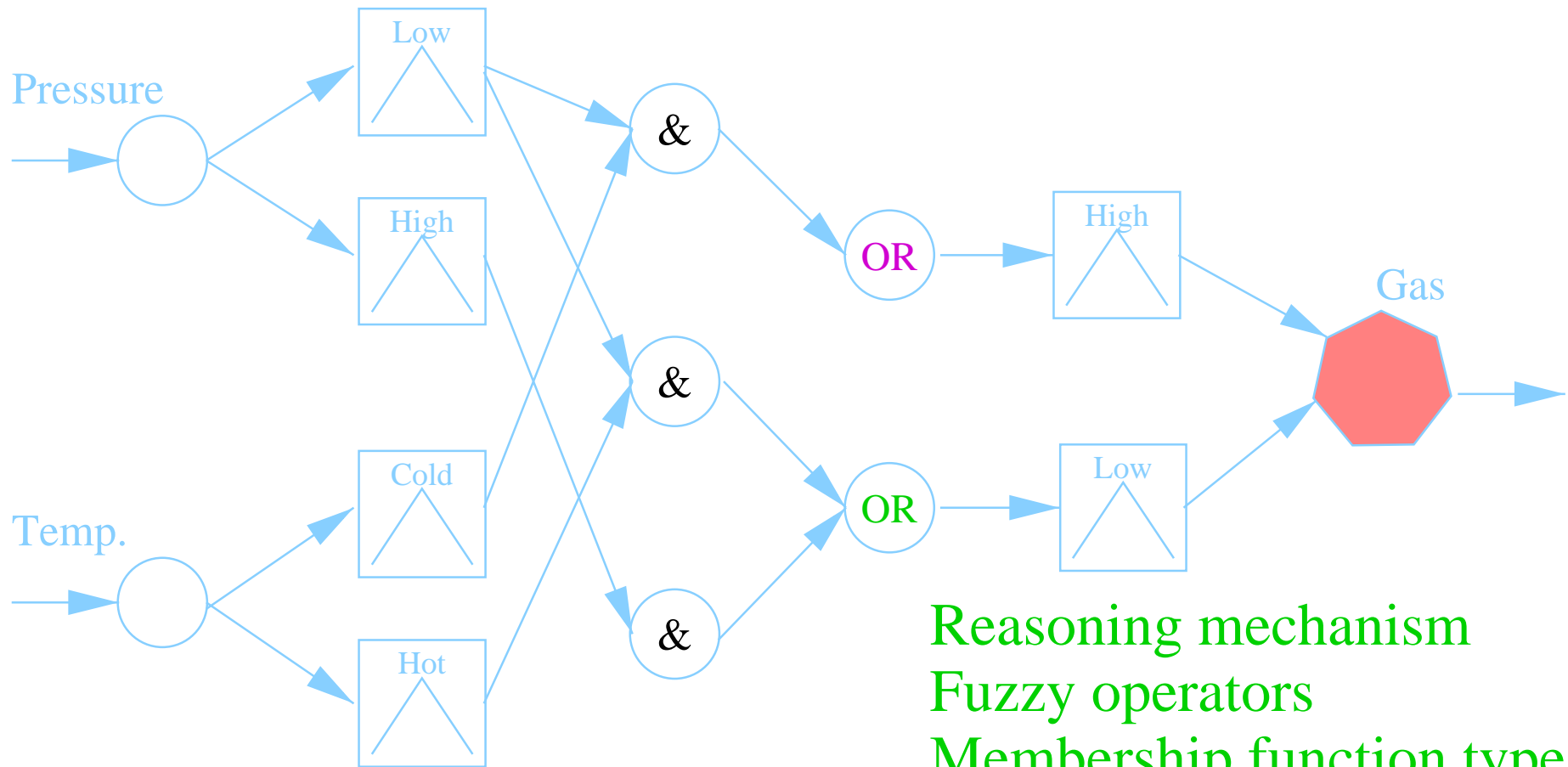
Connective parameters



Structural parameters



Logical parameters



Reasoning mechanism
Fuzzy operators
Membership function types
Defuzzification method

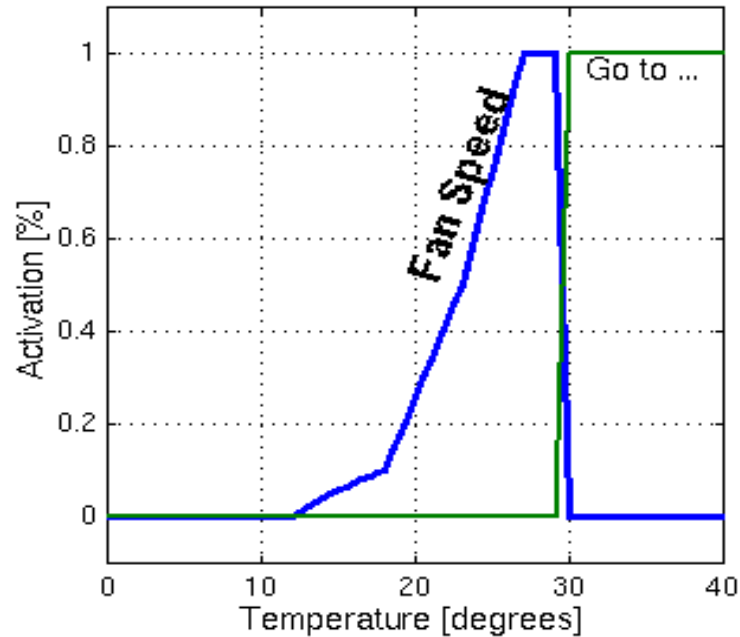
Parameters of a fuzzy system

Class	Parameters	Component
Logic	Reasoning mechanism	Inference engine
	Fuzzy operators	
	Membership function types	Fuzzi- and defuzzifier
	Defuzzification method	Defuzzifier
Structural	Relevant variables	Knowledge base
	Number of membership functions	
	Number of rules	
Connection	Antecedents of rules	Rulebase
	Consequents of rules	
	Rule weights	
Operational	Membership function values	Database

Dual external nature

Numeric

Linguistic



If Temperature is COOL then Ventilator is Off

If Temperature is WARM then Ventilator is Low

If Temperature is HOT then Ventilator is Medium

If Temperature is VERY-HOT then Ventilator is High

If Temperature is HELLISH then Ventilator is Off, and...

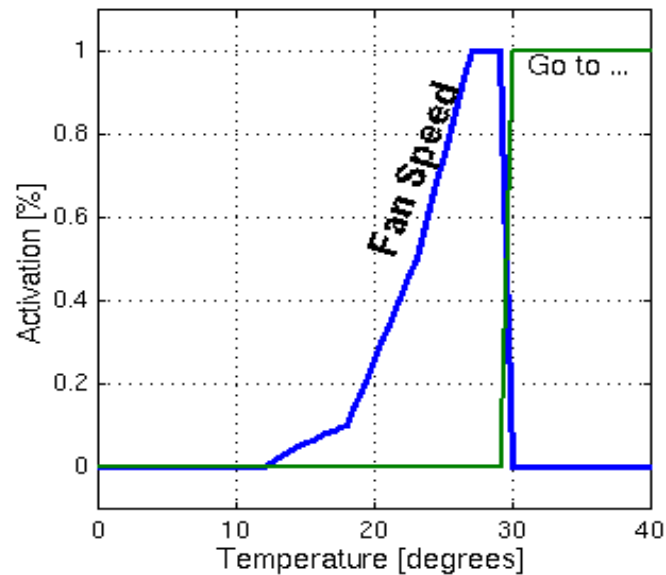
... Let's go to the lake!!!

Precision

Interpretability

Numeric issues

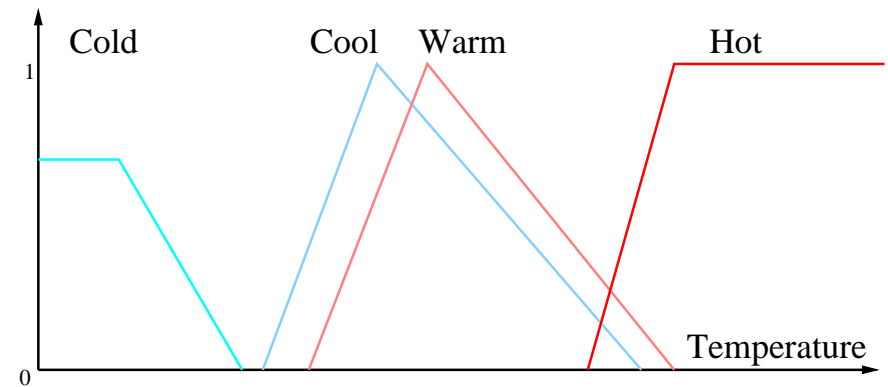
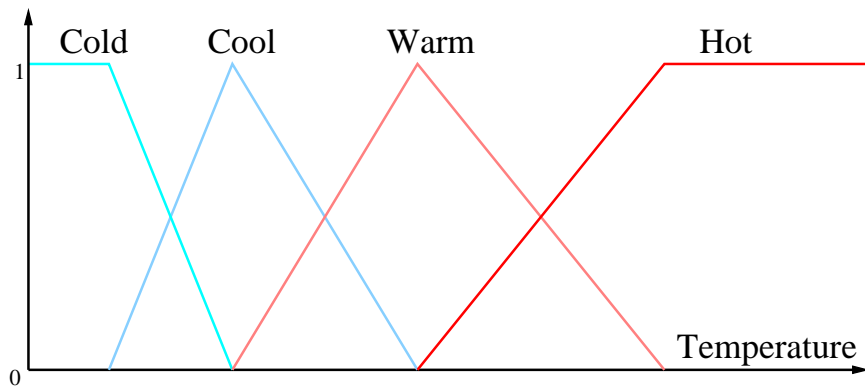
- Numeric mapping: Crisp inputs / Crisp outputs
- Nonlinear behavior, but linearity not excluded
- Universal approximator
- Uncertainty management: noise and low quality of data



Interpretability considerations: semantic criteria

Semantics: the study of meanings

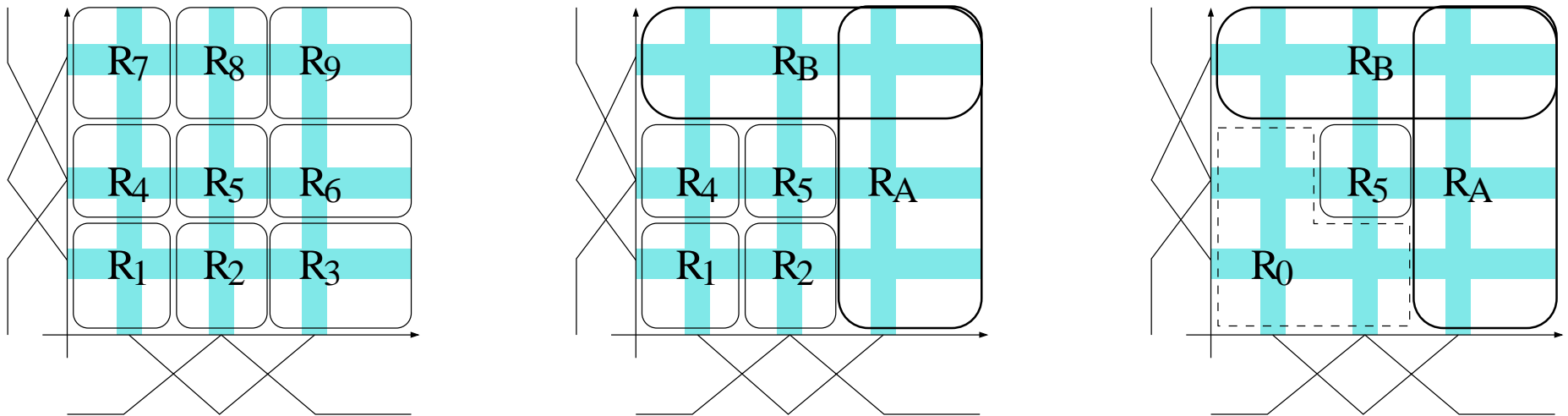
- Distinguishability: Each linguistic label has semantic meaning
- Number of elements: Compatible with human capabilities
- Coverage: Any element belongs to at least one fuzzy set
- Normalization: At least one element has unitary membership
- Complementarity: For each element, the sum of memberships is one



Interpretability considerations: syntactic criteria

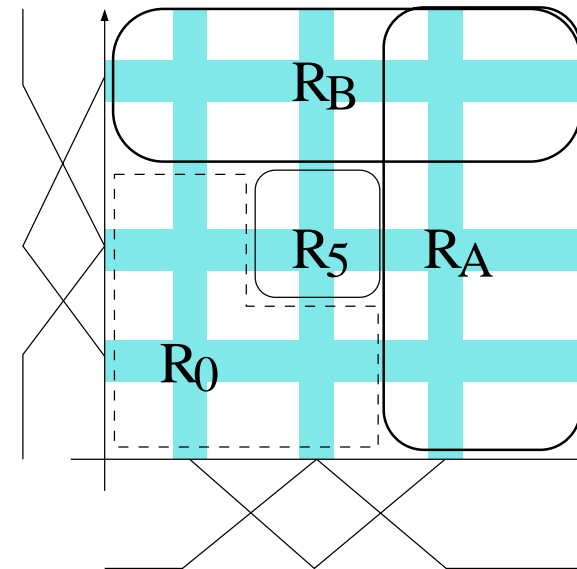
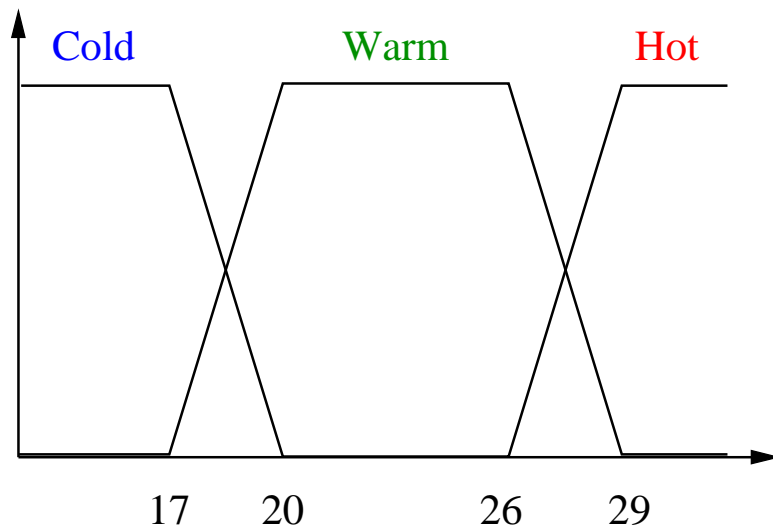
Syntax: the way in which linguistic elements are put together

- Completeness: for any input, at least one rule must fire
- Rule-base simplicity: Set of rules as small as possible
- Rule readability: small number of conditions in rule antecedents
- Consistency: rules firing simultaneously must have similar consequents



Strategies to satisfy interpretability criteria

- Linguistic labels shared by all rules
- Normal, orthogonal membership functions
- Don't care conditions
- Default rule



The general modeling problem

What do you know about the modeled system?

i.e. what is predefined and what looked for?

Search space

How do you search?

i.e. do a well suited and/or well known technique exists?

Search method

Have you preferences or restrictions to the model?

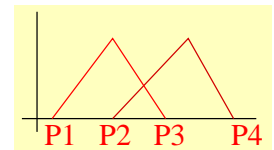
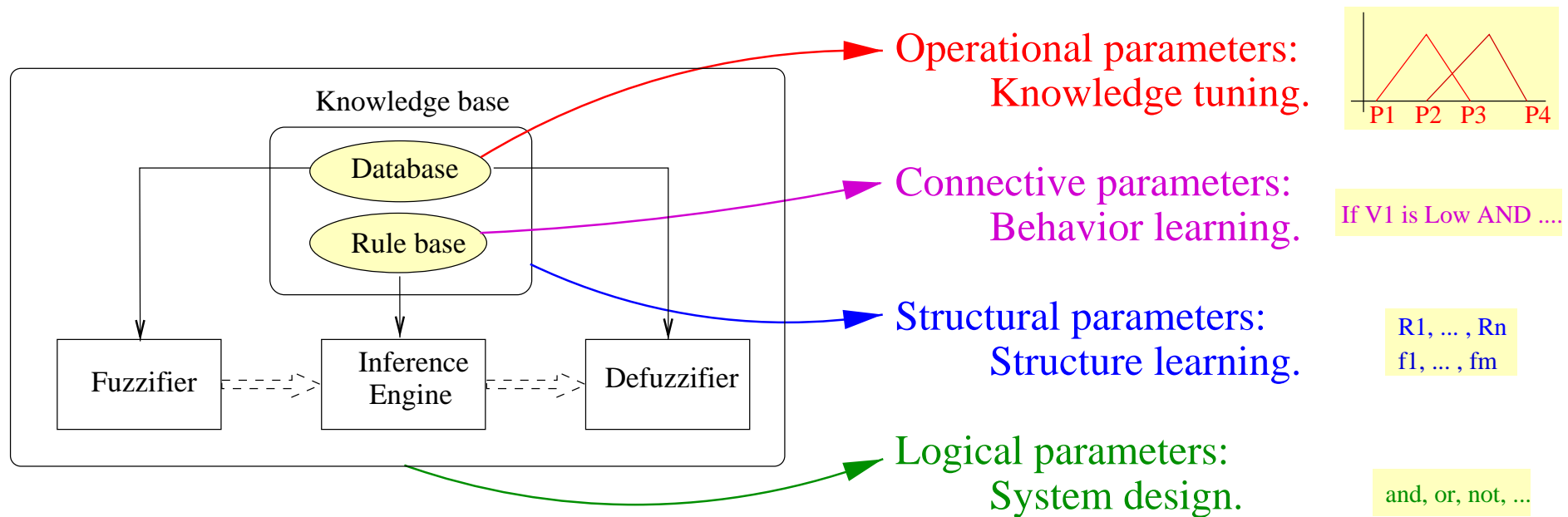
i.e. do issues like size, speed, or simplicity matter?

Constraints

Search space in fuzzy modeling

The number of parameters is too high to perform a full search, some parameter pre-definition is thus required.

According with the searched parameters we can have:



If V1 is Low AND

R1, ... , Rn
f1, ... , fm

and, or, not, ...

Search methods: Fuzzy modeling techniques

Knowledge engineering

"Classic" identification methods

Machine learning approaches

Neuro–fuzzy systems

Evolutionary fuzzy modeling techniques

Usual constraints in fuzzy modeling

What is the fuzzy system expected to do?

- Classification: Classification performance, quadratic error.
- Control: Dynamic response, adaptability, robustness, etc.
- Diagnostic: Overall performance, sensitivity, specificity
- Data mining: Completeness, complexity.

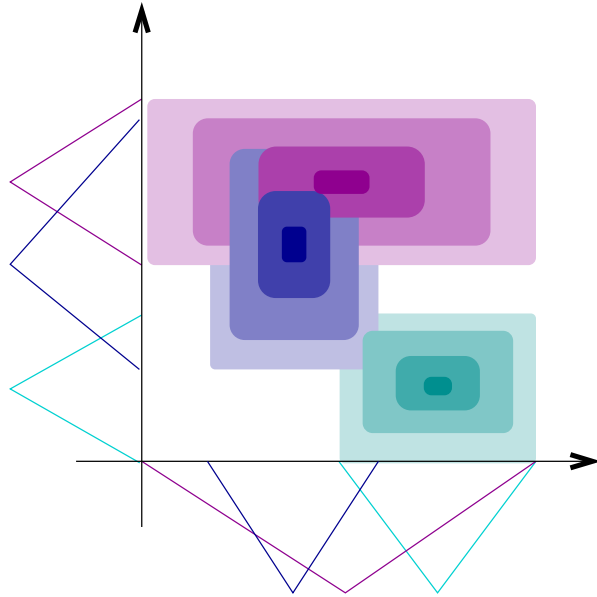
How is the system expected to do it?

- Speed: Real-time constraints, computing resources.
- Size: Available memory, computing platform.

Who is going to interact with the system?

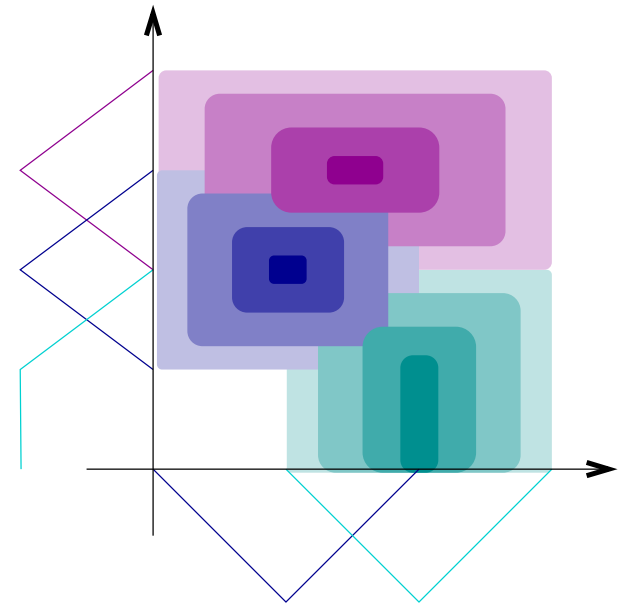
- Interpretability: Allowed complexity.
- Availability: Continuity of explanations (time to provide them)

Interpretability-related constraints

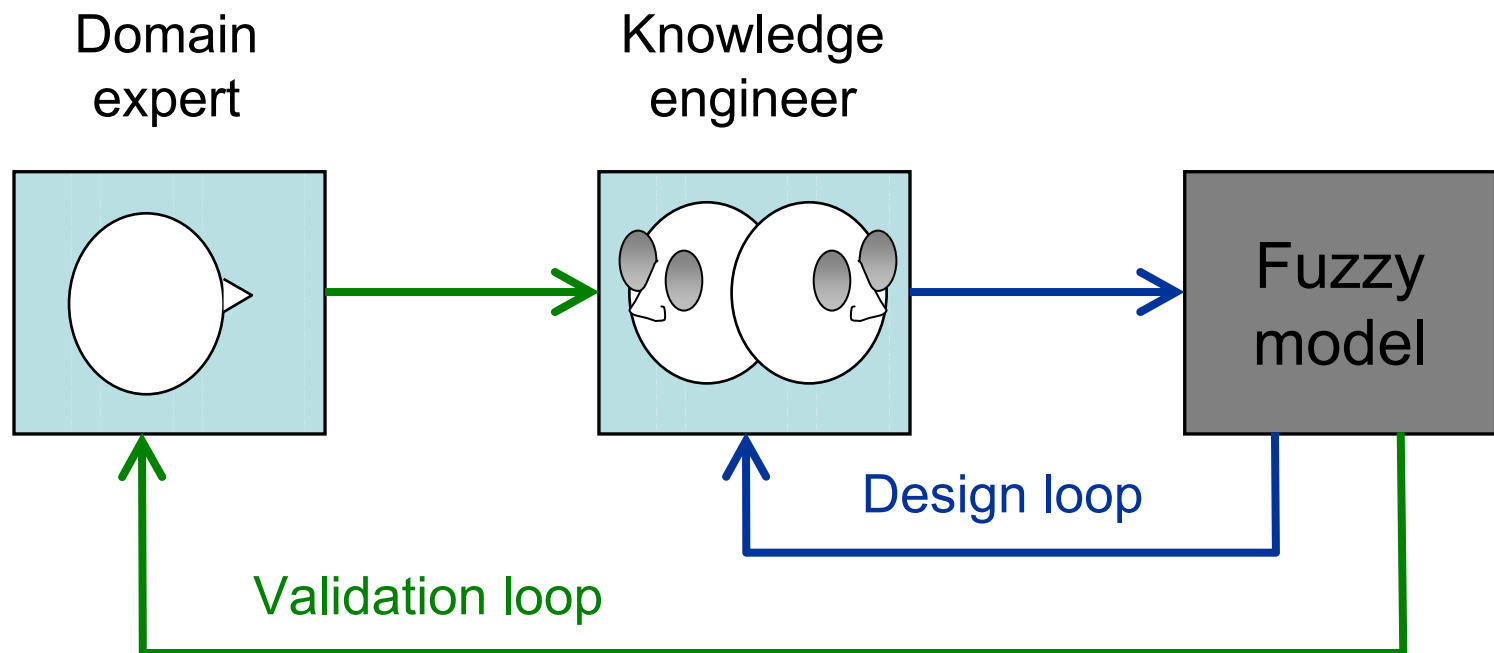


Rule-specific MFs are not allowed
all rules share the same MFs

Orthogonal MFs with well defined
null and unity membership

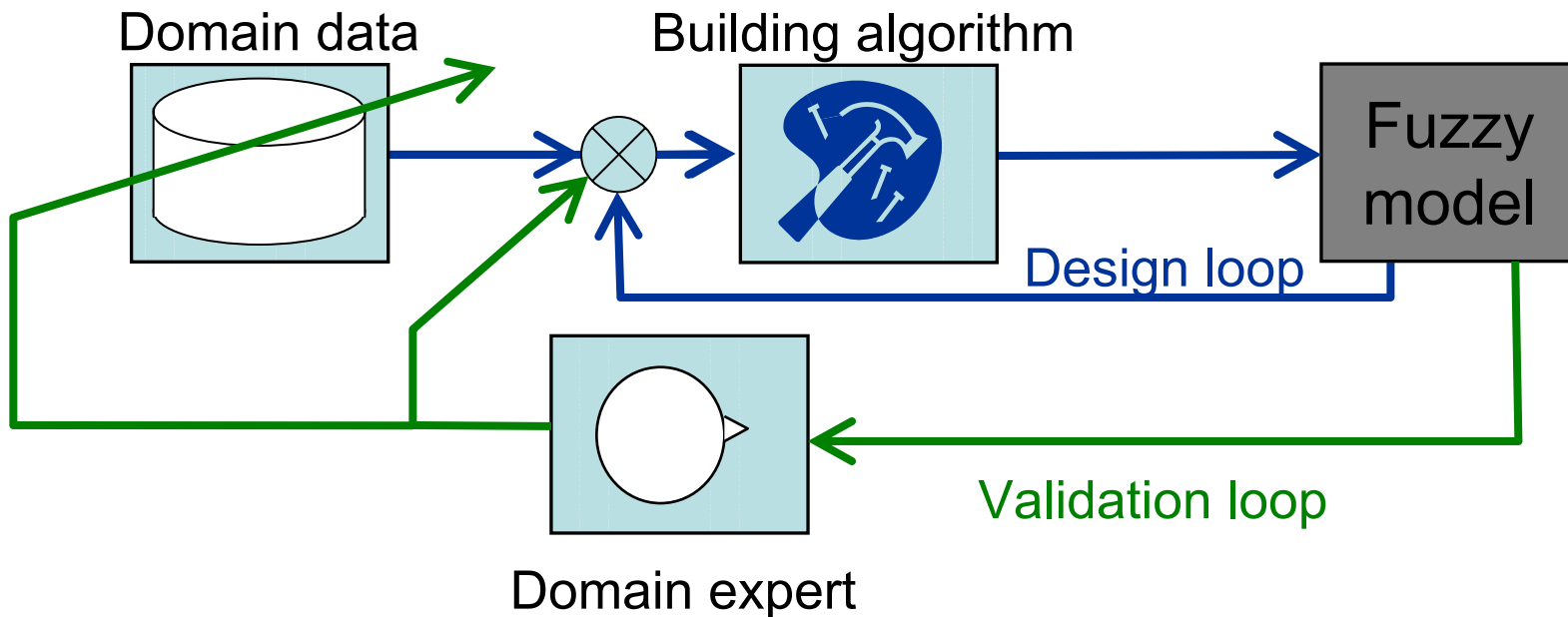


Fuzzy modeling: direct approach



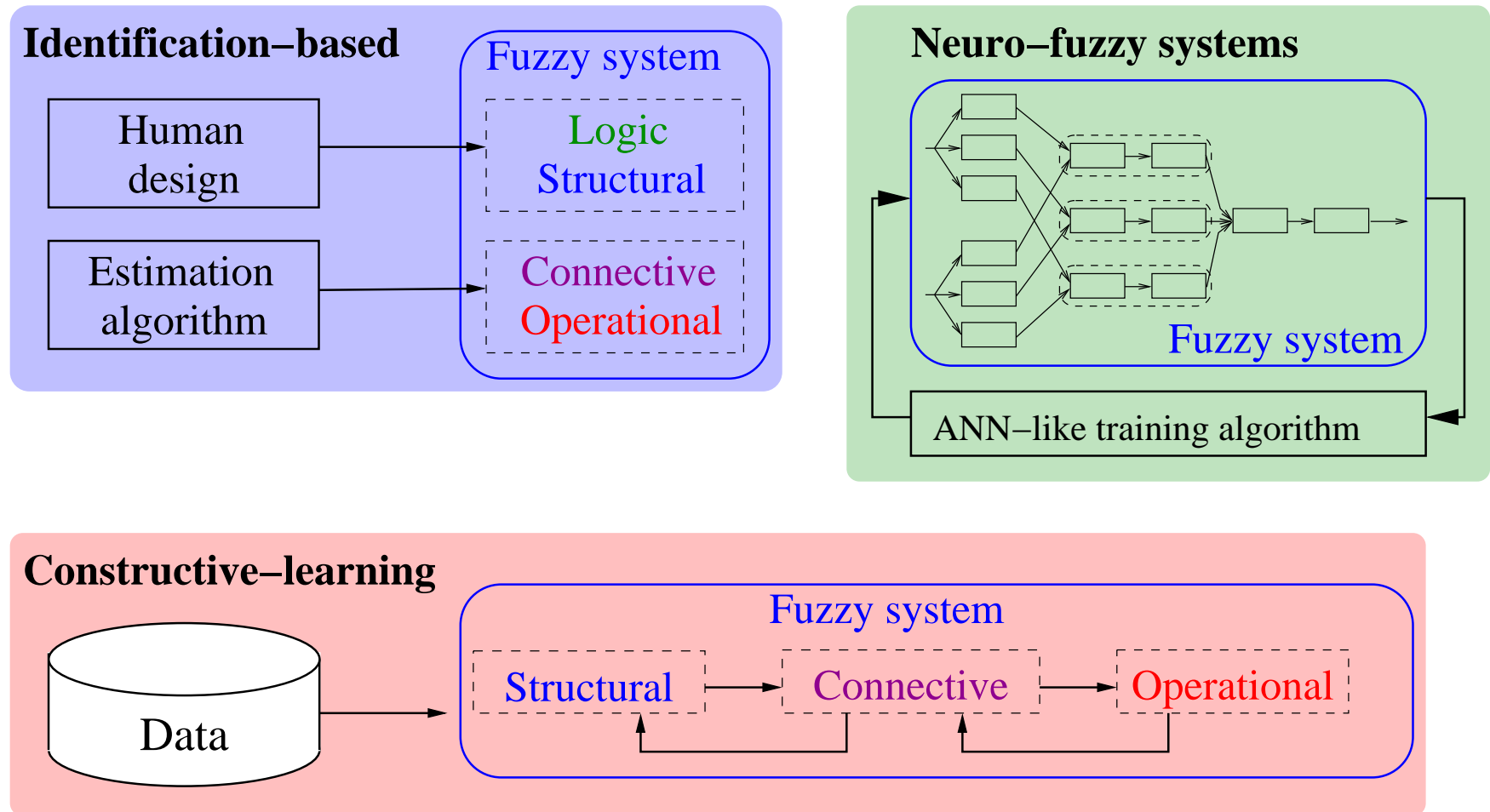
This approach is also called *knowledge engineering*

Fuzzy modeling: data-driven approaches

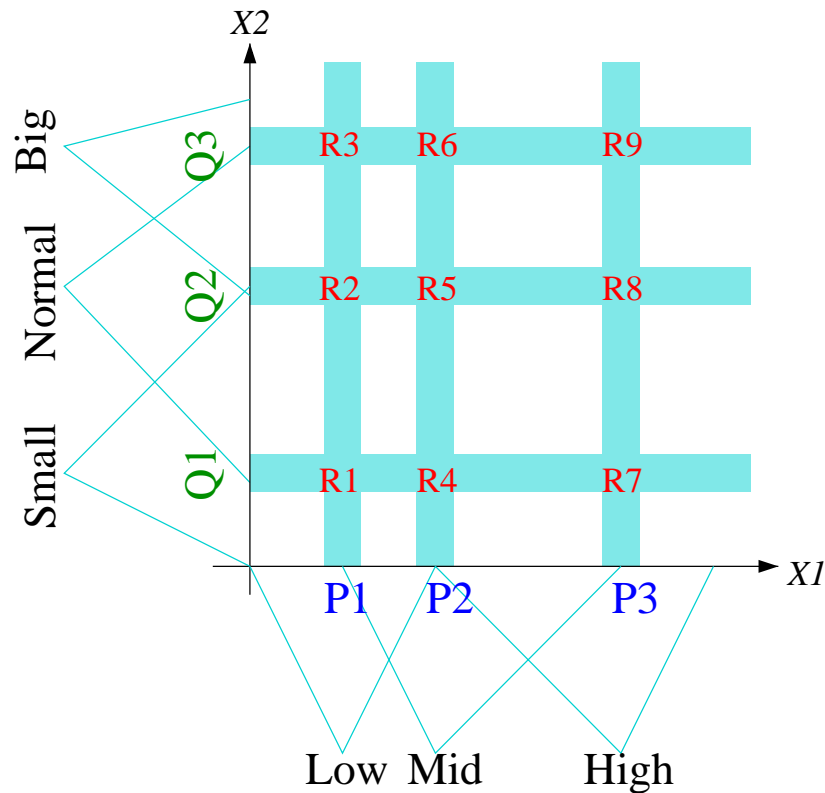


This approaches are also denominated
knowledge discovering

Fuzzy modeling: some data-driven approaches



Evolutionary knowledge tuning (database)



- Fixed rule base (completeness)

- Rules of type:

if $X_1 = \text{Low}$ and $X_2 = \text{Normal}$ then Output = C_i

- Knowledge is tuned by evolution,
which searches for membership function values

- Genome encodes values P, Q, and C

$$(3 * P + 3 * Q + 9 * C) * 5 \text{ bits} = 75 \text{ bits}$$

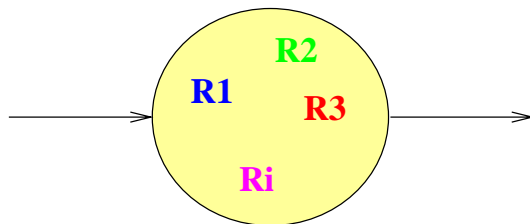
Evolutionary behavior learning (rule base)

- Number of rules explodes rapidly
- Two strategies for reducing this number:
 - Don't care conditions and default rule
- Evolution can be used to find a minimal (or fixed size) rule base,
- Three main approaches to evolutionary behavior learning

Michigan

Individual = One rule

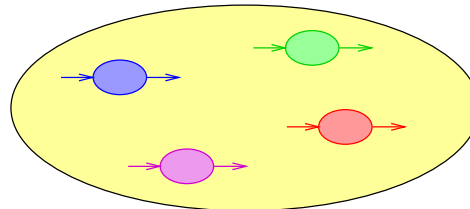
Population = Rule base
(i.e. fuzzy system)



Pittsburgh

Individual = Entire system
(rule base or knowledge base)

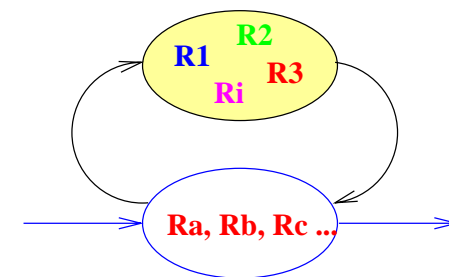
Population of systems



Iterative Rule Learning

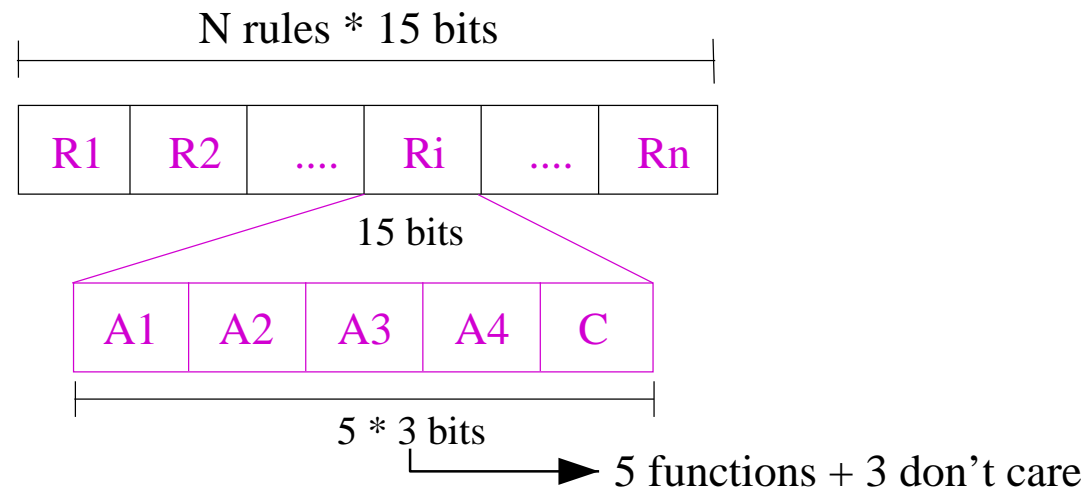
Evolution finds the best rule

Incremental construction of
the knowledge base



Evolutionary behavior learning: An example

- 4 input variables, 5 membership functions per variable
{Tiny, Small, Normal, Big, Huge}
- Space of 625 rules (1295 including don't care conditions)
IF V1 is Tiny AND ... AND V4 is Normal then Out = Huge
- Evolution searches for a subset of N rules (fixed by the designer)
- Genome encodes rules: Antecedents and consequents



Evolutionary knowledge base learning

(Knowledge base = rule base + database)

Parameter class	Modeling type	Usual quantity	Type of values	Fuzzy system attribute
Connective	Behavior learning	10 - 1000	Symbolic	Rule base
Operational	Knowledge tuning	10 - 1000	Real-valued	Database

Critical issues for applying evolution:

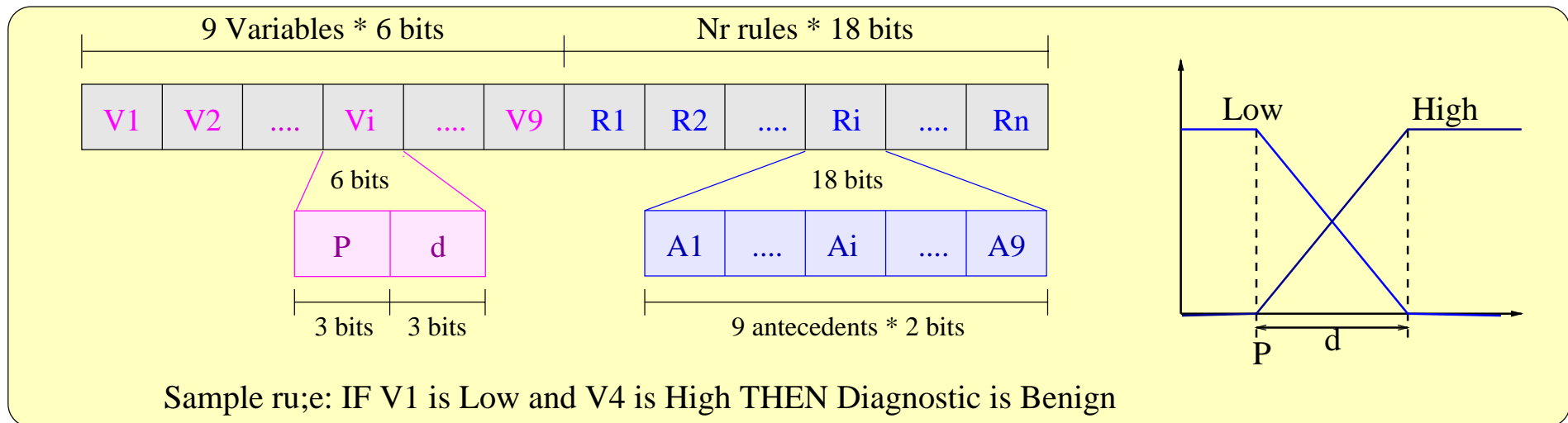
- Parameter representation
- Tight interdependency
- Size of the search space
- Computation requirements

Evolutionary knowledge base learning

- A basic approach: Single population, single evolution

Example: Breast cancer diagnosis problem (Peña and Sipper 99)

- 9 inputs, 1 output, 2 membership functions per variable
- A simple genetic algorithm searches for the knowledge base
- Genome encodes: Rule antecedents and membership function parameters

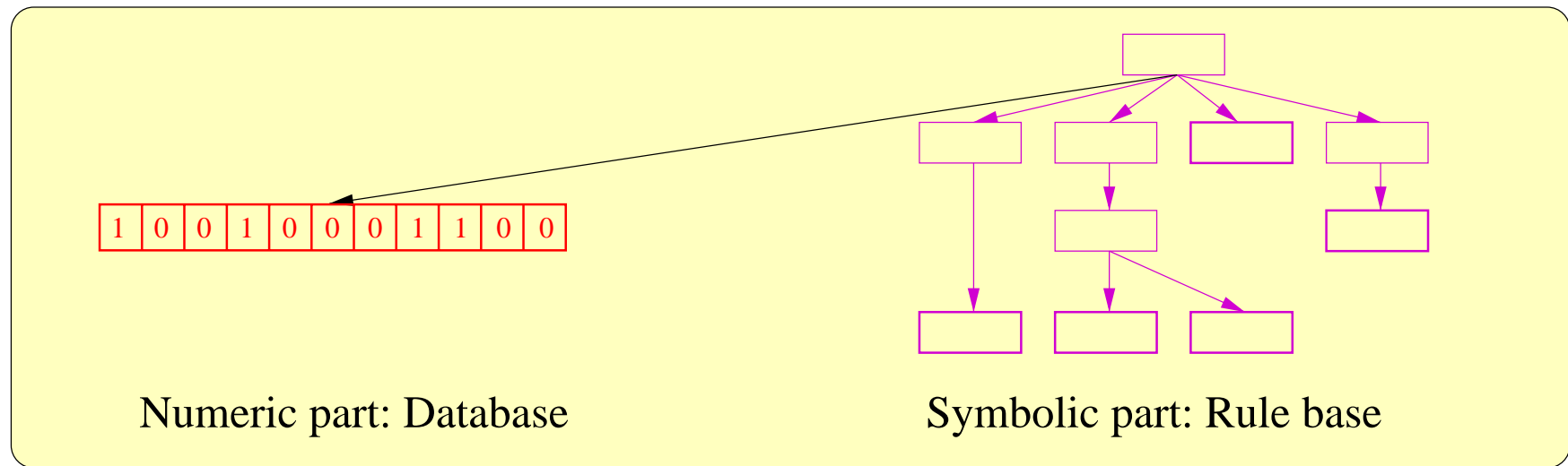


Evolutionary knowledge base learning

- A variation: Single population, double evolution

Example: Evolving fuzzy rule based classifiers with GA-P (García et al. 99)

- Genome encodes: Complete rule base and membership function parameters
- A simple genetic algorithm searches for the database
- The rule base is evolved using genetic programming

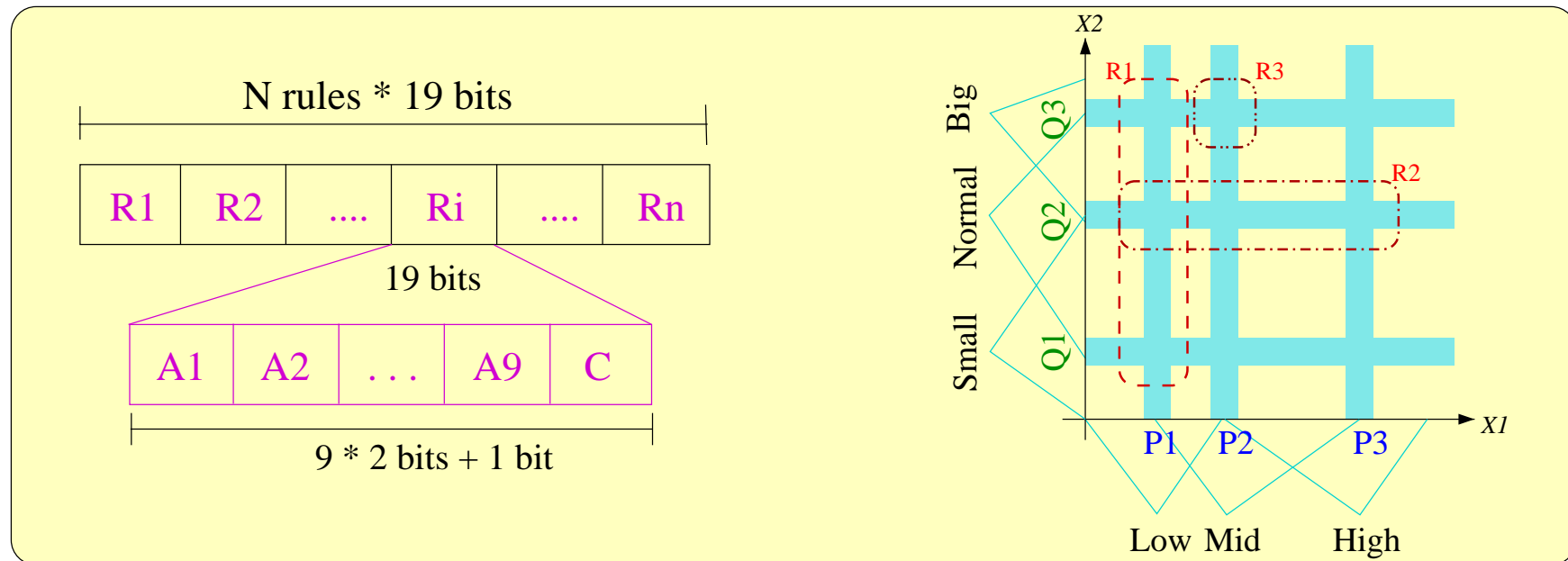


Evolutionary knowledge base learning

- Hybrid learning: Evolved rule base, learned database

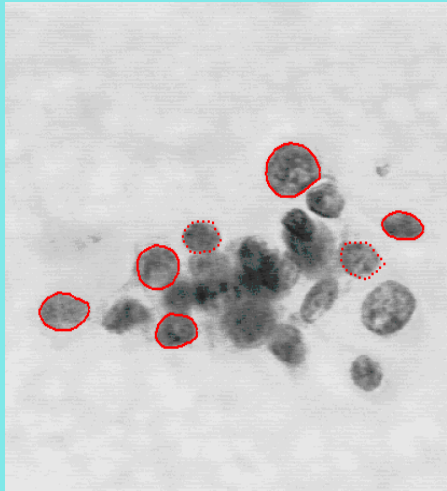
Example: Breast cancer diagnosis (J.-F. Philagor, student project SPG, 1999)

- Evolution searches for a fixed-size rule base
Genome encodes rules: Antecedents and consequent
- Database is tuned using a neuro-fuzzy approach
A fuzzy self-organizing map searches for P and Q values



The Wisconsin Breast Cancer Database

The test



The features

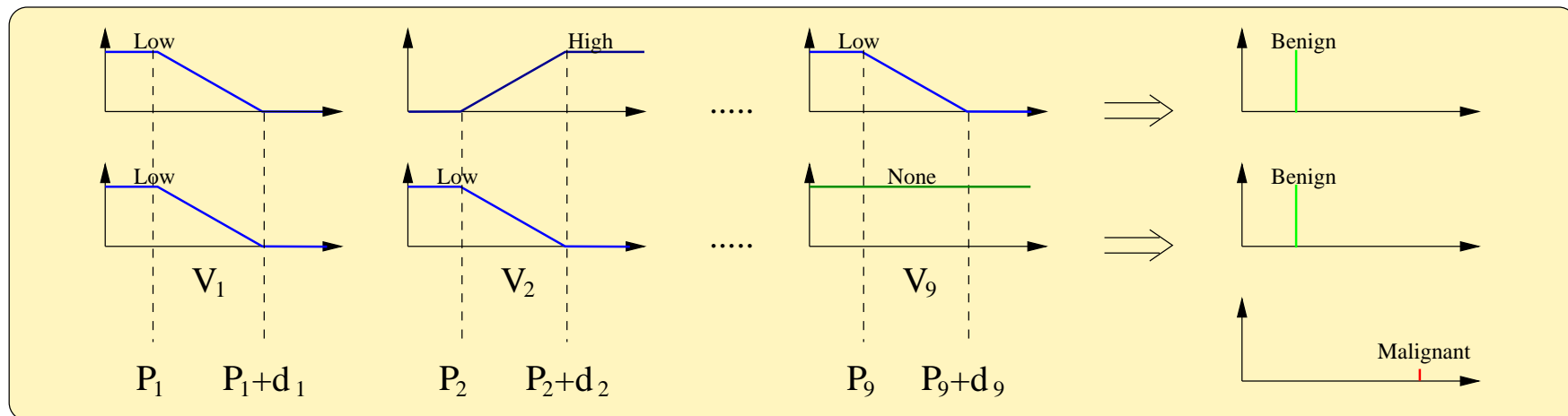
1. Clump Thickness (v_1)
2. Uniformity of Cell Size (v_2)
3. Uniformity of Cell Shape (v_3)
4. Marginal Adhesion (v_4)
5. Single Epithelial Cell Size (v_5)
6. Bare Nuclei (v_6)
7. Bland Chromatin (v_7)
8. Normal Nucleoli (v_8)
9. Mitosis (v_9)

The database

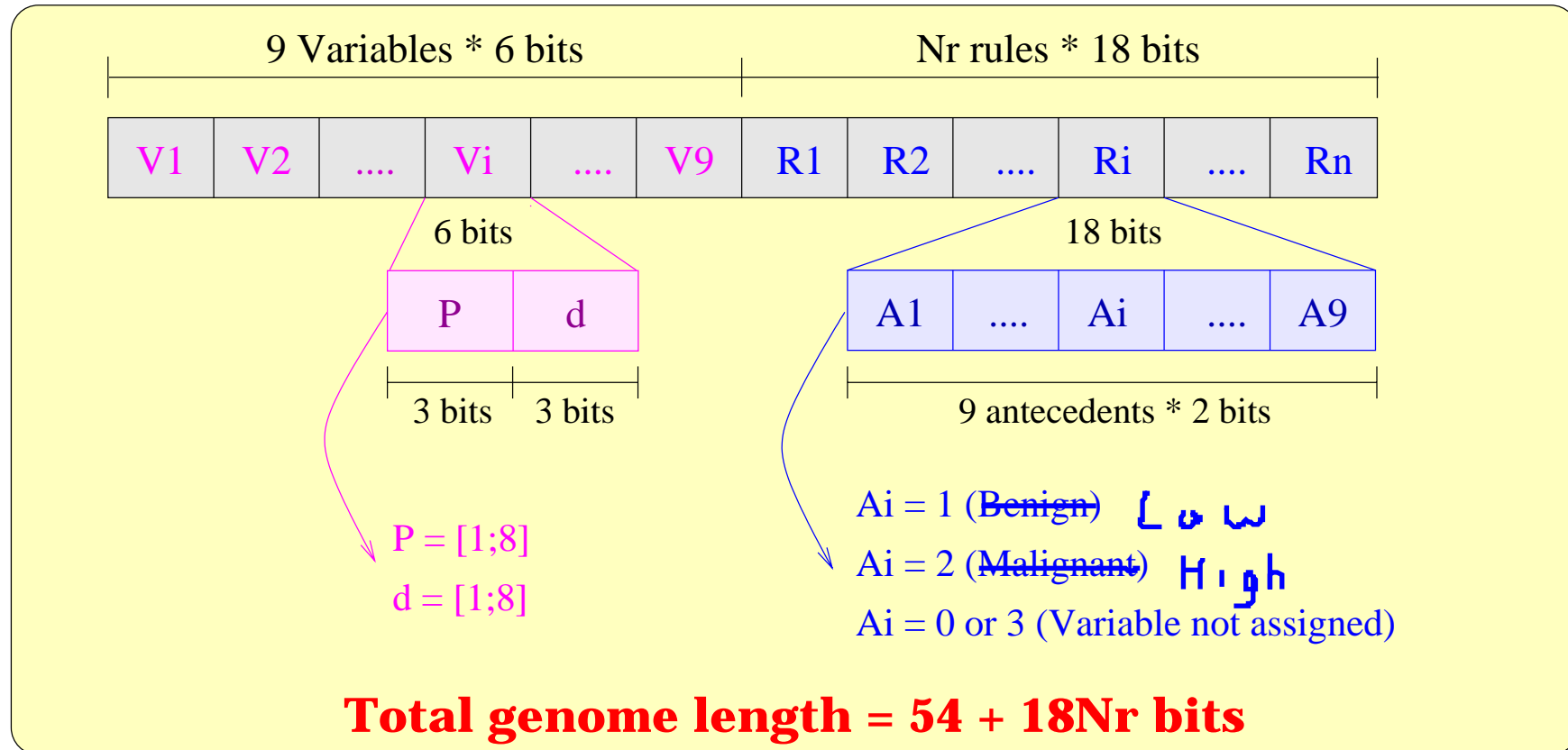
<i>case</i>	v_1	v_2	v_3	...	v_9	<i>diagnostic</i>
1	5	1	1	...	1	<i>Benign</i>
2	5	4	4	...	1	<i>Benign</i>
:	:	:	:	:	:	:
683	4	8	8	...	1	<i>Malignant</i>

Proposed Fuzzy System

R1: if (V1 is Low) and (V2 is High) and ... and (V9 is Low) then (output is Benign)
R2: if (V1 is Low) and (V2 is Low) and ... and (V9 is None) then (output is Benign)
⋮
⋮
⋮
⋮
⋮
else (Output is Malignant)



Genome encoding



Fitness function

$$F = F_c + a * F_v + b * F_e$$

F_c : Classification performance,
the most important performance measure

F_v : Number of variables
measures the interpretability

F_e : Quadratic error
selection pressure to fine tune parameters

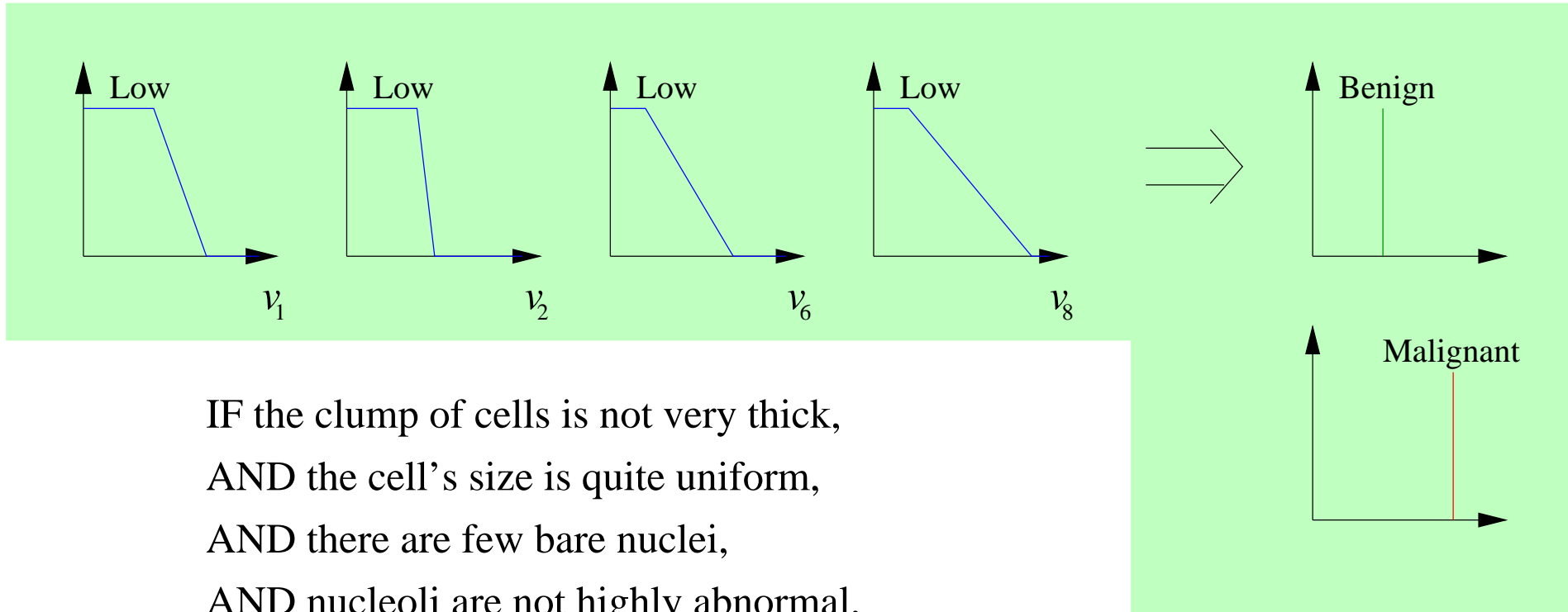
Results: Classification performance (Number of variables)

Rules	Setiono (96)	Setiono Liu (96)	Taha Ghosh (97)	Peña Sipper (98)	This work (99)
1	95.42% (2)			96.35% (3)	97.07% (4)
2				96.65% (7)	97.36% (3)
3	97.14% (4)	97.21% (4)			97.80% (4.7)
4					97.80% (4.8)
5			96.19% (1.8)		97.51% (3.4)

Learned Boolean rules

Evolved fuzzy rules

The best single-rule system



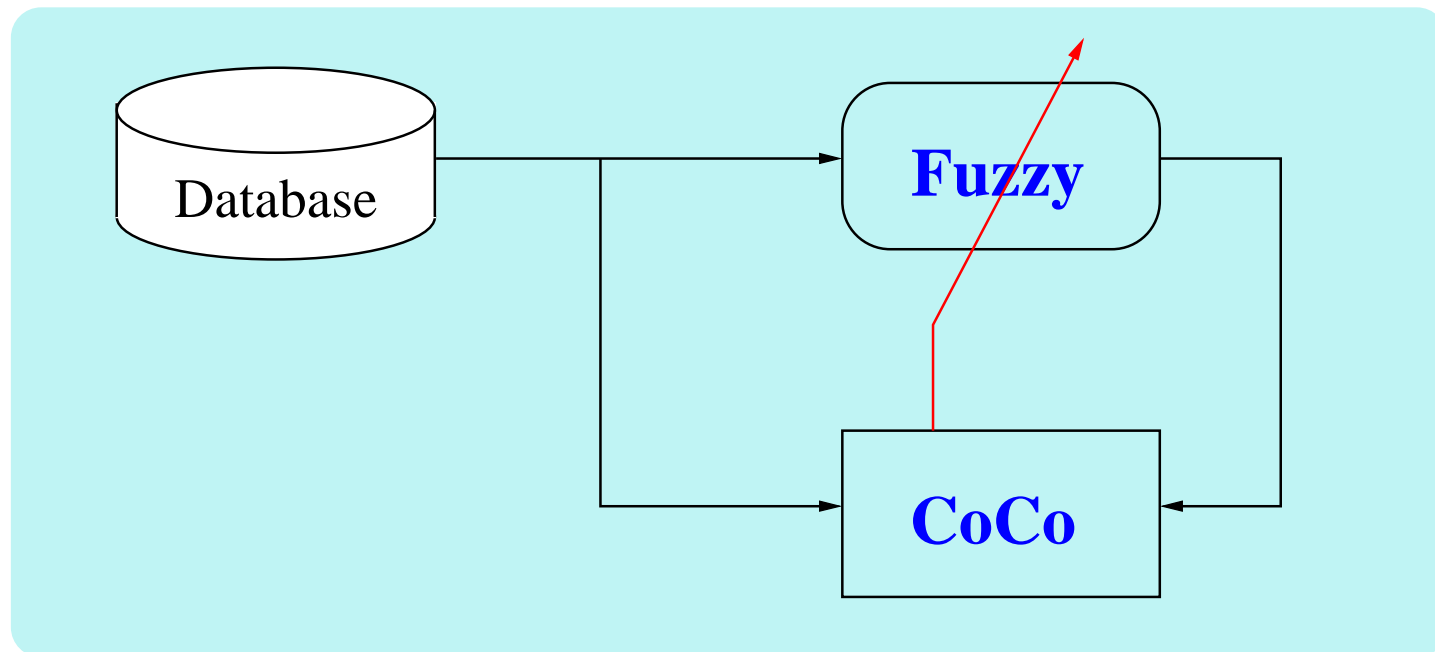
IF the clump of cells is not very thick,
AND the cell's size is quite uniform,
AND there are few bare nuclei,
AND nucleoli are not highly abnormal,

THEN the case is benign;

ELSE the case is malignant.

Proposed approach: two elements

- ① A system model: **Fuzzy systems**
- ② A building algorithm: **Cooperative coevolution**



Fuzzy modeling: a coevolvable problem

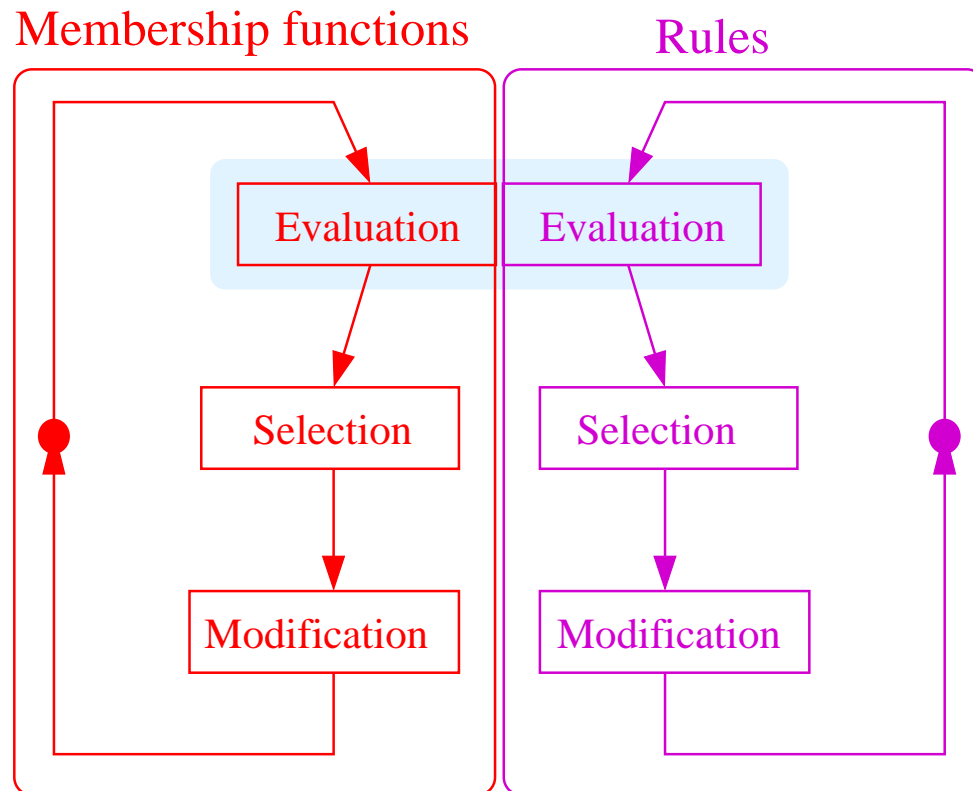
Parameter class	Modeling type	Usual number	Type of values
Logical	System design	3 – 10	
Structural (size)	Structure learning	5 – 20	Integer
Connective (rules)	Behavior learning	10 – 1000	Symbolic
Operational (labels)	Knowledge tuning	10 – 1000	Real-valued

The required solutions can be very complex, they can be decomposed in distinct components, represented by different types of values, and which are very interdependent.

These features render fuzzy modeling an adequate target for COOPERATIVE COEVOLUTION

Fuzzy CoCo:

A cooperative coevolutionary approach to fuzzy modeling



Two evolutionary algorithms searching for:

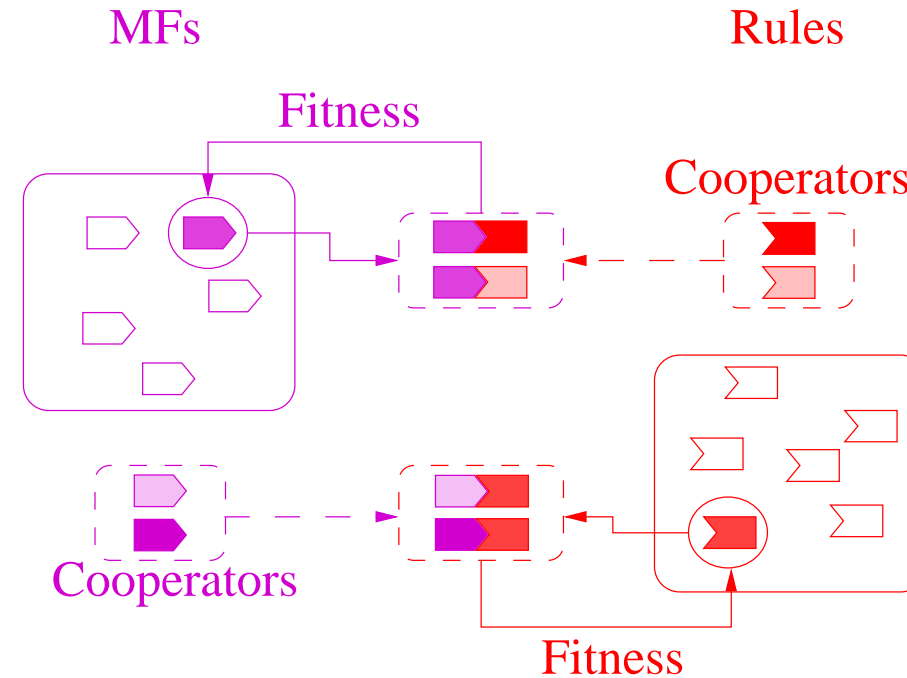
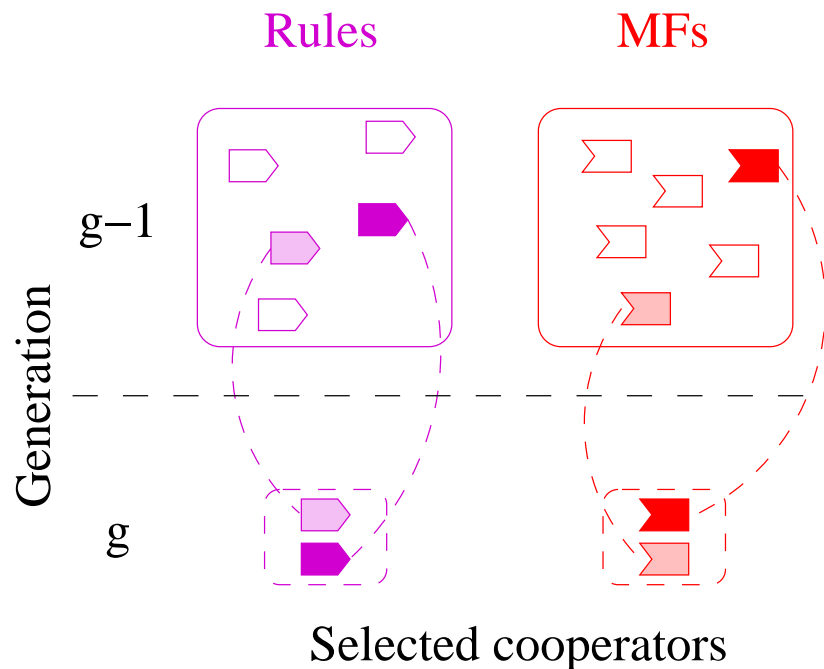
membership functions
and rules.

Advantages:

- Divide-and-conquer strategy
- Better search power
- Lesser computational cost
- More-flexible setup

Fitness evaluation in Fuzzy CoCo

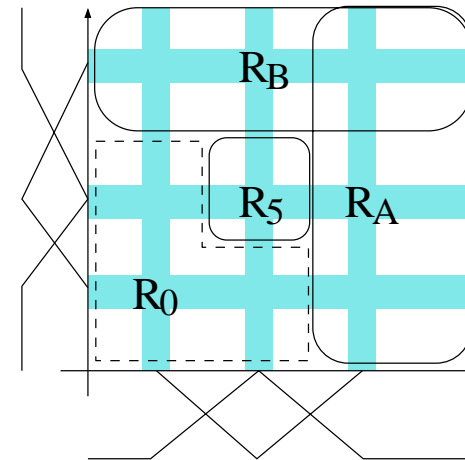
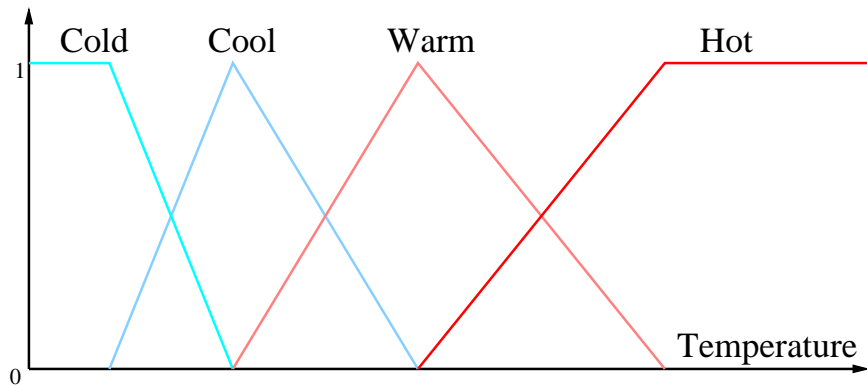
1. Cooperators for generation g are selected from generation $g-1$ both fitness-dependent and randomly



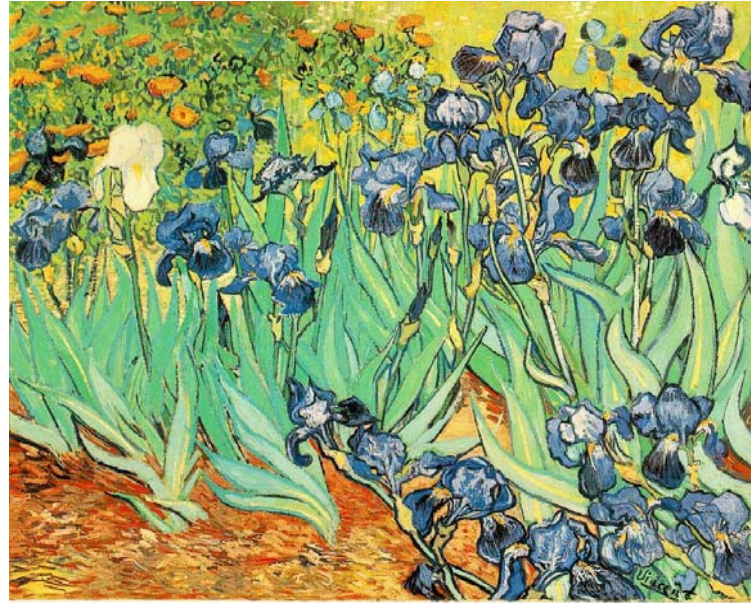
2. Individuals are combined with cooperators to form fuzzy systems.
3. These fuzzy systems are evaluated, and individual fitness is then calculated.

Interpretability strategies in Fuzzy CoCo

- Shared membership functions: reinforced by the existence of a separate species
- Normal, orthogonal membership functions: constrained representation
- Don't care conditions: encourage shorter rules
- Default rule: guarantees complete coverage of the input space
- Linguistic fitness: when used, increases selective pressure for interpretability



Fisher's Iris Data



The variables

Features

- (1) SL Sepal length
- (2) SW Sepal width
- (3) PL Petal length
- (4) PW Petal width

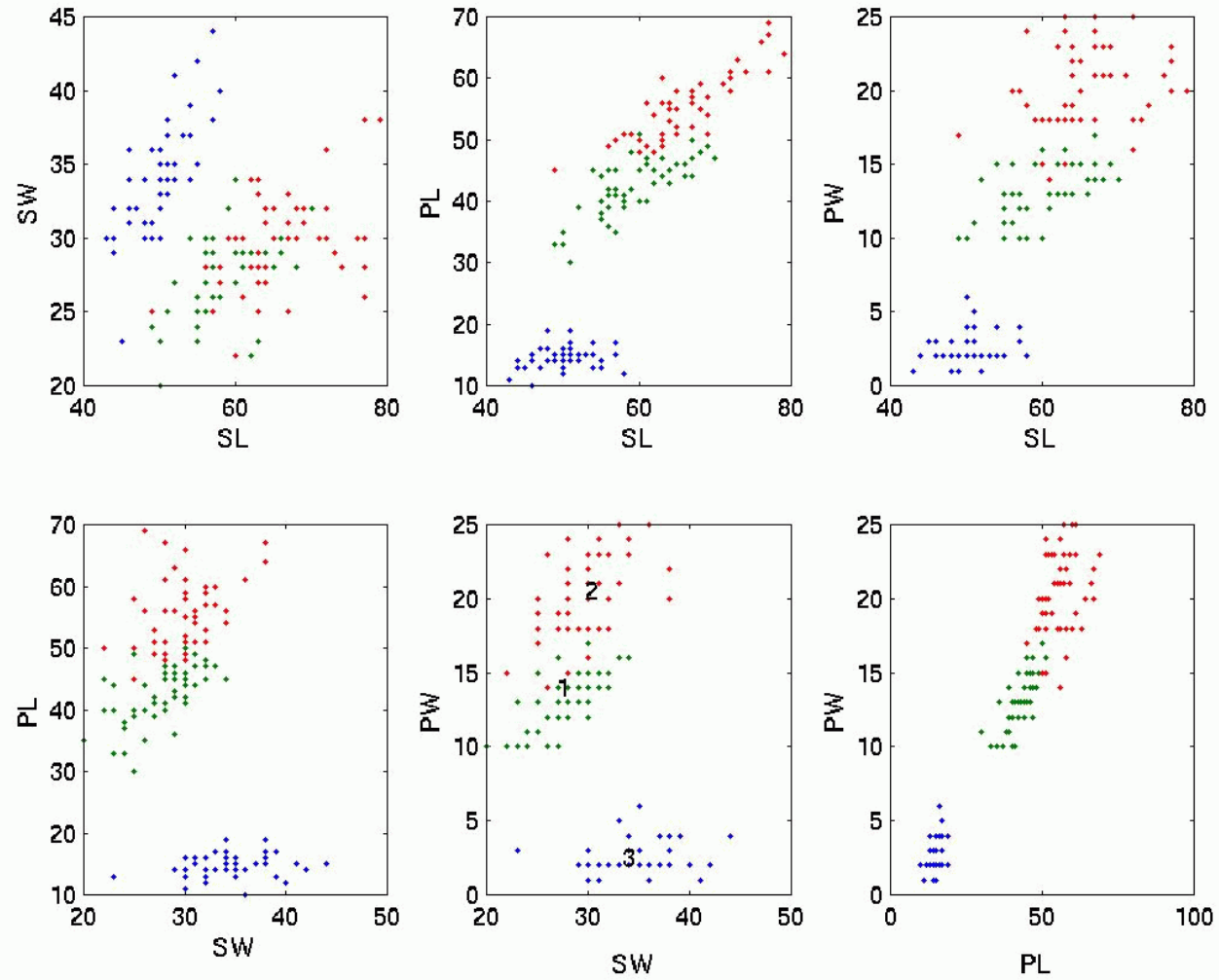
Classes

- (1) setosa
- (2) versicolor
- (3) virginica

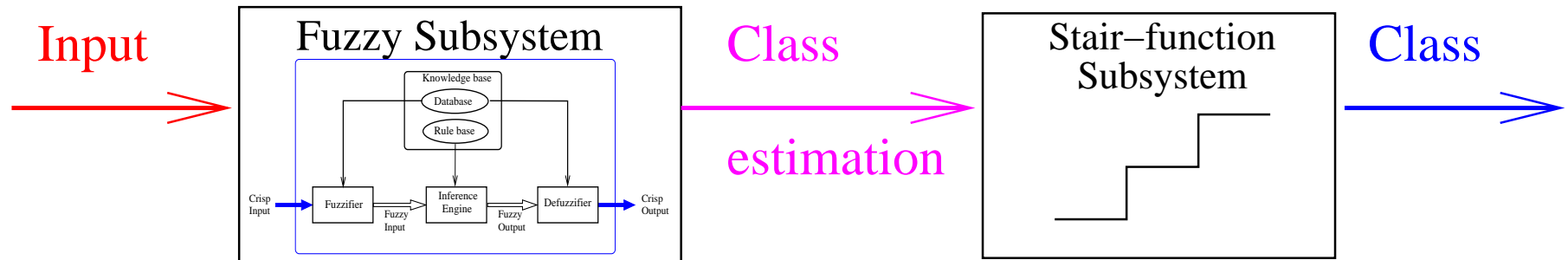
The database

Case	SL	SW	PL	PW	Class
1	5.1	3.5	1.4	0.2	Setosa
2	4.9	3.0	1.4	0.2	Setosa
150	5.9	3.0	5.1	1.8	Virginica

Iris: Variable analysis



Iris proposed solution: Controller-type

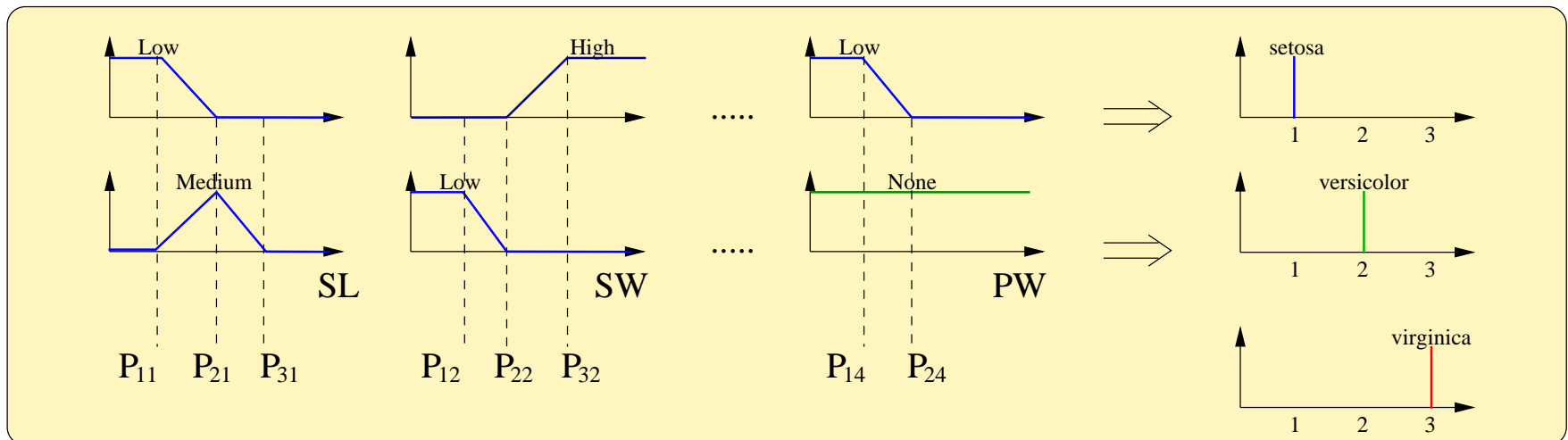


The fuzzy subsystem estimates a continuous "class" value

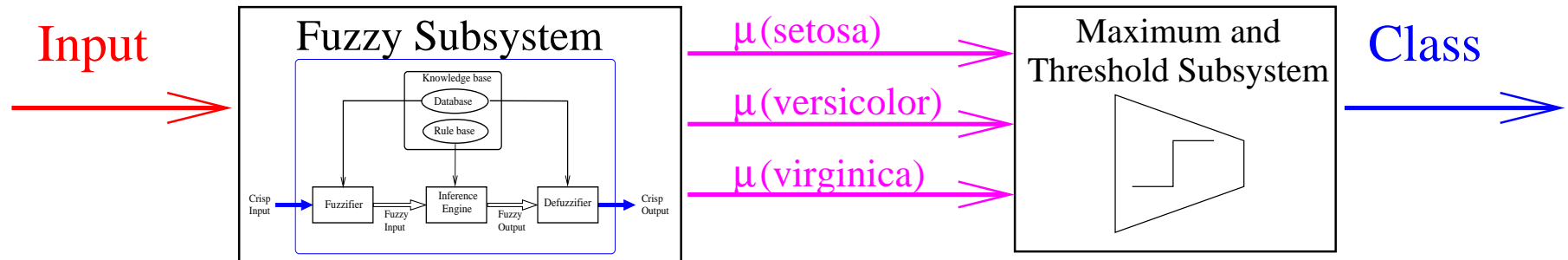
The selection unit approximates it to the nearest class

Iris controller–type: Proposed Fuzzy System

R1: if (SL is A11) and (SW is A12) and (PL is A13) and (PW is A14) then (output is Class1)
R2: if (SL is A21) and (SW is A22) and (PL is A23) and (PW is A24) then (output is Class2)
⋮ ⋮ ⋮ ⋮ ⋮
Rn: if (SL is An1) and (SW is An2) and (PL is An3) and (PW is An4) then (output is Classn)
else (Output is Class0)



Iris proposed solution: Classifier-type

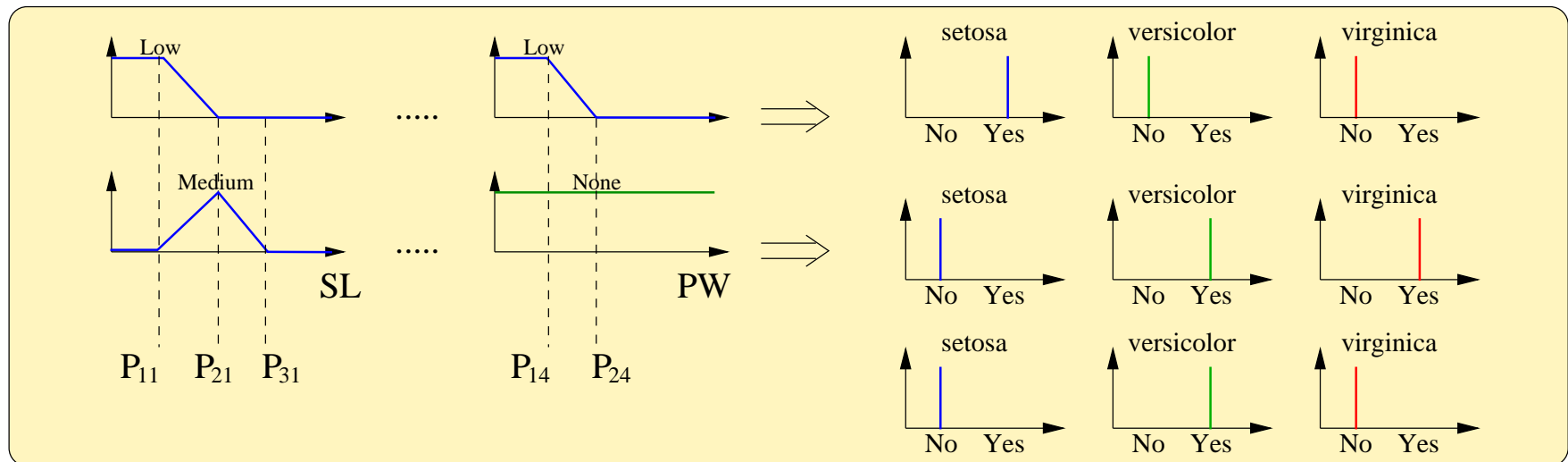


The fuzzy subsystem estimates a continuous membership value for each class

The selection unit chooses the most active class

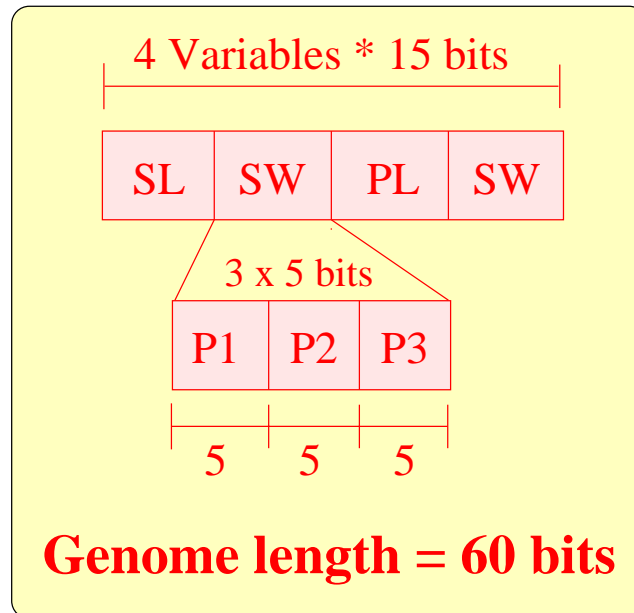
Iris classifier–type: Proposed Fuzzy System

R1: if (SL is A11) and ... and (PW is A14) then (setosa is Yes),(versicolor is No),(virginica is No)
 R2: if (SL is A21) and ... and (PW is A24) then (setosa is No),(versicolor is Yes),(virginica is Yes)
 ⋮ ⋮ ⋮ ⋮ ⋮
 Rn: if (SL is An1) and ... and (PW is An4) then (setosa is No),(versicolor is No),(virginica is Yes)
 else (setosa is No),(versicolor is Yes),(virginica is No)

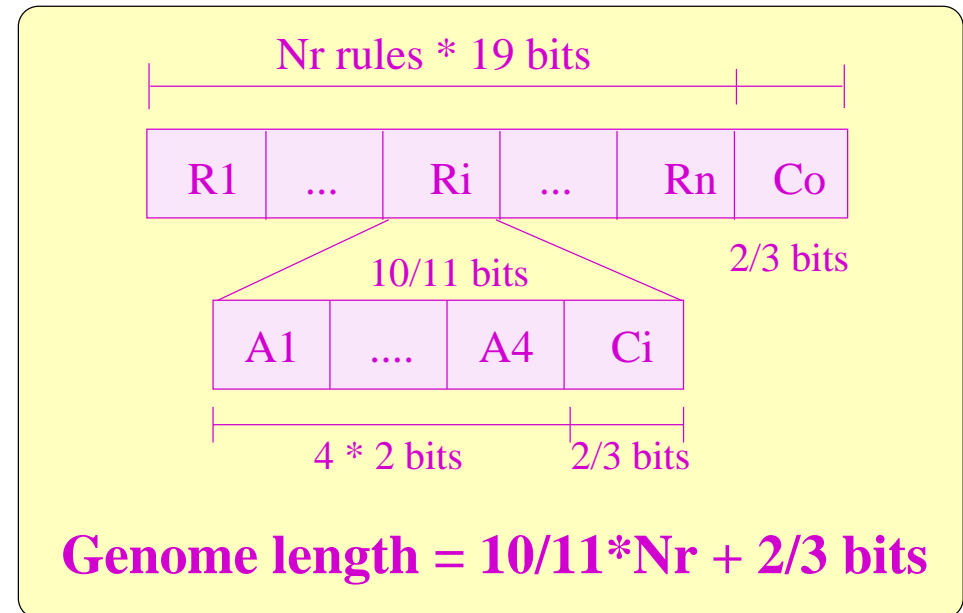


Iris: the genomes

Membership functions



Rules (Controller/Classifier)



Iris: Fuzzy CoCo set-up

1. Fitness function

$$F = \begin{cases} F_c * F_m^b \\ (F_c + a * F_v) * F_m^b \end{cases}$$

F_c : Classification performance,
the most important

F_m : $1 - \text{mse}$ (mean square error)
encourages not-so-bad errors

F_v : Number of variables
measures the interpretability

2. Fuzzy CoCo parameters

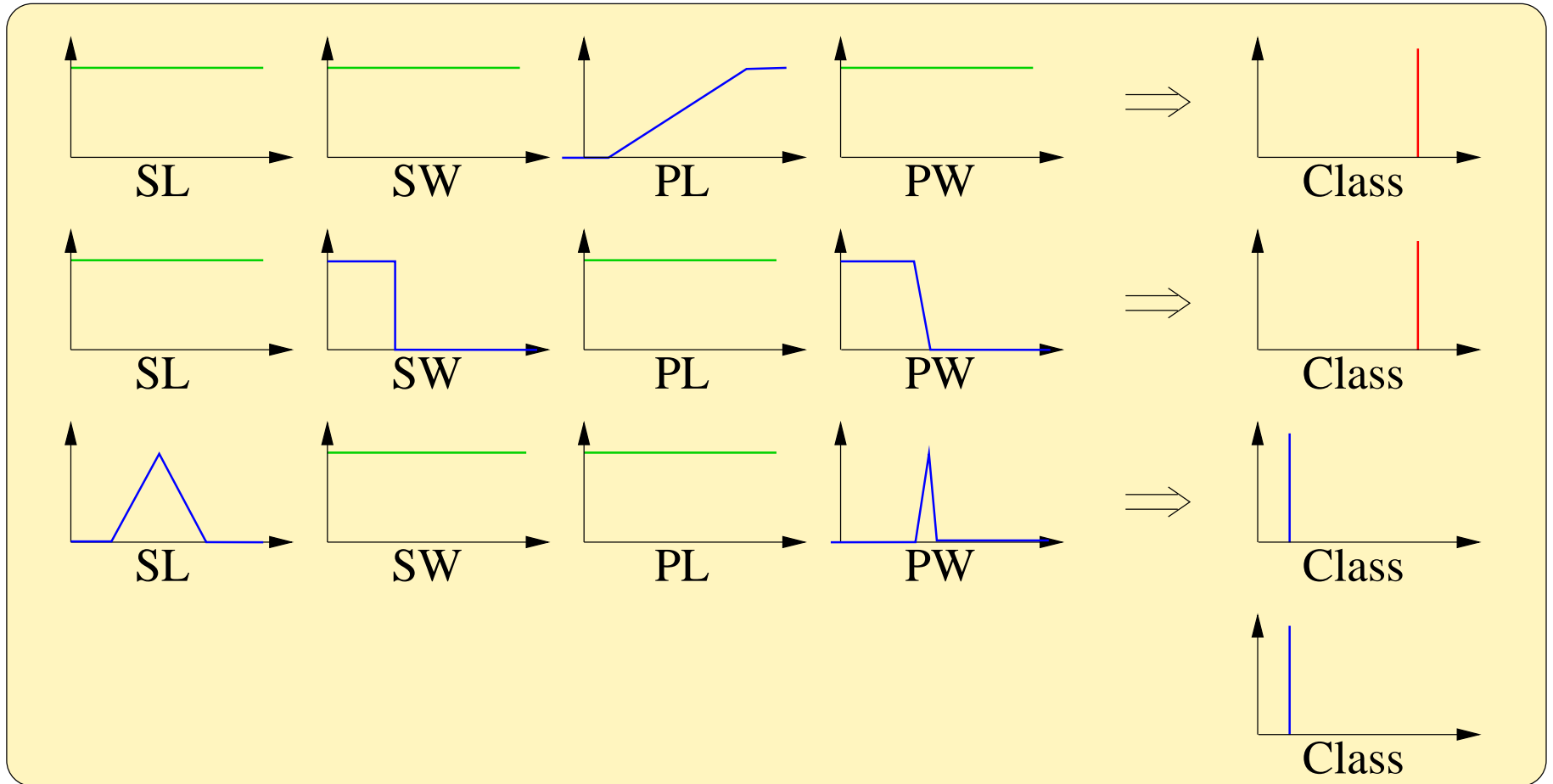
Population size	{60, 70}
Maximum generations	$500 + 100 * N_r$
Crossover probability	1
Mutation probability	{0.02, 0.05, 0.1}
Elitism rate	{0.1, 0.2}
"Fit" cooperators	1
Random cooperators	{1, 2}

Iris results: classification (average rule)

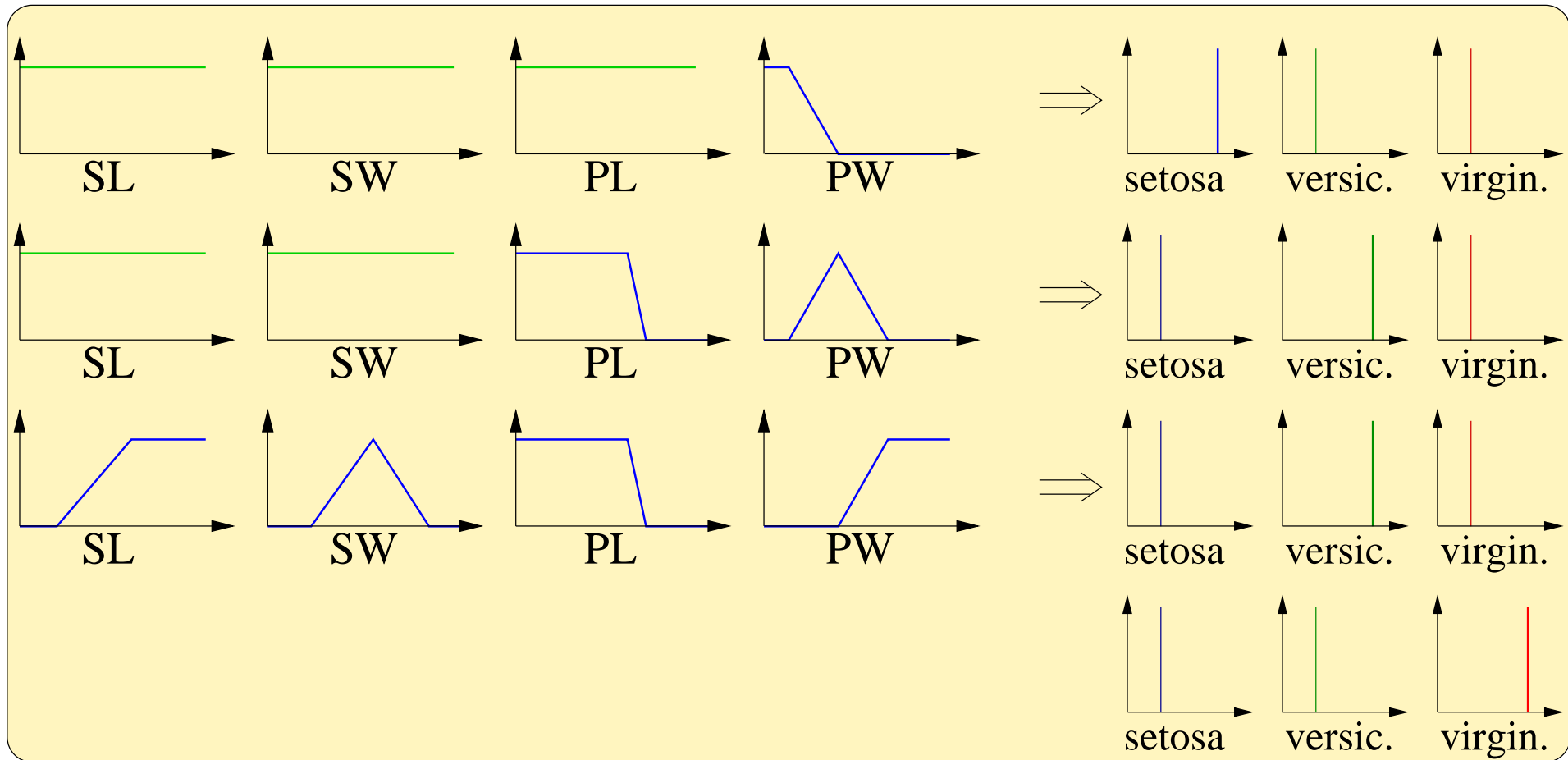
Controller	Rules	Simple GA Shi et al (1999)	FuGeNeSys Russo (1998)	Fuzzy CoCo ICANNGA'01
	2			99.33% (2)
	3			100 % (1.7)
	4	98 % (2.6)		100 % (2.5)
	5		100 % (3.3)	

Classifier	Rules	Constructive Learning Methods		Neurofuzzy Hung (99)	Fuzzy CoCo ICANNGA'01
		Hong (00)	Wu (99)		
	2				98 % (1.5)
	3		96.2 % (4)		99.33% (2.3)
	4			97.4 % (4)	99.33% (2)
8	97.3 % (2)				

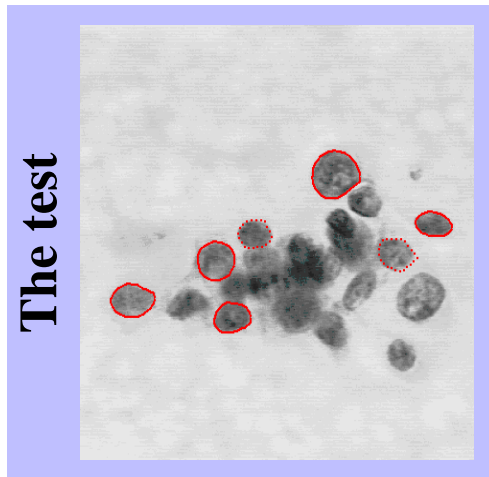
Iris controller-type: A three-rule system



Iris classifier-type: A three-rule system

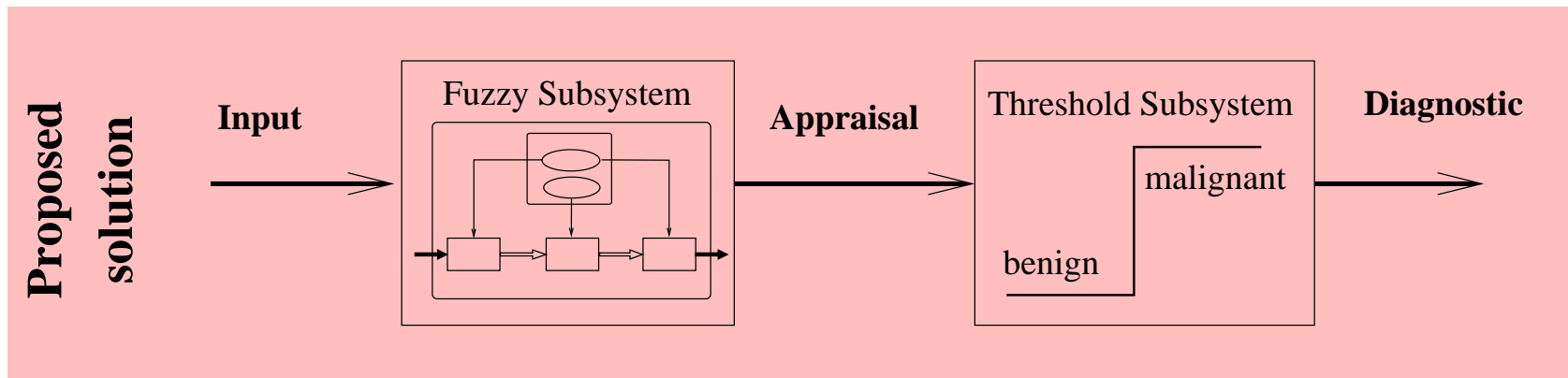


Breast cancer diagnosis: the WBCD problem



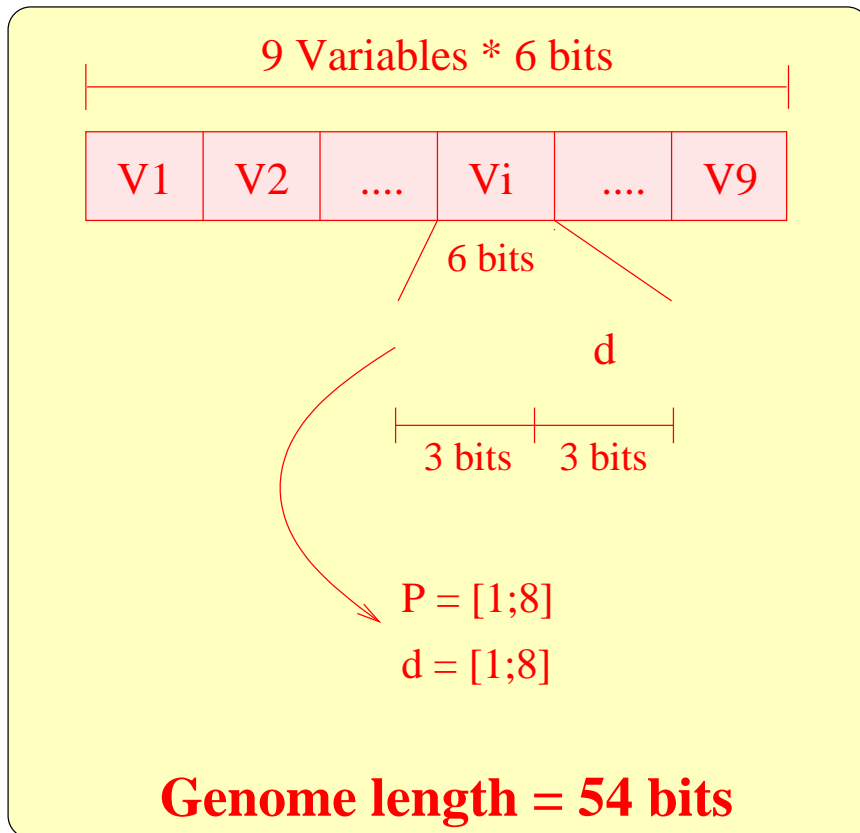
The database

<i>case</i>	v_1	v_2	v_3	...	v_9	<i>diagnostic</i>
1	5	1	1	...	1	<i>Benign</i>
2	5	4	4	...	1	<i>Benign</i>
:	:	:	:	:	:	:
683	4	8	8	...	1	<i>Malignant</i>

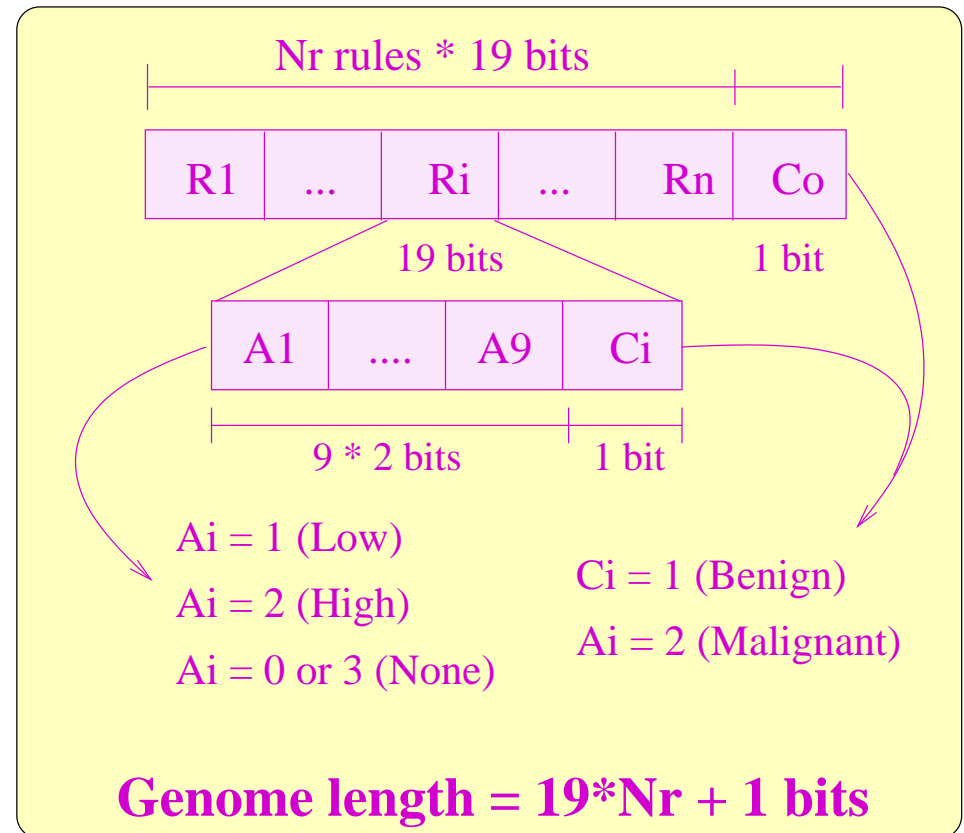


The genomes

Membership functions



Rules



Fuzzy CoCo set-up

Fitness function

$$F = F_c - a * F_v$$

F_c : Classification performance,
the most important performance measure

F_v : Number of variables
measures the interpretability

Fuzzy CoCo parameters

Population size	[30–90]
Maximum generations	1000 + 100* N_r
Crossover probability	1
Mutation probability	[0.02–0.3]
Elitism rate	{0.1–0.6}
"Fit" cooperators	1
Random cooperators	{1,2,3,4}

WBCD results: classification (longest rule)

Rules	NeuroRule Setiono (2000)	Fuzzy–genetic AIM 1999	Fuzzy CoCo – IEEE TFS 2001	
			Average	Best
1	97.36% (4)	97.07% (4)	97.36% (4.0)	97.36% (4)
2		97.36% (3)	97.73% (3.9)	98.54% (5)
3	98.10% (4)	97.80% (6)	97.91% (4.4)	98.54% (4)
4		97.80% (–)	98.12% (4.2)	98.68% (3)
5	98.24% (5)	97.51% (–)	98.18% (4.6)	98.83% (5)
6		98.10% (–)	98.18% (4.3)	98.83% (5)
7		97.95% (–)	98.25% (4.7)	98.98% (5)

Learned Boolean rules

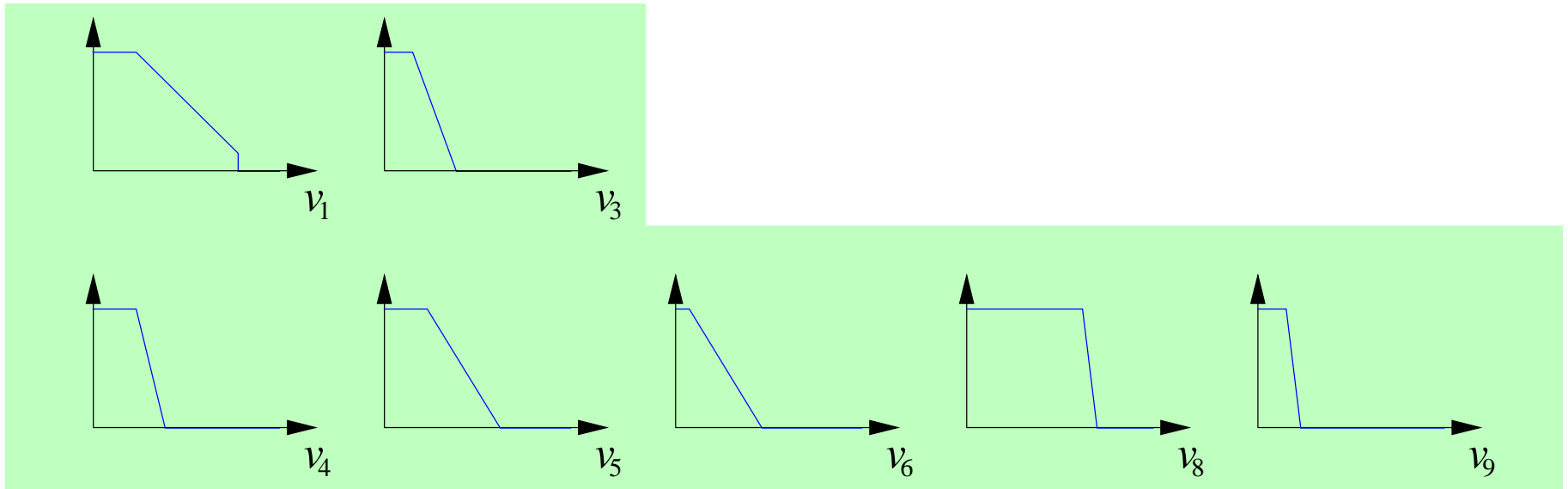
Evolved fuzzy rules

Two-rules evolved system

if (v1 is Low) and (v3 is Low) and (v5 is Low) then (output is Benign)

if (v1 is Low) and (v4 is Low) and (v6 is Low) and (v8 is Low) and (v9 is Low) then (output is Benign)

else (output is Malignant)



Classification rate = 98.54%

Computing requirements

Fuzzy GA: Single population (Peña & Sipper 99)

$$\begin{aligned} \text{Number of fitness evaluations} &= N_p * G_{\max} \\ &200 * (2000 + 500N_r) \end{aligned}$$

Single-rule systems: 500.000 fitness evaluations

Five-rule systems: 900.000 fitness evaluations

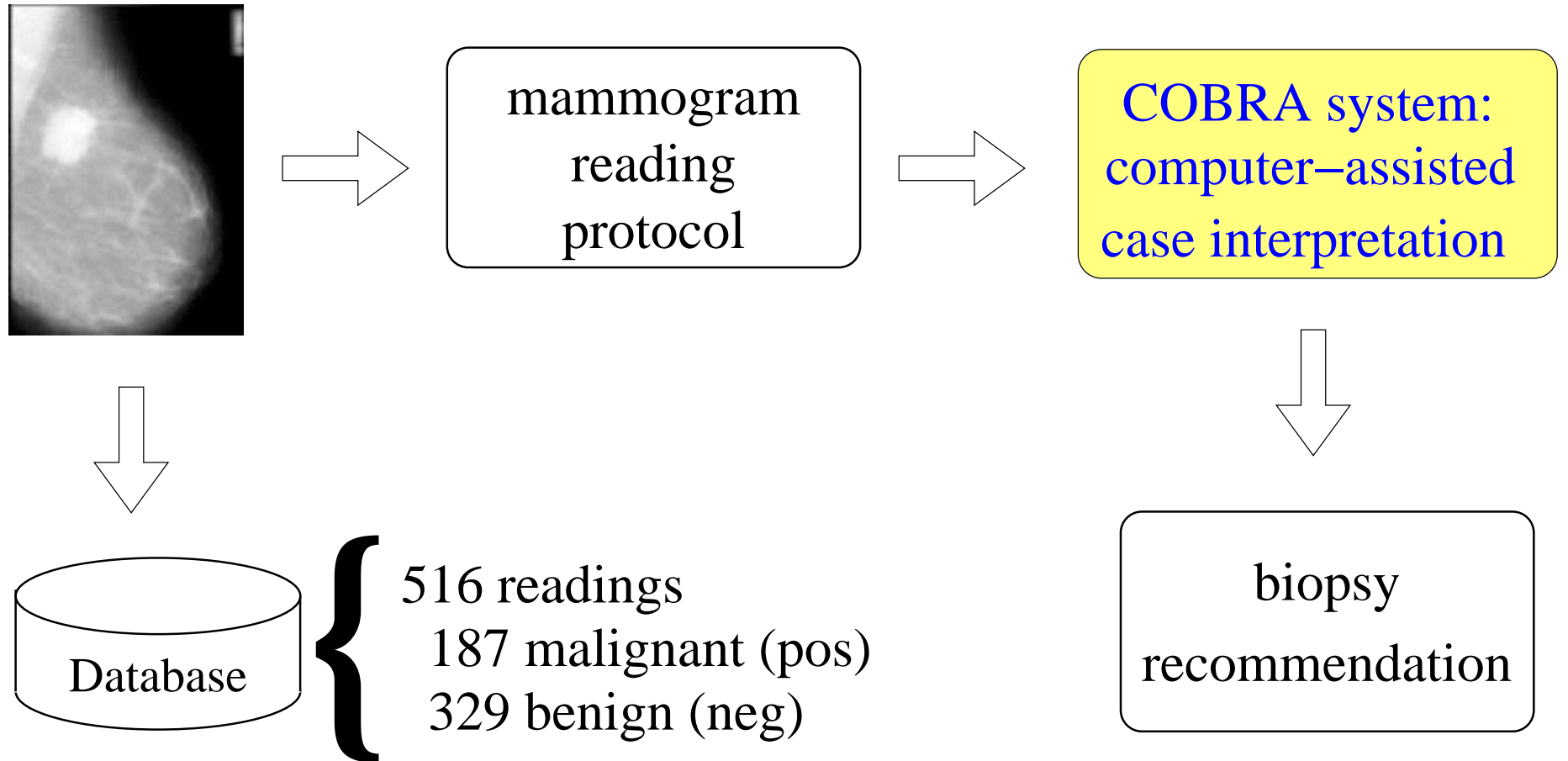
Fuzzy CoCo: Cooperative coevolution (CEC-2000)

$$\begin{aligned} \text{Number of fitness evaluations} &= 2 * N_p * G_{\max} * (N_{cf} + N_{cr}) \\ &32.000 * (1000 + 100N_r) \{ \text{worst case, } N_{cr}=3 \} \end{aligned}$$

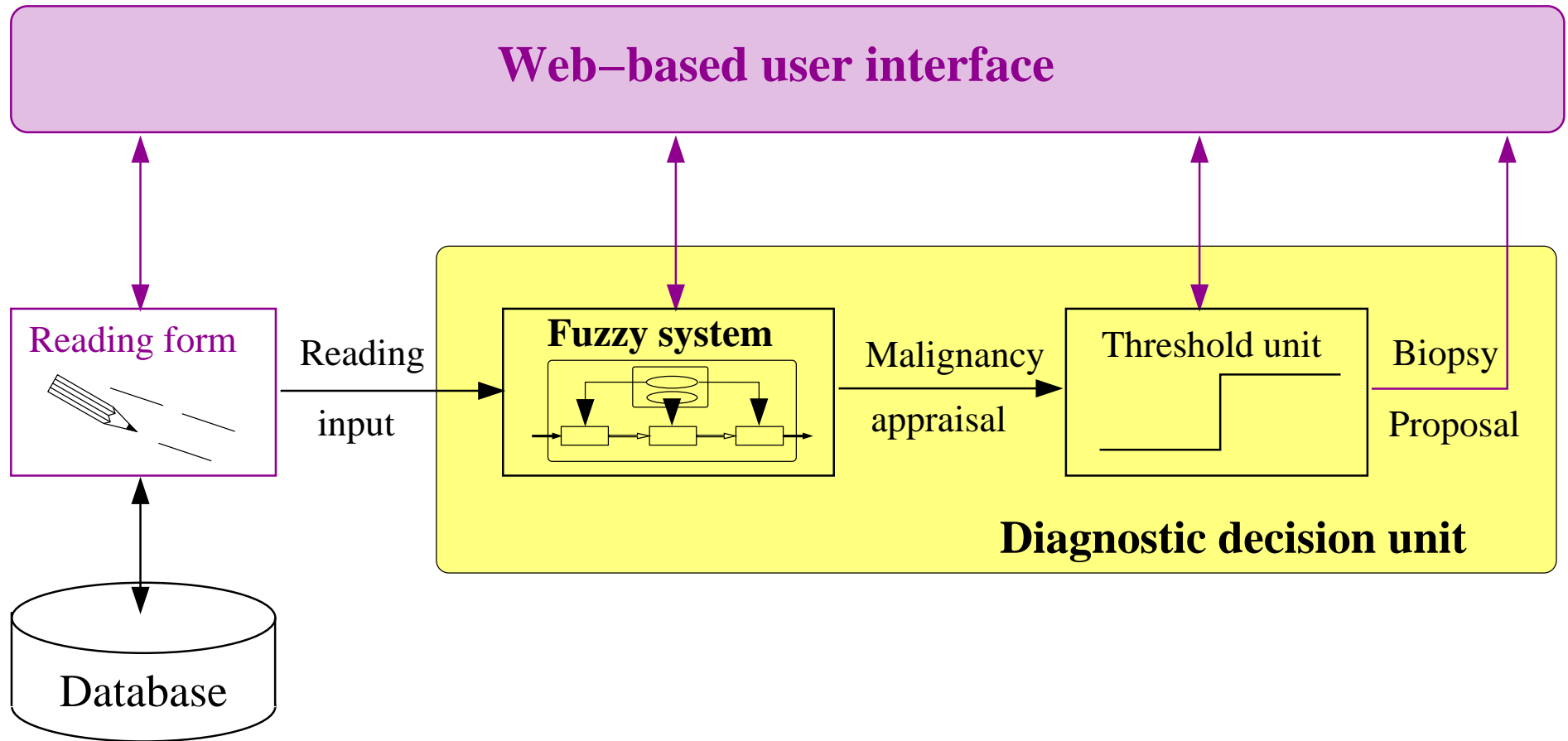
Single-rule systems: 352.000 fitness evaluations

Five-rule systems: 480.000 fitness evaluations

The problem: mammography interpretation



COBRA system: internal view



Understanding the database

CLINICAL DATA

Age :

Menopause :

Antecedents :

Variable type	Number
Binary	4
Continuous	3
Discrete	8

MICROCALCIFICATION

Group shape :

Other group features :

Group diameter (mm) :

Number of micros :

Morphology of micros :

Size irregularity :

MASS

Mass morphology :

Mass margins :

Density bigger than parenchyma :

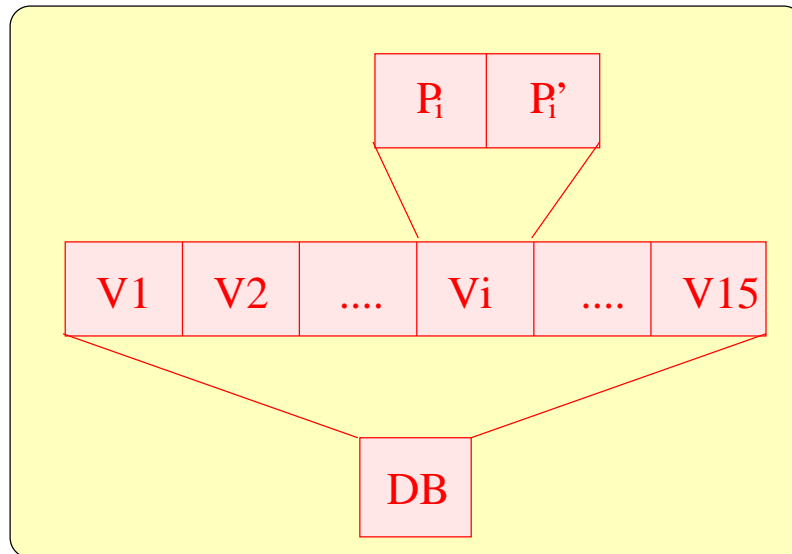
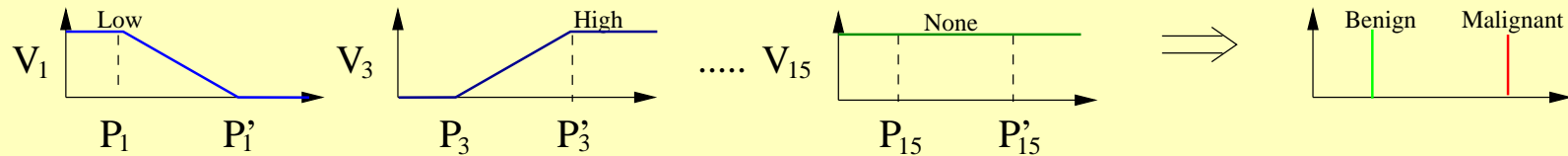
Focal distortion :

Focal symmetry :

Focal Area (mm) :

Genome encoding for linguistic labels

R_i : if (v_1 is A_{i1}) and (v_2 is A_{i2}) and (v_3 is A_{i3}) and ... and (v_{15} is A_{i15}) then (output is C_i)



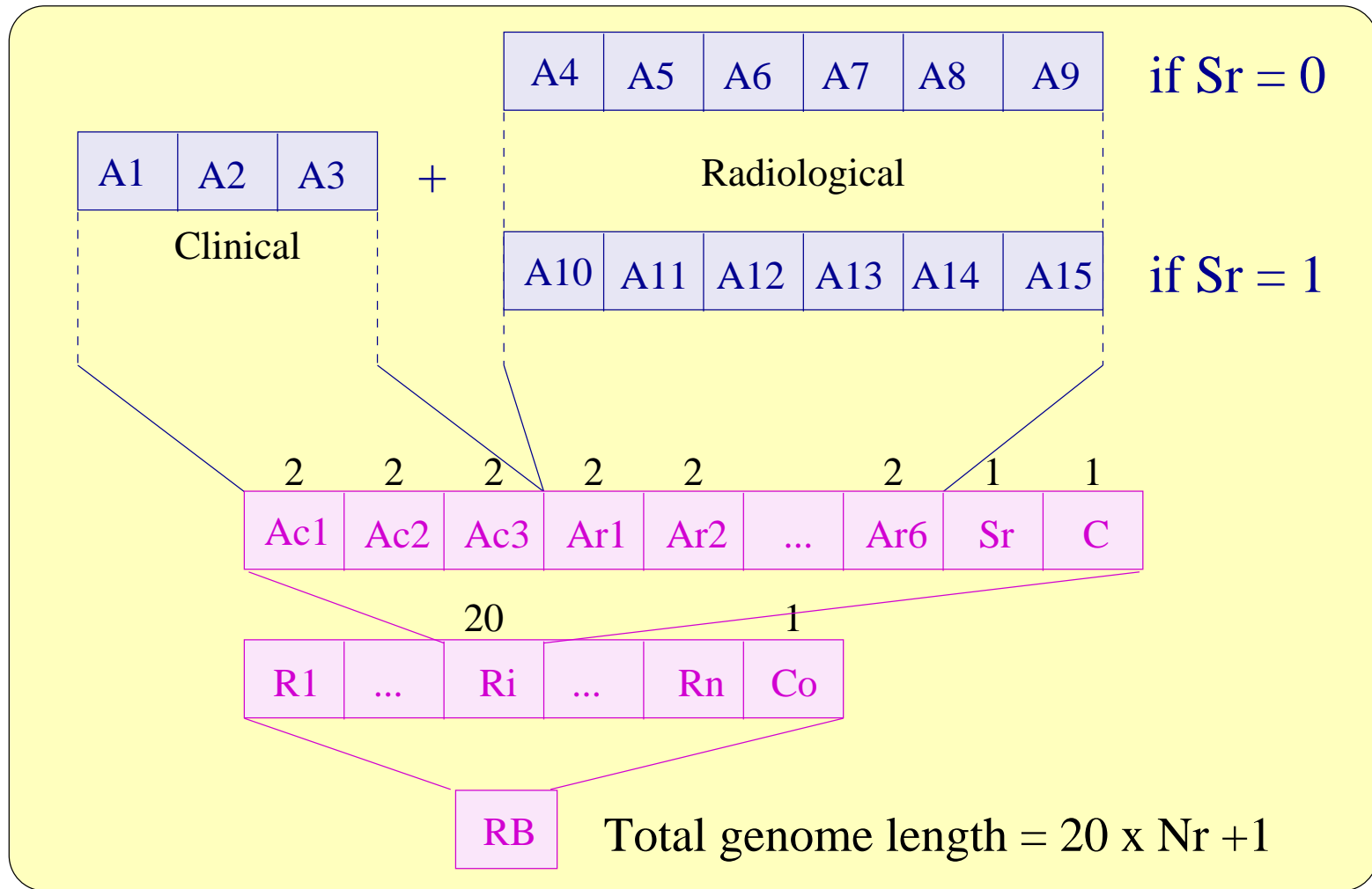
Binary variables (e.g., V_2):
not encoded

Continuous variables (e.g., V_1):
3 var. x 2 par. x 7 bits = 42 bits

Discrete variables (e.g., V_3):
8 var. x 2 par. x 4 bits = 64 bits

Total genome length = 106 bits

Genome encoding for rules



Performance measures and fitness function

Sensitivity $\frac{TP}{TP + FN}$

Specificity $\frac{TN}{TN + FP}$

Accuracy $\frac{TP + TN}{TP + TN + FP + FN}$

PPV $\frac{TP}{TP + FN}$

Basic fitness (Fbase)

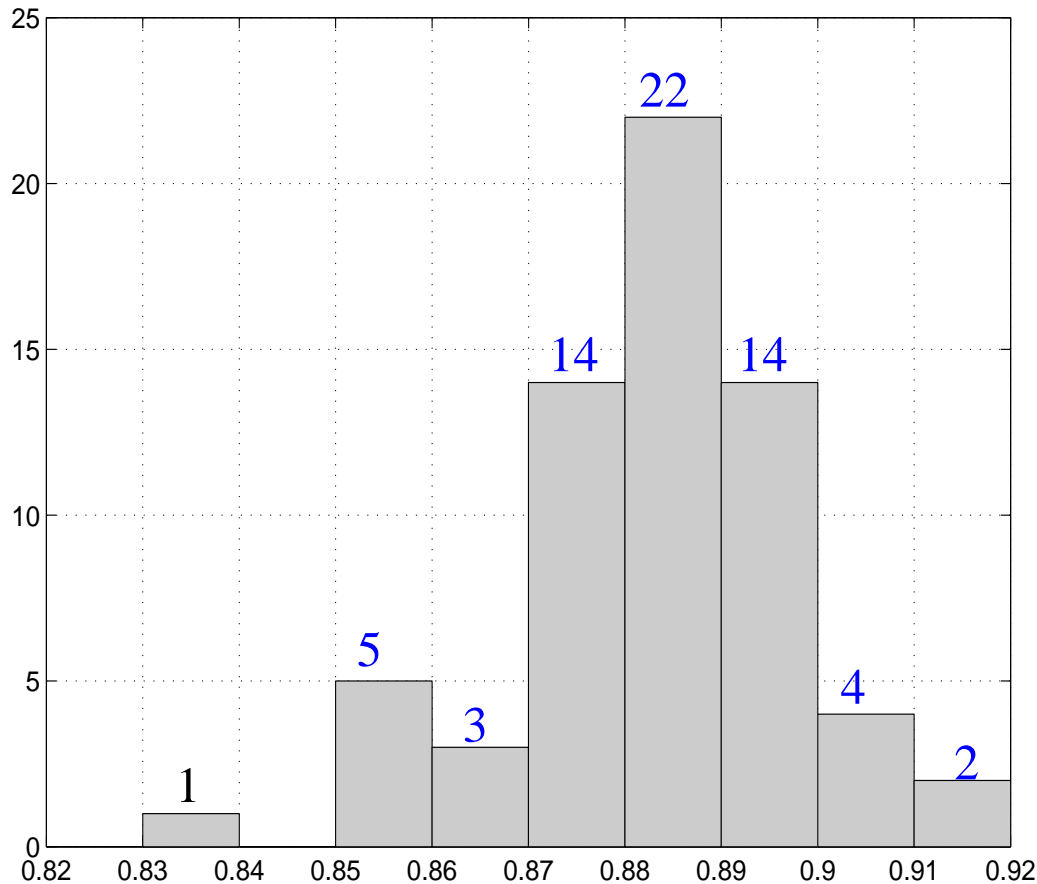
$$\frac{\text{Sensitivity} + \alpha \text{ Specificity}}{1 + \alpha}$$

Accuracy reinforcement

$$\frac{\text{Fbase} + \beta \text{ Accuracy}}{1 + \beta}$$

(note: done only if Accuracy > 0.7)

Fuzzy CoCo results on 65 runs



Average per class

Nr	Fitness	Reff	Vr
10	0.8754	9.17	2.52
15	0.8786	12.03	2.62
20	0.8934	14.15	2.59
25	0.8947	15.78	2.76

Best individual

Nr	Fitness	Reff	Vr
10	0.8910	9	2.22
15	0.8978	12	2.50
20	0.9109	17	2.41
25	0.9154	17	2.70

Performance of two selected systems

	17-rule		9-rule	
Measure	Figure	Ratio	Figure	Ratio
Sensitivity	99.47%	186/187	98.40%	184/187
Specificity	68.69%	226/329	64.13%	211/329
Accuracy	79.84%	412/516	76.55%	395/516
PPV	64.36%	186/289	60.93%	184/302

The 9-rule system with two different thresholds

	Threshold = 2		Threshold = 3	
Measure	Figure	Ratio	Figure	Ratio
Sensitivity	100.0%	187/187	98.40%	184/187
Specificity	63.22%	208/329	64.13%	211/329
Accuracy	76.55%	395/516	76.55%	395/516
PPV	60.71%	187/308	60.93%	184/302

COBRA system: reading form

The screenshot shows a web browser window with the address bar displaying `http://lslwww.epfl.ch/~cobra/`. The main content area is titled "NEW CASE" and is divided into two columns: "CLINICAL DATA" and "MICROCALCIFICATION" / "MASS".

CLINICAL DATA

- Age :
- Menopause :
- Antecedents :

MICROCALCIFICATION

- Group shape :
- Other group features :
- Group diameter (mm) :
- Number of micros :
- Morphology of micros :
- Size irregularity :

MASS

- Mass morphology :
- Mass margins :
- Density bigger than parenchyma :
- Focal distortion :
- Focal symmetry :
- Focal Area (mm) :

Buttons: **Reset** and **Diagnostic**

Footer navigation buttons: **New case**, **Test case**, **Cases List**, **Disclassified cases**, **View system**, **System performance**

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