The logical combinatorial approach to Pattern Recognition, an overview through selected works

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ABSTRACT. The so-called *logical combinatorial approach* to Pattern Recognition is presented, and works (mainly in Spanish and Russian) that are not ordinarily available, are exposed to the Western reader. The use of this approach for supervised and unsupervised pattern recognition, and for feature selection is reviewed. Also, an unified notation describing the original contributions is presented, thus rendering this important area more readable.

Our review is not exhaustive; nevertheless, most significant works are enclosed. Our hope is to motivate the reader to inquire further in these works.

This paper serves as an introduction to three articles on the *logical combinatorial approach* that appear in this issue of *Pattern Recognition*.

KEYWORDS: Pattern recognition, classification, feature selection, testor theory, logical combinatorial Pattern Recognition.

INTRODUCTION

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Several tools have been proposed since the beginning of Pattern Recognition, to solve three important problems: feature selection, supervised classification, and unsupervised classification or clustering. Fukunaga [1] defines *feature selection for representation* as a mapping from the original features (or variables) into those most efficient; tools are used for this purpose such as the discrete Karhunen-Loéve expansion, the same expansion for random processes, the estimation of eigenvalues and eigenvectors, principal component analysis [2], and factor analysis [3]. Also *Feature selection for classification* is defined in [1] as: given two or more classes, select those features that are most efficient to preserve class separability. To solve this problem, one resorts to discriminant analysis, non parametric discriminant analysis, sequential selection of quadratic variables, as well as several selection criteria of subsets of features that optimize some separability criteria among classes.

In *supervised classification*, the problem is to recognize, given a set of objects grouped into classes, in which one (or more than one) of these classes new objects or measurements belong. Several classifiers have been proposed: maximum likelihood; Bayesian, (1-nn), k nearest neighbors (k-nn), linear, quadratic, and other [1, 4-6].

In *unsupervised classification*, a sample of objects is at hand, but their clustering into groups or classes is unknown. The problem consists in defining such classes. To solve these problems, tools such as ISODATA [7], C-means [8], unsupervised Bayesian classifiers [6], grouping and dividing hierarchical classifiers [9], graph-theoretic classifiers [10, 11] are used, among others.

Work mentioned so far falls under Statistical Pattern Recognition, characterized because it works with the descriptions of the objects under study, contrasting with syntactic structural recognition [4], which works with the *parts* of the objects under study. This paper reviews

several articles developed under the Logical Combinatorial Pattern Recognition approach, in the three areas above mentioned. This approach, as the statistical approach, works with the descriptions of the objects. In this issue, a paper [12] introduces the concepts of testor, typical testor, logical-combinatorial approach, fuzzy testor, and others.

Let *U* be a (perhaps infinite) universe of objects, and, following the statistical approach, let us consider a given finite sample $M = \{O_1, ..., O_m\}$ of such (descriptions of the) objects. We shall denote by $R = \{x_1, ..., x_n\}$ the set of features or variables used to study these objects. Each of these features has associated a set of admissible values (its domain of definition) M_i , i=1,...,n. These sets of values, in contrast to the other approaches, can be of any nature: variables can be quantitative and qualitative simultaneously. Each of these sets contains a special symbol denoting absence of information (missing data). Thus, some variables are numeric; other, symbolic; incomplete information about some objects is allowed. This will turn out to be a fundamental feature of this Pattern Recognition paradigm. By a description of an object *O* we understand an n-tuple $I(O)=(x_I(O),...,x_n(O))$ where $x_i: M \rightarrow M_i$, for i=1,...,nare the variables or features used to describe it. Over M_i no algebraic, topologic or logic structure is assumed. Any of the Pattern Recognition problems mentioned above is formulated from a set of descriptions of *m* such objects.

FEATURE SELECTION

Let *M* be a given set of descriptions of *m* objects from *U* in terms of the features of *R*. We assume that the descriptions of *M* are structured in *r* subsets (classes) $K_1,...,K_r$ (not necessarily disjoint, not necessarily crisp). Under these conditions two kinds of problems can be formulated: feature selection for classification; same for description. The first case

tries to find the subset(s) of features that determine representation subspaces of the objects which are most appropriate for a later problem of supervised classification. In the second case the objective is not classification of new objects, but to determine the informational relevance of each of the features, or of subsets of them. Feature selection in this field was initiated by Dmitriev, Zhuravliov and Krendeleiev [13]. A recent book [25] goes further into this area.

Ortiz-Posadas [14] applies the logical-combinatorial model of Pattern Recognition Theory [15] to the computer-assisted medical diagnostic and prognostic. She uses three models: a medical model, of Heathfield *et al.* [16], for histopathologic diagnosis of breast diseases; a mathematical model, provided by the logical-combinatorial model of Pattern Recognition; and a computational model. Her dissertation explains how to select a suitable set of variables for the case in question. Some medical results appear in [17-18].

A CINVESTAV report [19] presents an extension to *typical testor* using analogy between patterns, and an algorithm to detect *fuzzy typical testors* of a training matrix.

A short letter [20] extends the expression to determine the feature relevance (that is, how relevant or discriminant a given feature of an object is) in crisp and fuzzy environments, and introduces an algorithm (different to that of [19]) to compute all the fuzzy typical testors of a training matrix.

Carrasco-Ochoa [21] analyzes how changes in the training matrix affect its typical testors. He does this for three kind of typical testors: (1) those of Zhuravlev [22, 23]; (2) the ε -typical testors, which use a similarity function between two objects that makes two objects similar if they differ in ε (an integer \ge 0) or less feature values; (3) those of Goldman [24].

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Lazo-Cortés *et al.* [12], in this issue, give several contrasting definitions of *testor*, and their application in supervised classification. This introduces the *logical combinatorial approach* to Pattern Recognition. An excellent introduction is also in a book [25].

Ozols and Borisov [26], in this issue, introduce *shadows composition* in order to solve fuzzy supervised classification.

Martínez-Trinidad and Ruiz-Shulcloper [27], in this issue, use the fuzzy clustering criteria of the logical-combinatorial approach, to do unsupervised pattern classification of non-numerical terms belonging to a vocabulary.

SUPERVISED CLASSIFICATION

In this kind of problems, we assume that the universe U is structured in a finite number $K_I,...,K_r$ of proper subsets, called classes, and from each of them we have a sample of descriptions of objects, the so-called training matrix $MA = \{K_I \cup,..., \cup K_r\}$. The problem is to find the membership relations from a new object from U (outside the given samples) with the *r* classes. This relationship does not have to be all or nothing. The logical combinatorial approach deals with spaces without algebraic (or of any other kind of) structure. The representation space is simply a Cartesian product, which also has the peculiarity of being heterogeneous, that is, each of the sets forming it can be of different nature: a set of real numbers, a set of labels, a set of truth values from a given logic, etc. An example of this appears in medical diagnosis problems, where descriptions take the form I(O)=(black, female, 45, 38.6⁰, 1500, ?, slight,...), where ? means absence of information. In other disciplines, too, such as Geo Sciences, Sociology, Pedagogy, Marketing, etc. (the so called

soft sciences), objects are described in terms of qualitative and quantitative variables. Thus, the tools herein presented.

Most significant models of supervised classification are those works based on *partial precedences*. There are three models: voting algorithms, Kora model, and algorithms based on representative sets.

Voting algorithms

These were conceived in a simplified manner by Zhuravlev and his group [28]. The main idea is partial precedence, that is, partial analogies. An object can resemble another object not in its totality; those parts which *do* resemble each other can give information about possible regularities; of course not all of the same magnitude. For practical problems in the disciples cited above, this model was very limited, since it only allows object descriptions based in Boolean variables. A later paper [25, 29] added some results, producing an improved parametric model comprising six steps: 1) defining the system of support sets; 2) defining the similarity function; 3) row evaluation, given a fixed support set; 4) class evaluation for a fixed support set; 5) class evaluation for all the system of support sets, and 6) resolution rule. Thus, to define a voting algorithm, is to define a set of parameters for each of the above six steps.

A support set is a non empty subset $\Omega = \{x_{i_1}, ..., x_{i_s}\}$ of features which shall be used to analyze the objects. We denote as ΩO the subdescription in terms of the features of Ω . Thus, a system of support sets denoted by Ω_A are several support sets which together will allow analysis of the objects to be classified, comparing them with objects in each one of the classes K_i i=1,...,r. Note that said analysis is done paying attention to different parts or subdescriptions of the objects, and not analyzing the complete descriptions. Examples of systems of support sets are the set of typical testors, combinations with a fixed cardinality, combinations with variable cardinality, the power set of features, etc.

A similarity function defines the form to compare the descriptions (subdescriptions) of the objects. Ruiz-Shulcloper and his group [25, 29] propose to do this starting by determining the comparison among the values of features, and evaluating these comparisons through a similarity function. A simple example of similarity function is

$$\Gamma(\Omega O_i, \Omega O_j) = \sum_{x_p \in \Omega} C_p(x_p(O_i), x_p(O_j)).$$

When the systems of support sets and the similarity function have been defined, the voting process starts in the stage of row evaluation; that is, the similarity between the different parts of the objects already classified and those to be classified is analyzed. Each row of *MA* (each object $O_i \in MA$) is compared with object *O* to be classified using the similarity function Γ . This evaluation is a function of the similarity values among the different parts being compared. An example of this evaluation is $\varphi_{\Omega}(O_i, O) = \gamma(O_i)P(\Omega)\Gamma(\Omega O_i, \Omega O)$ where we consider a weight $\gamma(O_i)$ associated to each object O_i from *MA* and a weight $P(\Omega)$ for the support set Ω .

The class evaluation for a fixed support set Ω consists in totaling the evaluations obtained for each of the objects *MA* with respect to the object *O* to be classified. This total evaluation is a function of the row evaluations already obtained. An example of this evaluation is:

$$\varphi_{\Omega}^{j}(O) = \frac{1}{|K_{j}|} \sum_{t=1,\dots,|K_{j}|} \varphi_{\Omega}(O_{t}, O), \text{ the upper index refers to the class } K_{j}.$$

In the class evaluation for all the system of support sets, evaluations are totaled for all the system of support sets. Following our example, this step could be expressed as follows:

$$\varphi^{j}(O) = \frac{1}{|\Omega_{A}|} \sum_{\Omega \in \Omega_{A}} \varphi_{\Omega}^{j}(O).$$

Finally, the resolution rule is a function that establishes a criterion taking into account each voting thus obtained, and reaches a decision concerning the relations of the object to be classified with every class of the posed problem. In general, this function has the following form: $r(\varphi^1(O),...,\varphi^r(O)) = (\alpha_1(O),...,\alpha_r(O))$. A similar manner to compute the values of

$$\alpha_{i}(O) \ i=1,...,r., \ \text{can be:} \ \alpha_{i}(O) = \begin{cases} 1 & if \varphi^{i}(O) > \varphi^{j}(O) \forall i \neq j \\ 0 & otherwise \end{cases}$$

This model for supervised classification has been applied to several practical problems [18, 30-33].

An extension to the above model is proposed by Lazo [34], where he lets the features be numeric or linguistic; that is, values can be some labels or words of natural languages (fuzzy variables). The similarity between values of the same feature is evaluated in the interval [0,1], considering as a special case a two-valued comparison criteria. An important element of this new model is that it covers problems considering fuzzy membership (in Zadeh's sense [35]) of the objects to the classes. In this manner, he can work with fuzzy support sets in which each feature belongs to a certain degree to the support set. For instance, fuzzy Goldman testors can be used as support sets. The evaluation of the similarity between objects, proposed in this work, is summarized in the following general expression: $\Gamma(\Omega O_i, \Omega O_j) = |\Omega|^{-1} \sum_{x_p \in \Omega} \mu_\Omega(x_p) C_p(x_p(O_i), x_p(O_j))$. This expression considers

the degree of membership of feature x_p to the support set Ω . In addition, if Ω is a fuzzy set, the scalar cardinality (see Zimmermann [36]) is considered. In the case of linguistic variables, one can use as criteria for comparing values of features the Hamming, Euclid or Kolmogorov-Fomin criteria, which are obtained from the homonymous distances between fuzzy sets, see [37]. To evaluate by class using a fixed support set, one works with

$$\varphi_{\Omega}^{j}(O) = \frac{1}{|K_{j}|} \sum_{t=1,\dots,|K_{j}|} \varphi_{\Omega}(O_{t},O)$$
, if in the formulation of the problem each object belongs to

only one class. To consider degrees of membership of each object to each class, the expression $\varphi_{\Omega}^{j}(O) = \frac{1}{|K_{j}|} \sum_{t=1,\dots,|K_{j}|} \mu_{K_{j}}(O_{t})\varphi_{\Omega}(O_{t},O)$ is used, where $|K_{j}|$ denotes the scalar

cardinality of class K_{j} , and $\mu_{K_{j}}(O_{t})$ denotes the degree of membership of object O_{t} to class K_{j} , j=1,...,r. For evaluation by class for all the system of support sets, one considers the average of the evaluations, in similar manner as the former model. Finally, in the resolution rule can be considered each $\varphi^{j}(O) j=1,...,r$, for each class as the fuzzy membership degrees to the classes, or as the votings for belonging to each class.

Finally we can mention the voting algorithm for linguistic variables [34]. In this model the problem of classification is posed in similar terms to defining the testors to certain degree, that is, considering that objects belong with certain degree to each class. Similarity functions are defined between object descriptions in terms of linguistic variables, as well as expressions for row evaluation of a fixed support set and for all the system of support sets.

Kora Model

These models start with Bongard's group [38], who propose the Kora-3 algorithm for solution of Geophysics and Geology problems. This was the first algorithm to be used for solving supervised classification in the Geological environment.

Kora-3 rests on the idea that classes are formed by objects fulfilling certain complex properties, composed by the conjunction of three simple properties. This algorithm is very limited since it only allows Boolean description of objects, and works only with two nonintersecting classes. The basic idea is partial precedence (in similar form to the voting algorithms), but restricted to the use of relations of any three features. A subset of three features $\Omega = \{x_{i_1}, x_{i_2}, x_{i_3}\}$ and a combination of values (a_1, a_2, a_3) for Ω , form a *complex feature* of class K_i if and only if the triplet (a_1, a_2, a_3) appears at least δ_i times in the subdescriptions corresponding to the objects of K_i and does not appear in the subdescriptions of objects of the other class. Those objects in which this combination appears are called *objects characterized* by the complex feature. On the other hand, objects characterized by less than δ_i complex features form *the reminder of class K_i*. In this manner, if two complex features r_1 y r_2 characterize exactly the same objects, they are called equivalent, and if r_1 characterizes all objects characterized by r_2 and at least one more, then it is said that r_1 is stronger than r_2 . Based on these concepts the Kora-3 algorithm is defined in three steps: (1) learning step; (2) re-learning step; (3) classification step. In the learning step the complex features are computed for each class using parameters δ_1 and δ_2 . In the relearning step *complementary complex features* are computed using new thresholds $\delta'_{I} < \delta_{I}$ and $\delta'_2 < \delta_2$ in the remainder of the classes. Finally, the classification step counts how many complex features of each class characterize or vote in favor of the new object to be classified, and selects the class that provides the largest number of complex features. This algorithm became very limited given the characteristics of practical problems, hence the first extension proposed [39, 40] modified the concept of a complex feature, allowing now a combination of three values to appear δ_i times (sufficiently) in the classes and δ' times (sufficiently little) in the other class. Later, De la Vega-Doria *et al.* [41] propose an extension which allows the use of any system of support sets and not only support sets formed by three features, it is allowed to work with more than two classes, overlap among classes is allowed, the type of variables can be of any sort, lack of information is allowed, and in the most general case, classes can be fuzzy, that is, object belong to a class to a certain degree. To reach this extension the concept of complex feature was modified once more, giving rise to the concept of fuzzy δ_i -complex feature. A combination of values $a=(a_{i1},...,a_{ip})$ for the features of a support set $\Omega = \{x_{i_1},...,x_{i_s}\}$, forms a δ_i -complex fuzzy feature (a,Ω) for K_i , with degree $\mu_i((a,\Omega))$, i=1,...,r, if and only if: a) $\exists O \in K'_i | x_{i1}(O) = a_{i1}$

^ ... ^
$$x_{ip}(O) = a_{ip}$$
; b) $\sum_{j=1}^{m} \Gamma(\Omega O_j, a) \mu_i(O_j) \ge \delta_i$; c) $\sum_{j=1}^{m} \Gamma(\Omega O_j, a) (1 - \mu_i(O_j)) \le \delta_i$; and d)

$$\mu_i((a,\Omega)) = \frac{\sum_{j=1}^m \Gamma(\Omega O_j, a) \mu_i(O_j)}{\sum_{j=1}^m \mu_i(O_j)}, \text{ where } \delta i > 0 \text{ and } \delta' i > 0 \text{ are thresholds. When working with}$$

crisp classes, the assumption is that $\mu_i((a,\Omega))=1$ for all (a,Ω) associated with K_i i=1,...,r. This redefinition allows (as formerly mentioned) to handle any system of support sets. In addition, it is considered in the expression the case of classes being fuzzy and each δ_i complex feature has associated a degree of membership to said set, calculated in the basis of the similarity of the combination of values a with the corresponding subdescriptions in the respective class. The set of all δ_i -complex fuzzy sets for K_i is denoted as $RC(K_i)$. On the other hand, the set of all objects $O \in K_i$ such that $\sum_{(a,\Omega) \in RC(K_i)} \Gamma(\Omega O, a) < \eta_i$ will be called η_i -

fuzzy reminder of the class K_i and is denoted by $r(K_i)$. In the same manner as in the Kora-3

algorithm, the complementary complex features are computed in the remainder of the classes, but now using the new formulation. To each complex feature and each complementary complex feature, one computes a weight or importance $P((a,\Omega))$ as

$$P((a,\Omega)) = \sum_{x_k \in \Omega} P(x_k) \sum_{O_j \in K_i} \Gamma(\Omega O, a) P(O_j) \text{ where } P(x_k) \text{ and } P(O_j) \text{ are the weights of feature } x_k$$

and of object O_j . Finally, to classify a new object O, it is compared with all δ_i -complex fuzzy features of each class K_i i=1,...,r, in the following manner:

$$\mu_i(O) = max \frac{\sum_{(a,\Omega) \in RC(K_i)} \Gamma(\Omega O, a) \mu_i((a,\Omega)) P((a,\Omega))}{\sum_{(a,\Omega) \in RC(K_i)} \Gamma(\Omega O, a) P((a,\Omega))} \cdot$$

Model based on representative sets

This model [42] is based in partial precedences and representative sets. The idea is to evaluate information in favor and against objects belonging to classes, and to consider that the parameters used for classification should be associated with each class. In contrast with previous models, this model can use a different system of support sets for each class. The rationale is that, for a particular class, a combination of features or ranges in values can provide valuable information to characterize such class, but one can not conclude that the same combination behaves similarly for another class. Hence, the model proposes to determine the system of support sets for each class. { Ω }_{*j*} denotes the system of support sets associated with class K_j and CK_j represents the complement of the class. We define the *positive representatives set* M_j^1 for class K_j with respect to $\Omega \in {\Omega}_j$ as the set of all values corresponding to Ω in the subdescriptions of the objects in K_j that are present η_j times and are not present in in CK_j . Similarly, the *negative representatives set* M_j^0 for class K_j with respect to Ω is defined as the set of all values corresponding to Ω in the subdescriptions of the objects in CK_j that are present η_j times and are not present in K_j . The set of all values corresponding to Ω in the subdescriptions of the objects of K_j that are not elements neither of the negative representatives set nor of the positive representatives set form the *neutral set of combinations* for class K_j denoted by M_j^{Δ} . The algorithm rests on these three concepts. This method differentiates between the informational value of features and of objects, which can be calculated in some other way, see [20, 25], or be given by the expert. The first proposal of this model is Boolean, that is, considers descriptions of objects formed by Boolean features only. Thus, the algorithm is: 1) define the systems of support sets (and their associated parameters) for each class; 2) compute the sets M_j^1 , M_j^0 and M_j^{Δ} for each class; 3) to classify a new object O, compute $\varphi_j(\Omega O)$ for each support set Ω of each class

 K_j j=1,...,r. This expression is defined in three cases: a) $\varphi_j(\Omega O) = (p(x_{i_1})+,...,+p(x_{i_s}))(p(\Omega O_{i_1})+,...,+p(\Omega O_{i_t}))$ if $\Omega O \in M_j^1$, where $p(x_i)$ is the weight of the variables conforming Ω and $p(\Omega O_i)$ means the weight of object $O_i \in K_j$ that coincides with O according to Ω ;

b)
$$\varphi_j(\Omega O) = (p(x_{i_1}) + \dots + p(x_{i_s}))(p(\Omega O_{i_1}) + \dots + p(\Omega O_{i_t}))$$
 if $\Omega O \in M_j^0$, where $p(x_i)$ is as

before, and $p(\Omega O_i)$ means the weight of object $O_i \in CK_j$ that coincides with O according to Ω ; c) $\varphi_i(\Omega O) = 0$ if $\Omega O \in M_i^{\Delta}$. 4) In this step the total evaluation $\varphi_i(O)$ is computed for

each class
$$K_j = 1, ..., r$$
, as $\varphi_j(O) = \frac{1}{|\{\Omega\}_j|} \sum_{\Omega \in \{\Omega\}_j} \varphi_j(\Omega O)$. 5) Finally, compute the r-tuple

 $(\alpha_I(O),...,\alpha_r(O))$ in which one finds the membership relation $\alpha_i(O)$ of object O for each one

of the *r* classes, as follows: $\alpha_j(O) = \begin{cases} 1 & \text{if } \varphi_j(O) > 0 \\ 0 & \text{if } \varphi_j(O) < 0 \end{cases}$. This model proved to be of limited * $\text{if } \varphi_j(O) = 0$

use for solution of practical problems, where objects are described by features that can take any kind of values. Thus, an improvement is proposed in [25], which allows such descriptions (and values) but defines the comparison criteria to yield Boolean values. In this manner, a similarity function among two subdescriptions is defined, which can be used to determine if a subdescription belongs to M_{j}^{1} , M_{j}^{0} or M_{j}^{Δ} in the same manner as in the Boolean case, but now verifying if they are similar or not. For this, the following expressions are evaluated:

$$\varphi_{j}^{+}(\Omega O) = \left(p(x_{i_{1}})+,\ldots,+p(x_{i_{s}})\right)\left[\chi_{1}\sum_{O_{i}\in K_{j}}p(O_{i})\Gamma(\Omega O,\Omega O_{i})+\chi_{0}\sum_{O_{i}\in CK_{j}}p(O_{i})\overline{\Gamma}(\Omega O,\Omega O_{i})\right] \quad \text{and}$$

$$\varphi_{j}^{-}(\Omega O) = \left(p(x_{i_{1}})+,...,+p(x_{i_{s}})\right)\left[\chi_{1}\sum_{O_{i}\in CK_{j}}p(O_{i})\Gamma(\Omega O,\Omega O_{i})+\chi_{0}\sum_{O_{i}\in K_{j}}p(O_{i})\overline{\Gamma}(\Omega O,\Omega O_{i})\right], \text{ where }$$

 $p(x_i)$, $p(O_i)$ are weights associated to the features and objects, χ_0 , χ_1 are weight factors and $\overline{\Gamma}$ takes values opposite to those of Γ . We then decide if ΩO is in M_j^1 , M_j^0 or M_j^{Δ} as follows: a) $\Omega O \in M_j^1$ if $\varphi_j^+(\Omega O) - \varphi_j^-(\Omega O) > \delta$; b) $\Omega O \in M_j^0$ if $\varphi_j^-(\Omega O) - \varphi_j^+(\Omega O) > \delta$; and c) $\Omega O \in M_j^{\Delta}$ if $|\varphi_j^+(\Omega O) - \varphi_j^-(\Omega O)| < \delta$. Except for these changes, the previous algorithm remains the same.

Carrasco [43] proposes an extension to [42], in which objects can be described by features of any kind, classes do not need to be disjoint; they can be fuzzy, although imposing the restriction that for every class there must exist at least one object that is closer to totally

belonging to the class than of not belonging to the class, which is reasonable, since this is a problem of supervised classification. This extension is reached by extending the concept of positive, negative and neutral representatives sets, as follows. Let Ω be a support set for class K_j . The positive representatives set for class K_j with respect to Ω , denoted by M_j^1 , is defined such that: ∃O∈MA | as the set of all ΩO_p 1) $\Omega O = \Omega O_p;$ 3) $\sum_{i=1}^{m} \Gamma(\Omega O_i, \Omega O_p)(1 - \alpha_j(O_i)) < \delta_j;$ 2) $\sum_{i=1}^{m} \Gamma(\Omega O_i, \Omega O_p) \alpha_j(O_i) \ge \eta_j;$

4)
$$\mu_j(\Omega O_p) = \frac{\sum_{i=1}^m \Gamma(\Omega O_i, \Omega O_p) \alpha_j(O_i)}{\sum_{i=1}^m \alpha_j(O_i)}$$
, where $\mu_j(\Omega O_p)$ denotes the degree by which the

combination of values ΩO_p is positive representative of K_j ; $\alpha_j(O)$ is the degree of membership of O to the class K_j and $(1-\alpha_j(O))$ represents the degree of membership to the complement of K_j , η_j represents the minimum similarity threshold of ΩO_p with all the objects in K_j ; δ_j represents the maximum similarity threshold of ΩO_p with respect to all objects in the complement of K_j . Analogously, the negative representatives set M_j^0 is redefined by interchanging in expressions 2), 3) and 4), $\alpha_j(O)$ by $(1-\alpha_j(O))$. The combinations ΩO_p that are not in M_j^1 nor in M_j^0 will be neutral combinations and shall be denoted by M_j^{Δ} . The informational relevance of a positive (negative) representative ΩO_p for a class K_j , is denoted by $P(\Omega O_p)$, and it is computed by the expression $P(\Omega O_p) = (p(x_{i_1})+,...,+p(x_{i_n})) \sum_{i=1}^{m} p(O_i) \alpha_i(O_i) \Gamma(\Omega a, \Omega O_i)$. Given an object O to classify,

the total positive evidence, denoted by $\varphi_j^+(O)$, and the total negative evidence, denoted by $\varphi_j^-(O)$, are computed for each one of the classes K_j , j=1,..,r as follows:

$$\varphi_{j}^{+}(O) = \begin{cases} \sum_{\Omega \in \{\Omega_{1}^{i}, \dots, \Omega_{s_{j}}^{i}\}} \sum_{\Omega O_{p} \in \Omega M_{1}^{i}} \Gamma(\Omega O, \Omega O_{p}) P(\Omega O_{p}) & \text{if the classes are crisp sets} \\ \sum_{\Omega \in \{\Omega_{1}^{i}, \dots, \Omega_{s_{j}}^{i}\}} \sum_{\Omega O_{p} \in \Omega M_{1}^{i}} \Gamma(\Omega O, \Omega O_{p}) P(\Omega O_{p}) \mu(\Omega O_{p}) & \text{if the classes are fuzzy sets} \\ \varphi_{j}^{-}(O) = \begin{cases} -\sum_{\Omega \in \{\Omega_{1}^{i}, \dots, \Omega_{s_{j}}^{i}\}} \sum_{\Omega O_{p} \in \Omega M_{1}^{0}} \Gamma(\Omega O, \Omega O_{p}) P(\Omega O_{p}) & \text{if the classes are crisp sets} \\ -\sum_{\Omega \in \{\Omega_{1}^{i}, \dots, \Omega_{s_{j}}^{i}\}} \sum_{\Omega O_{p} \in \Omega M_{1}^{0}} \Gamma(\Omega O, \Omega O_{p}) P(\Omega O_{p}) & \text{if the classes are fuzzy sets} \end{cases} \end{cases}$$

and then the total evaluation $\varphi_j(O)$ is computed for each class K_j , j = 1, ..., r, as follows:

$$\varphi_{j}(O) = \begin{cases} 0 & \text{if } \varphi_{j}^{+}(O) = \varphi_{j}^{-}(O) = 0 \\ \\ \frac{\varphi_{j}^{+}(O) + \varphi_{j}^{-}(O)}{2max\{\varphi_{j}^{+}(O), |\varphi_{j}^{-}(O)|\}} & \text{otherwise} \end{cases}$$

This evaluation estimates the membership of O in class K_j . 1) If working with crisp classes,

use:
$$\alpha_j(O) = \begin{cases} 1 & \text{if } \varphi_j(O) > 0 \\ 0 & \text{if } \varphi_j(O) < 0 \text{ in this case * denotes abstention.} \\ * & \text{if } \varphi_j(O) = 0 \end{cases}$$

2) For fuzzy classes use:

$$\alpha_{j}(O) = \begin{cases} 0.5 + \varphi_{j}(O)(2\max_{O_{i} \in MA} \{\mu_{j}(O_{i})\} - 1) & \text{if } \varphi_{j}(O) \ge 0\\ \\ 0.5 + \varphi_{j}(O) & \text{if } \varphi_{j}(O) < 0 \end{cases}.$$

A detailed study of supervised classification algorithms based in partial fuzzy precedence is given in [44].

In Logical Combinatorial Pattern Recognition, some other models of supervised classification have been proposed. Such is the case of algorithms based in the weight of the patterns [25, 45], among which we find: voting algorithms; those based in the ideal of the class, and algorithms based in exactness thresholds.

Another model of supervised classification is the algorithms based in typicality and contrast. This model was initially proposed by Diordenko and Vaskovskii [46] for the solution of problems with geologic anomalies, its purpose was not to classify geological regions, but to structure their data so as to facilitate the classification. Based on these ideas, Vega [47] extended the algorithm and developed a supervised classification algorithm based in notions of typicality and contrast. This particular algorithm has been used to classify patients with cleft palate [48].

UNSUPERVISED CLASSIFICATION

In this problem it is assumed that the structure of universe U is not known. To find such structure, in general a subset MI of U, the initial sample, is given. On occasions the number of groups that there should result (or the number desired), is known, and the problem is termed *restricted unsupervised classification*. On other occasions, this number is also unknown (*free unsupervised classification*). Here, only a representation space that is a Cartesian product without further properties is used. The problem is precisely to find the classes, the groupings. Such groups need not be formed by crisp sets.

Free unsupervised classification

The first tools in this field appear in Sirotinskaia [49], who proposes the CLASS algorithm. This is a hierarchical algorithm and was written to solve geology prospecting problems. It is based in the concept of a compact set. A set NU_r is a compact set if and only if

a) $\forall O_i \in MI$

 $O_t \neq O_{i_{p+1}}$

$$\begin{split} & [\mathbf{O}_{i} \in \mathrm{NU}_{r} \land (\max_{\substack{\mathbf{O}_{i} \in \mathrm{MI} \\ \mathbf{O}_{i} \neq \mathbf{O}_{i}}} \left\{ \Gamma(\mathbf{O}_{i}, \mathbf{O}_{t}) \right\} = \Gamma(\mathbf{O}_{i}, \mathbf{O}_{j}) \geq \beta_{0} \lor \max_{\substack{\mathbf{O}_{i} \in \mathrm{MI} \\ \mathbf{O}_{i} \neq \mathbf{O}_{j}}} \left\{ \Gamma(\mathbf{O}_{j}, \mathbf{O}_{i}) \right\} = \Gamma(\mathbf{O}_{i}, \mathbf{O}_{i}) \geq \beta_{0}] \end{split}$$

$$b) \forall \mathbf{O}_{i}, \mathbf{O}_{j} \in \mathrm{NU}_{r} \exists \mathbf{O}_{i_{1}}, \dots, \mathbf{O}_{i_{q}} \in \mathrm{NU}_{r}$$

$$[\mathbf{O}_{i} = \mathbf{O}_{i_{1}} \land \mathbf{O}_{j} = \mathbf{O}_{i_{q}} \land \forall p \in \{1, \dots, q-1\}; \left[\max_{\substack{\mathbf{O}_{i} \in \mathrm{MI} \\ \mathbf{O}_{i} \neq \mathbf{O}_{i_{p}}}} \left\{ \Gamma(\mathbf{O}_{i_{p}}, \mathbf{O}_{i}) \right\} = \Gamma(\mathbf{O}_{i_{p}}, \mathbf{O}_{i_{p+1}}) \geq \beta_{0}]]$$

c) Every isolated object constitutes a β_0 -compact (degenerate) set. This definition was proposed by Martínez [50,51]. The definition of [49] does not appear here.

The CLASS algorithm is summarized as follows: step 1) Compute the values of all possible pairs of MI; that is, form a similarity matrix $MS = \|\Gamma(O_i, O_j)\|_{m \times m}$ where *m* is the number of objects of *MI*; step 2) Compute $\beta_0 = \frac{2}{m(m-1)} \sum_{i=1}^{m-1} \sum_{j=i+1}^{m} \Gamma(O_i, O_j)$; step 3) Compute the β_0 -compact sets; step 4) Compute the similarity measures among all compact sets

$$NU_1,...,NU_c$$
 using the expression: $\Gamma(NU_s, NU_v) = \frac{1}{|NU_s|} \sum_{t=1}^{|NU_v|} \sum_{q=1}^{|NU_v|} \Gamma(O_t, O_q)$; step

5) compute a new β_0 ' from the matrix of comparisons among nuclei, and repeat steps 4 and 5. In this same sense Ruiz *et al.* [29] propose a modification based in the HOLOTYPE algorithm (originally proposed by Y.A. Voronin *et al.* [52] for supervised classification) giving rise to a new algorithm analogous to CLASS but now using the concept of connected set –which is the one used by HOLOTYPE. A set NU_r is a *connected set* if and only if: a) $\forall O_i, O_j \in NU_r \exists O_{i_1}, ..., O_{i_q} \in NU_r$

$$[O_{i}=O_{i_{1}} \land O_{j}=O_{i_{q}} \land \forall p \in \{1,...,q-1\} \ \Gamma(O_{i_{p}}, O_{i_{p+1}}) \geq \beta_{0}]$$

b) $\forall O_i \in MI [(O_j \in NU_r \land \Gamma(O_i, O_j) \ge \beta_0) \Rightarrow O_i \in NU_r]$

c) Every β_0 -isolated object is a β_0 -connected (degenerate) set. This definition was proposed by Montellano [53, 54]. The definition of connected set of [52] does not appear here.

Also, the TAXDIF model [53, 54], based on graphs, which allows the descriptions of objects to be given by features of any kind, was developed. It proposes crisp and fuzzy clustering criteria for grouping of objects, similar to the compact and connected criteria used in CLASS and HOLOTYPE. A significant contribution in this new model is the definition of crisp grouping criterion Π . This definition generates several grouping criteria, including the connected and compact criteria. Similarly, the model defines the concept of fuzzy grouping criterion Π_d which allows, from a crisp structuring generated by a crisp criterion Π , the generation of a fuzzy structuring of the original sample. That is, in the result, each object belongs in different degree to each of the clusters (clouds). Also, TAXDIF is flexible in handling several similarity functions, which can be partial functions and be defined using support sets, such as those used in supervised classification, see [25]. As example, membership function defined for connected the sets is:

criteria is given in [51]. Membership of an object in a crisp connected set is given by the

fulfillment of certain similarity properties, which establish the definition of crisp grouping criterion. Similarly, the degree of membership of an object O_i to a connected (fuzzy) cloud is given in terms of the degree of fulfillment of said properties.

This model is an alternative to the C-means fuzzy algorithm, which is defined for numeric features. The model of TAXDIF algorithms can be described by the following steps: step 1) Define the similarity function Γ ; step 2) Determine threshold β_0 ; step 3) Define the fuzzy grouping criterion Π_d ; step 4) Generate the family of crisp groupings τ ; step 5) Define the family of fuzzy groupings or clouds corresponding to τ . Varying the parameters results in different algorithms.

Due to the features of the model and its potential in the solution of problems such as those present in soft-sciences, a theoretical study was performed, about the set relations among the groupings generated by several grouping criteria.

Fig. 1

This study shows that for the same β_0 value, the structures (groupings) found for several grouping criteria are related (see figure 1). For instance, connected groupings are unions of compact groupings. Other relations are in Martínez *et al.* [51]. This hierarchy imposes an order among the structures or groupings, following their generality, which can be of use in practice. Low levels contain very specific categories (including isolated prominent objects); upper levels are formed by more general structures. This disposition of said structures can provide, for a same universe of objects, different visions, with different abstraction levels. [55] proposes these tools for the solution of semantic grouping problems, in this case

objects are related key words from documents, and the problem is to find word groupings based in semantic similarity. [27] proposes an algorithm for constructing classes of a thesaurus.

Now a days, unsupervised classification tools for large information volumes are being developed [56]. Also, conceptual unsupervised clustering has been proposed [57-60]. In these works one finds contributions to the so-called (in machine learning) unsupervised learning or conceptual clustering. The problem is: from a set of objects, find how these objects are grouped and, in addition, find the properties or concepts that objects in each group satisfy. This research line has aroused much interest, in recent years, for its potential in data mining problems.

Restricted unsupervised classification

Logical Combinatorial Pattern Recognition has devoted less effort to these problems. Nevertheless, algorithms have been proposed where the number of desired clusters is specified *a priori*. For instance, García and Martínez [61, 62] proposed a modification to the C-means algorithm allowing object descriptions to be of any nature. Keeping the original structure of the algorithm, they use, instead of a distance measure, a similarity measure among objects, such as those of [25]. The similarity function must fulfil: 1) $\Gamma(O_j, O_k) \in [0,1]$ for $1 \le j \le m$ and $1 \le k \le m$ (*m* is the number of objects to group); 2) $\Gamma(O_j, O_j) = 1$ for $1 \le j \le m$; and 3) $\Gamma(O_j, O_k) = \Gamma(O_k, O_j)$ for $1 \le j \le m$ and $1 \le k \le m$. Let u_{ik} be the membership degree of object O_k in grouping C_i , and let R^{com} be the set of all real matrices $c \le n$. Any c-partition [61] of the data set is represented by a matrix $U = [u_{ik}] \in R^{com}$, which satisfies: 1) $u_{ik} \in \{0,1\}$ for $1 \le j \le m$ and $1 \le k \le m$; 2) $\sum_{n=1}^{c} u_{ik} = 1$ for $1 \le k \le m$; and 3) $\sum_{n=1}^{m} u_{ik} > 0$ for

 $1 \le i \le c$. Then, the partition matrix U is determined by the maximization of the objective function given by $J(U) = \sum_{i=1}^{c} \sum_{k=1}^{m} u_{ik} \Gamma(O_i^r, O_k)$ where $\Gamma(O_i^r, O_k)$ is the similarity between the most representative object O_i^r ("the center") in grouping C_i and object O_k . Notice that in this case "the center" is an object belonging to the sample, instead of a fictitious element such as in classical C-means. Working with a similarity measure (and not with a distance) implies that the calculation from seeds can not be realized by using centroids (as the original work does, since it works in multidimensional metric spaces). [61, 62] proposes a way to compute these representative objects, using similarity. Once these elements are found the functional J(U) is maximized when u_{ik} is determined as follows: $\begin{bmatrix} 1 & if \Gamma(O_i^r, O_i) = max \{\Gamma(O_i^r, O_i)\} \end{bmatrix}$

$$u_{ik} = \begin{cases} 1 & \text{if } f(o_i, o_k) & \text{man}_{1 \le q \le c} f(o_q, o_k) \\ 0 & \text{otherwise} \end{cases}$$
. That is, object O_k shall be assigned to the cluster

having its representative object most similar with O_k .

Thus, the algorithm is: step 1) Select *c* objects in the sample as initial seeds. Fix *ni*', the maximum number of iterations, and set *ni*=0; step 2) Compute the partition matrix $U=U^{(ni)}$ using the previously mentioned initial seeds; step 3) Determine the representative objects of the groupings of matrix $U^{(ni)}$; step 4) Compute the partition matrix $U^{(ni+1)}$; step 5) If the set of representative objects is the same as in the last iteration, stop. Otherwise, increment ni=ni+1; step 6) If ni>ni' stop. Else, go to step 3.

More recently, other restricted unsupervised classification algorithm [64] has been proposed, based in the concept of connected set. By introducing suitable concepts and results, this algorithm works with any similarity function and with data described by features of any kind. Also, it determines theoretically how many different ways there are to partition a space into connected components, the value of thresholds β_0 characterizing them, and the cardinality of each of these partitions.

Acknowledgments

One of the authors (A. G.) acknowledges the hospitality of Dr. Robert S. Ledley, Editor of *Pattern Recognition*, for the opportunity to be Guest Editor of the Special Section on non conventional pattern classifiers, in this issue.

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CINVESTAV stands for Centro de Investigación y Estudios Avanzados del IPN.

CIC stands for Centro de Investigación en Computación del IPN.

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