

# A CLUSTER FRAMEWORK FOR DATA MINING MODELS

## *An Application to Intensive Medicine*

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**Abstract:** Clustering is a technique widely applied in Data Mining problems due to the granularity, accuracy and adjustment of the models induced. Although the referred results, this approach generates a considerable large set of models, which difficult the comprehension, the visualization and the application to new cases. This paper presents a framework to deal with the enounced problem supported by a three-dimensional matrix structure. The usability and benefits of this instrument are demonstrated trough a case study in the area of intensive medicine.

## 1 INTRODUCTION

Medical prognosis has played an increasing role in health, namely in the critical care medicine. This fact induced the medical community to take a more active interest in developing models for mortality prediction and organ failure diagnosis based on Artificial Intelligence (AI) techniques (Hanson et al, 2001), that make possible the doctors pro-active action. This is, as it can be easily understood, a critical task, since the premature detection of malfunctions in the organism may allow physicians to respond quickly with therapy. In this context, the existence of large Databases (DB) containing Intensive Care Units (ICU) clinical information, motivate and enable the application of Data Mining (DM) techniques (Cios et al, 2002), in a Knowledge Discovery Database process (KDD), to induce prediction models of organ failure in a more efficient way than other approaches (e.g., Logistic Regression) (Gilles et al, 2001). The Sequential Organ Failure Assessment (SOFA) (Vincent et al,

1998; Moreno et al, 1999) scores the dysfunction degree of an organ. It can be set to values from 0 to 4 representing the organ state. Moreover, multiple organ failure (Goris et al, 1985) highly increases the probability of the patient's death. This score is evaluated by the doctors on a daily basis taking considerable costs and time to be obtained. Obviously, this process is fallible and dependent on the doctor's expertise.

Previous work in this area provided predictive models characterized by its generality, consequently, associated to limited values of accuracy, specificity and sensitivity. The major question concerning the efficiency of such models is the patient individual adjustment. This work envisages the resolution of that bottleneck, proposing a framework for clustering the patient's prediction models, allowing the disposition of a set of predictive models (e.g., decision trees, artificial neural networks) in a three dimensional matrix.

Considering the admission data and other variables taken on the admission day, as well as

Clinical Adverse *Events* (CAEs) occurred during the patient's stay in the ICU, it is possible to predict the failure of each organ for the day following the last day of collected data (time series). A total of 72 models were created using a data set created from the EURICUS II study made in 42 ICUs on 9 UE countries, between 1997 and 1999 (<http://www.frice.nl>). The results showed the effectiveness of the proposed approach. Five of the clusters presented maximum values (100%) simultaneously for the accuracy, specificity and sensitivity. In these kinds of patients the doctors will get very useful support to their decisions.

The paper is organized as follows: after this introductory considerations, the second and third sections present the clinical data and some definitions about events and critical events; the fourth and fifth sections introduce the process of data preparation, transformation and model generation; the last two sections, preceding the eighth one that concludes the article, are dedicated to the results (presenting the achieved accuracies) and to the contributions (the framework to organize the models).

## 2 CLINICAL DATA

In this study a database was created based on EURICUS II, a study made in 42 ICUs on 9 UE countries, between 1997 and 1999. For a period of 10 months every admission to the ICU was included. This database integrates the features related to the *case-mix* (Fetter et al, 1980), namely the *Age*, the *Type of Admission* (unscheduled surgery, scheduled surgery and medical), the *Admission Source* (Operating Bloc, Recovery Room, Emergency Room, Infirmary, other ICU, other Hospital, other sources), *Diagnosis*, Gravity Index defined by *SAPSII* (Le Gall et al, 1993), *SOFA* of each Organ System (Respiratory, Coagulation, Liver, Cardiovascular, Central Nervous and Renal), *Mortality in the ICU and in the Hospital*; *Number of CAEs* for each of the parameters monitored continuously, *Length of Stay* and *Admission Day*.

By definition, an organ is considered to fail when its *SOFA* score is higher or equal than 3 in a 0 to 4 scale.

In this study, from the 5355 patients admitted to the ICUs only 4425 (82.63%) stayed for two or more days, 3105 (57.98%) stayed three or more days and 2329 (43.49%) four days or over. For the data concerning the fifth day of stay, only 1845 (34.35%) patients were considered.

## 3 CLINICAL ADVERSE EVENTS

*Events* (Ev) or *Critical Events* (CrEv) are the occurrences of values out of the established limits for the four physiologic variables that are monitored continuously. These variables are the Heart Rate (*HR*), the Systolic Blood Pressure (*BP*), the Oxygen Saturation (*SaO2*) and the Urine Output (*Diur*). A group of clinical specialists determined the intervals considered normal for each one of these parameters.

Adverse events were defined as binary variables, whose values correspond to one of two situations, in that the variable is within or not of the established limits (if yes, by how long). We considered as an *Event* when the value of the analyzed parameter maintains out of the limits, for a period equal or superior to a continuous period of 10 min. (1 h. in the case of *Diur*) and less than 60 min. (2 h in the case of *Diur*).

It is still considered an *Event* when, in a discontinuous way, values are verified out of the limits, but that are inferior to 10 min. and in a period of time of 30 min. maximum, since the sum of those is greater or equal to 10 min.

The definition of *Critical Event* is similar to the *Event*, but with different values. The times of 10 min. referred in the definition of *Events*, should be replaced by 1 hour, the 30 min. for 2 hours and when we refer to *Diur*, we consider 2 hours instead of 1 respectively.

A *Critical Event* can also be defined in some special situations, i.e., when the value of the analyzed parameter places among certain values.

We only can consider a new event, after a recovery period of 30 min. or more for *BP*, *SaO2* and *HR*, and of 2 hours or more for *Diur*, with values inside of the intervals. In *Critical Events*, it should be considered a period greater than 2 hours for *Diur* and greater than 60 minutes for the remaining ones.

## 4 DATA PREPARATION

A data preparation phase has been necessary to treat the wrong or omitted data. Besides, not all the variables were considered to generate the prediction models, as it is the case of the age, once it is already considered within *SAPSII* score.

Table 1 shows the variables that were considered in this study and their description. For modelling purposes, six new binary variables were created, based in the six *SOFA* values, according to the expression:

0	, if $SOFA_{Org} < 3$	(false, no organ failure)
1	, else	(true, organ dysfunction)

where  $org \in \{\text{Respiratory, Coagulation, Liver, Cardiovascular, Central Nervous, Renal}\}$  stands for the organ system.

Table 1: Variables Description

Variable	Description	Domain	
ID	Patient number		*
Respirat	Respiration System	{0,1,2,3,4}	**
Coagulat	Coagulation System	{0,1,2,3,4}	**
Liver	Liver System	{0,1,2,3,4}	**
Cardiova	Cardiovascular System	{0,1,2,3,4}	**
Cns	Central Nervous System	{0,1,2,3,4}	**
Renal	Renal System	{0,1,2,3,4}	**
Nrbpvt	Number of <i>BP Events/day</i>	{0,1,...,24}	
Nrbpcriv	Number of <i>BP Critical</i>	{0,1,...,10}	
Nrofhrv	Number of <i>HR Events/day</i>	{0,1,...,24}	
Nrofhr	Number of <i>HR Critical</i>	{0,1,...,10}	
Nrofo2ev	Number of <i>O2 Events/day</i>	{0,1,...,24}	
Nrofo2cr	Number of <i>O2 Critical</i>	{0,1,...,7}	
Nrofurev	Number of <i>Diur Events/day</i>	{0,1,...,24}	
Nrofur	Number of <i>Diur Critical</i>	{0,1,...,7}	
Admfrom	Admission From	{1,2,...,7}	***
Admtype	Admission Type	{1,2,3}	***
SapsII	Simplified Acute Physiology	{0,1,...,118}	***
Diagn	Diagnostic	{0,1}	***

- \* Not considered for the prediction models. They were only considered to build the clustering framework.
- \*\* Dependent variables.
- \*\*\* Variables just considered in the first day.

Once we intend to predict an organ failure in a certain day, based in the data of previous days, it was necessary to transform the database structure, in order to capture a temporary sequence of the variables (time series).

The variables *AdmFrom*, *AdmType*, *SapsII* and *Diagn* are obtained once (in the first day) but their values are considered in all situations of organ failure predictions.

For the construction of the various models, the *SOFA* values were not considered as input. Instead the number of *Events* and *Critical Events* registered for these two days were considered for the prediction in cause.

We just considered a temporary horizon of five days, because, in medical terms, the fifth day of stay in an ICU is considered a critical point in terms of the evolution of the patient's clinical state. The first day was not considered for prediction purposes, once the goal is to predict organ failure based on the data collected in the previous days. However, it is considered as input for all the other ones.

## 5 MODELS DEFINITIONS AND CONSTRUCTION

Making use of SPSS Clementine tool (<http://www.spss.com>), we submitted the database to a Kohonen Network (Kohonen, 1995), to segment it in three distinct groups. Later, it was fallen back upon the C5 (<http://www.rulequest.com>) algorithm in way to generate a model of decision trees to understand each one of those clusters. This way, we obtained 3 models for each one of the dependent variables, and for each one of the days of stay in the ICU (18 models for each day).

After having selected the most appropriated variables to the generation of models, a Kohonen Network was applied to the database, in order to create two additional variables, which correspond to the coordinates assigned at each record (identifying the cluster that it belongs). These coordinates make possible the partition of the patients into three clusters. Later, applying the C5 algorithm to each cluster is possible to generate the respective decision tree.

The validation of those models was made through a 10-fold cross validation method (Dubitzky et al, 2001). Finally, the achieved results were analysed by means of a confusion matrix, a matrix of size  $L \times L$ , where  $L$  denotes the number of possible classes (Kohavi et al, 1998),

## 6 ACHIEVED ACCURACIES

The confusion matrix is a common tool for classification analysis, this matrix is created by matching the predicted and actual values. When  $L = 2$ , there are four possibilities (Table 2): the number of correct positive - True Positive (TP), correct negative - True Negative (TN), incorrect positive - False Positive (FP); and incorrect negative - False Negative (FN) classifications.

Table 2: The  $2 \times 2$  confusion matrix

↓ actual / predicted →	Negative	Positive
Negative	TN	FP
Positive	FN	TP

From this table, three accuracy measures can be defined (Essex, 1995): the true Positive Rate (PR), also known as sensitivity, recall and Type II Error; the true Negative Rate (NR), also known as specificity, precision and Type I Error; and the Predictive Accuracy (PA), which gives an overall evaluation.

These metrics can be computed using the following equations:

$$PR = \frac{TP}{FN + TP} \times 100\%$$

$$NR = \frac{TN}{TN + FP} \times 100\%$$

$$PA = \frac{TN + TP}{TN + FP + FN + TP} \times 100\%$$

In the Table 3, we can see the results of the predicting models of the fifth day of stay.

Table 3: Results for the fifth day  
Cluster 0 Cluster 1 Cluster 2

	PR	92,45%	100,00%	97,37%
Respiratory System	NR	95,04%	100,00%	93,76%
	PA	94,43%	100,00%	94,61%
	PR	100,00%	100,00%	91,67%
Coagulation System	NR	98,51%	100,00%	99,44%
	PA	98,61%	100,00%	99,03%
	PR	100,00%	100,00%	88,24%
Liver System	NR	99,91%	100,00%	99,84%
	PA	99,91%	100,00%	99,54%
	PR	94,26%	100,00%	95,73%
Cardiovascular System	NR	93,07%	100,00%	96,62%
	PA	93,29%	100,00%	96,46%
	PR	93,88%	100,00%	88,98%
Central Nervous System	NR	95,51%	98,31%	95,98%
	PA	95,23%	98,44%	94,61%
	PR	92,31%	100,00%	98,28%
Renal System	NR	98,06%	100,00%	98,14%
	PA	97,53%	100,00%	98,15%

As we can see, the accuracies achieved are quite good, in some situations, we achieved the maximum values (100%). However, these are the clusters that have fewer patients, between 62 and 65 in a universe of 1845. Clusters 0 and 2 contain between 649 and 1134 patients.

These results were possible due to the approach adopted, as well as the use of misclassification costs that allow us to specify the relative importance of different kinds of prediction errors.

## 7 CLUSTERING FRAMEWORK

The Data Mining process created 72 prediction models plus a higher order classification model (based on a decision tree) that matches a patient to the respective prediction model.

To deal with this complexity and to make more explicit the relation patient vs. organ failure prediction model, was considered a visualization framework. In this framework, the prediction models are denoted by:

$$m(d, o, c, pa, se, sp)$$

where  $d \in \{2,3,4,5\}$  stands for the day of the stay,  $o \in \{\text{Renal, Central Nervous, Cardiovascular, Liver, Coagulation, Respiratory}\}$  stands for the organ, and  $c \in \{0,1,2\}$  for the cluster. The last arguments are the Predictive Accuracy (pa), the Sensitivity (se), and the Specificity (sp). These models may be organized in a cube that makes possible the graphical presentation of the patient course along the stay in ICU as we can see in the Figure 1.

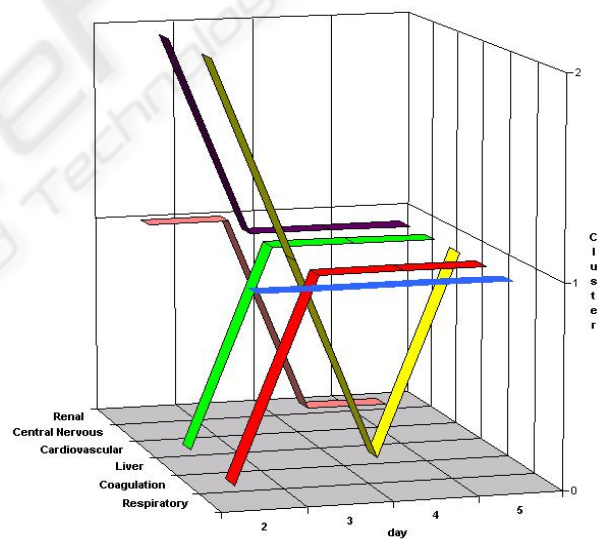


Figure 1: Clustering Framework

For a given patient we have a prediction model for each one of the 6 organs (o) indexed to the day of stay (d) and the correspondent cluster (c). Be noticed that, in the same day, the correspondent models of a particular patient may belong to different clusters.



In the example presented in the Figure 1, the prediction models for day 2 are given by:

- m(2, Renal, 1, 97.74, 90.91, 98.19);
- m(2, Central Nervous, 2, 89.10, 68.56, 93.72);
- m(2, Cardiovascular, 0, 84.95, 73.91, 87.68 );
- m(2, Liver, 2, 99.18, 74.19, 99.52);
- m(2, Coagulation, 0, 97.35, 71.70, 98.59);
- m(2, Respiratory, 1, 96.30, 90.00, 97.18).

As we can see, the prediction model for the Central Nervous system of this patient changed from the cluster two to cluster one, and the Predictive Accuracy, Sensitivity and Specificity changed also.

For perception convenience, this cube can be split into three layers, one for each cluster, where the validity of each model is represented by a grey scale (Table 4). The darkest tone revealed higher accuracies (as in the cluster 1).

The Figure 2 shows the Predictive Accuracies transformed in a four tones of grey scale.

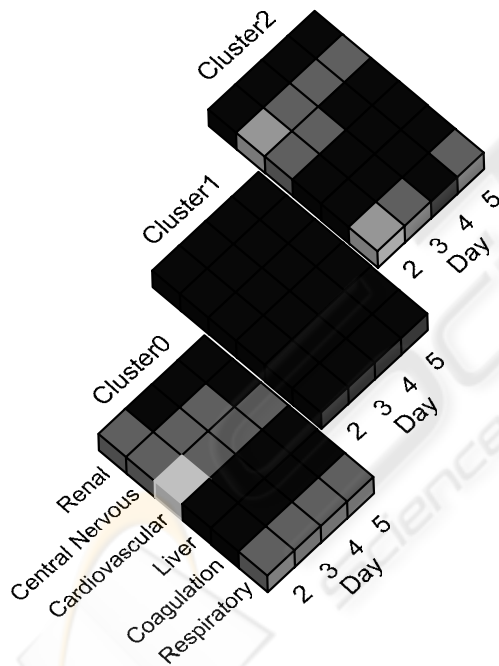


Figure 2: Predictive Accuracies

The cells correspondent to the day 5 represents the values indicated in Table 3. The rest represent the remaining Predictive Accuracies of the others days of stay in the ICU. This way we can, in a visual and easy form, distinguish the zones of interest for prediction.

Table 4: Predictive Accuracies scale

	From 95,01% to 100%
	From 90,01% to 95,00%
	From 85,01% to 90,00%
	Up to 85,00%

The database segmentation criteria, for the third day of stay related with the Renal system failure, can be visualized under the form a decision tree as following:

```

admtpe <= 2 (1329)
  admfrom <= 1 (922, 1.0) -> 0
  admfrom > 1 (407)
    admtpe <= 1 (309, 1.0) -> 0
    admtpe > 1 (98, 1.0) -> 1
  admtpe > 2 (1776)
    admfrom <= 1 (12, 1.0) -> 1
    admfrom > 1 (1764, 1.0) -> 2
    
```

In this case, the variables that determined the classification in three clusters were the Admission Type and Admission From. The values presented between parentheses stand for the support level and the confidence level, respectively.

As we can see, there is only one rule that respect to cluster 2, and two rules for each one of the clusters 0 and 1. If the admission type is medical, and the admission from is other then Operating Bloc, the patient will be in the cluster 2. This rule was applied to 1764 cases.

As we could see in the framework, the first day of stay was not considered, once it doesn't make sense to predict organ failure for this day, because the only data we have was collected in the same day.

Each of the 72 models referred in the framework correspond to decision trees generated by the C5 algorithm. Consider for example the decision tree that predicts the Central Nervous system for the fifth day, in the cluster 0:

```

sapsii <= 54 (905)
  nrofhrcr4 <= 1 (870)
    nrofo2ev1 <= 0 (687)
      .....
    nrofo2ev1 > 0 (183)
      diagn <= 0 (140)
        nrofhrev3 <= 0 (119)
          admfrom1 <= 3 (43, 1.0) -> 0
          admfrom1 > 3 (76)
            nrofo2ev1 <= 4 (70)
            
```

As we could see in the tree, we could say that if a patient has a *SAPSII* score less than 54, equal or less than one critical events of heart rate in the fourth day, at least one event of O<sub>2</sub> in the first day, with a diagnostic non operative, no events of heart rate in the third day, and with admission Source of Operating Bloc, Recovery Room or Emergency Room, the central nervous system will not be in failure.

This is the kind of information that is really important in an ICU environment in a decision support context.

## 8 CONCLUSIONS AND FURTHER WORK

In this study, we presented a clustering framework, with the purpose of identifying and applying the model generated for the cluster in which a patient frames to, according to his characteristics. The majority of the models revealed high accuracies, which is very useful in a decision support context. The gains of this approach can be summarized as follows:

- A matrix to dispose and explore the models;
- A system to apply the models to a particular patient through a process based on three indexes: the day, the organ and the cluster;
- An explicit way to declare the best and the worst predictive zones (models) based on assessment metrics such the accuracy, the specificity and sensitivity. The doctors know exactly what is the value and usability of the models and its prediction.
- An alternative or complementary formalism of knowledge representation and visualization for decision support.

Further work will include the study of meta-learning techniques in order to maintain the matrix in dynamic environments (as the ICU), as well the graphic technologies to support the visualization and interaction with the framework, enabling the construction of intelligent decision support systems.

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