



Article

# Early Prediction of Seven-Day Mortality in Intensive Care Unit Using a Machine Learning Model: Results from the SPIN-UTI Project

Martina Barchitta <sup>1,2</sup>, Andrea Maugeri <sup>1,2</sup>, Giuliana Favara <sup>1</sup>, Paolo Marco Riela <sup>3</sup>, Giovanni Gallo <sup>3</sup>, Ida Mura <sup>2,4</sup> and Antonella Agodi <sup>1,2,\*,†</sup> on behalf of the SPIN-UTI network

- Department of Medical and Surgical Sciences and Advanced Technologies "GF Ingrassia", University of Catania, 95123 Catania, Italy; martina.barchitta@unict.it (M.B.); andrea.maugeri@unict.it (A.M.); giuliana.favara@unict.it (G.F.)
- GISIO-SItI—Italian Study Group of Hospital Hygiene—Italian Society of Hygiene, Preventive Medicine and Public Health, 00144 Roma, Italy; idamura@uniss.it
- Department of Mathematics and Informatics, University of Catania, 95123 Catania, Italy; paolo.riela@unict.it (P.M.R.); gallo@dmi.unict.it (G.G.);
- Department of Biomedical Sciences, University of Sassari, 07100 Sassari, Italy
- \* Correspondence: agodia@unict.it
- † Collaborators of the SPIN-UTI network are indicated in the Acknowledgment section.

Abstract: Patients in intensive care units (ICUs) were at higher risk of worsen prognosis and mortality. Here, we aimed to evaluate the ability of the Simplified Acute Physiology Score (SAPS II) to predict the risk of 7-day mortality, and to test a machine learning algorithm which combines the SAPS II with additional patients' characteristics at ICU admission. We used data from the "Italian Nosocomial Infections Surveillance in Intensive Care Units" network. Support Vector Machines (SVM) algorithm was used to classify 3782 patients according to sex, patient's origin, type of ICU admission, nonsurgical treatment for acute coronary disease, surgical intervention, SAPS II, presence of invasive devices, trauma, impaired immunity, antibiotic therapy and onset of HAI. The accuracy of SAPS II for predicting patients who died from those who did not was 69.3%, with an Area Under the Curve (AUC) of 0.678. Using the SVM algorithm, instead, we achieved an accuracy of 83.5% and AUC of 0.896. Notably, SAPS II was the variable that weighted more on the model and its removal resulted in an AUC of 0.653 and an accuracy of 68.4%. Overall, these findings suggest the present SVM model as a useful tool to early predict patients at higher risk of death at ICU admission.

**Keywords:** healthcare-associated infections; machine learning; intensive care unit; risk prediction; mortality; adverse outcomes



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### 1. Introduction

Healthcare-associated infections (HAIs) are the most frequent adverse outcome occurring when patients stay in hospital wards, especially in intensive care units (ICUs) [1–9]. Due to their impact on patients' morbidity and mortality, as well as on antimicrobial resistance and assistance healthcare costs, these infections represent a major concern for public health [5,10–15]. As reported by the World Health Organization (WHO), the global burden of HAIs raises up to 15% among all hospitalized patients, with a proportion that achieves more than 30% in those who stay in ICUs [16–18].

Indeed, patients admitted to ICUs generally had a worse clinical prognosis, including prolonged hospital stays, sepsis and mortality [19]. Particularly, mortality in ICUs is two time higher among infected patients than those not infected [6,8]. This in turn depends on several patients' factor such as the use of invasive procedures, their severity, type of infection, therapy, and microorganisms' characteristics, including clonal spread [19–22]. Nowadays, several early warning scores have been proposed as a helpful instrument to monitor

patients' clinical deterioration and disease severity during their stay in ICUs [23–27]. In clinical practice, the Simplified Acute Physiology Score (SAPS) II represents the most commonly used score. Specifically, it is able to predict patients' prognosis and to estimate their risk of HAIs, sepsis and dying, according to 17 physiological variables at ICU admission [28–37].

With this in mind, the complexity of HAI burden suggested the need of novel approaches aimed at early identifying patients at higher risk of adverse events in ICU [38]. Indeed, the prediction of patients at higher risk of mortality in ICU play a key role in improving patients' survival and in implementing their management [39]. Although several traditional statistical approaches are widely used in clinical practice, modern machine learning models showed more accurate results in the early identification of patients who are more likely to die during their stay in ICU, considering different sets of risk factors [9,39–42]. For instance, support vector machine (SVM) algorithm is a supervised machine learning method with a high performance in discriminating two different classes of events, especially in many medical fields where a large amount of patient' variables are collected [43,44].

In the present study, we aimed to identify and predict patients at higher risk of dying, considering their clinical and pathological characteristics at ICU admission. The primary purpose of this study is to evaluate the ability of the SAPS II to predict the risk of death after 7 days from their admission to ICU. The secondary purpose is to develop and test a machine learning algorithm, which combines the SAPS II with additional patients' characteristics, to further improve the predicting performance. Here, we used data collected during the seven editions of the "Italian Nosocomial Infections Surveillance in Intensive Care Units" (SPIN-UTI) network, an ongoing surveillance system established by the Italian Study Group of Hospital Hygiene (GISIO) of the Italian Society of Hygiene, Preventive Medicine and Public Health (SItI) [45–51]. From 2006 to date, the SPIN-UTI network has collected data of nearly 20,000 patients, 4300 infections and 5300 microorganisms, which helped to feed the European surveillance network with data of HAIs in Italian ICUs [45–51].

#### 2. Materials and Methods

#### 2.1. The SPIN-UTI Project

Here, we used data collected during the seven editions of the SPIN-UTI project, which were conducted from 2016 to 2019 in Italian ICUs, using the ECDC protocol [52]. The protocols used and characteristics of 20,060 patients surveyed were fully described elsewhere [45–51]. In general, the SPIN-UTI project prospectively surveys patients staying in ICU for more than two days and collects data at hospital, ICU and patient level. By contrast, patients who stay in ICU less than two days are excluded a priori. The reason for their exclusion is because the primary outcome of the SPIN-UTI project is the incidence of HAIs, which by definition develop after two days of ICU stay. The study was approved by the Ethics Committee "Catania 1", Catania, Italy (protocol numbers 111/2018/PO and 295/2019/EMPO).

#### 2.2. Definition of SAPS II and Other Predictors

In the present work, SAPS II at ICU admission was initially used as the main predictor. As previously described [35], the computation of SAPS II included the following components: Age; heart rate; systolic blood pressure; temperature; Glasgow Coma Scale; continuous positive airway pressure; PaO2; FiO2; urine output; blood urea nitrogen; sodium; potassium; bicarbonate; bilirubin; white Blood Cell; chronic diseases; type of admission. Each component was assessed within 24 h from ICU admission and the worst value was recorded. The total SAPS II was finally computed as the sum of weighted values for each component [35]. The SPIN-UTI project also collected information on patients who underwent non-surgical treatment for signs and symptoms related to the acute coronary syndrome. Moreover, we defined admission with trauma those resulting from blunt or penetrating traumatic injury to the patient, with or without surgical intervention.

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Instead, impaired immunity was defined as an impairment due to treatment, diseases or <500 PMN/mm3. Finally, we also collected if any antibiotic therapy was administered in the 48 h preceding ICU admission and/or during the first two days of ICU stay. The occurrence of HAI was defined according to a set of clinical and laboratory criteria that are fully described in the ECDC protocol [53].

#### 2.3. Dataset of "Real" Records

We first worked only on patients with a complete assessment of the following information: Sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, SAPS II, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission and onset of HAI. The primary outcome of the current analysis was mortality within seven days from ICU admission. Accordingly, the current analysis included: (1) Patients who stayed in ICU for at least seven days and (2) those who died within two to seven days after ICU admission. By contrast, patients who stayed in ICU for less than two days, those who were discharged prior to 7 days, and those who died with the first two days were excluded. The matrix of missing values and the selection of "real" records are showed in Figures S1 and S2. Specifically, this dataset of "real" records consisted on a total of 3782 patients with complete data and meeting the inclusion criteria. The dataset described above was used for traditional statistical analyses and as test set for the machine learning algorithm.

#### 2.4. Dataset of Synthetic Records

The missing data is arguably the most common problem encountered by machine learning professionals when analyzing real-world data. The same applies for the SPIN-UTI project, where a lot of records present missing information. For instance, in the current analysis there were 61% (n = 12,237) of records with missing data for variables defined in the paragraph above. As many statistical models and machine learning algorithms rely on complete datasets, it is key to handle the missing data appropriately. Moreover, machine learning algorithms generally requires large datasets to be trained. For these reasons, we created a dataset of synthetic records that was used as training set for the machine learning algorithm. Accordingly, we first imputed missing data from incomplete records of the original dataset, using the K-Nearest Neighbor (K-NN) imputation method described by Malarvizhi and Thanamani [54]. Specifically, we applied the K-NN algorithm considering the Euclidean distance in the feature space for non-binary variables and the Jaccard distance for dichotomic variables. After applying two cycles of K-NN imputation to the two classes of data (i.e., alive or died patients), we recovered 3258 records. More details on methods used for data imputation are reported in the Supplementary Materials. Moreover, synthetic data were generated to balance the two classes of died and alive patients, using the Synthetic Minority Over-sampling Technique (SMOTE). In contrast to under-sampling, the oversampling approach is useful to increase the cardinality of the minority class by duplicating records. In general, the SMOTE first selects random records from the minority class population, then identifies the K-NN, and finally creates synthetic records from the random ones and the randomly selected K-NN [55]. After applying the SMOTE to 3782 "real" records, we obtained 1131 synthetic records for the class of died patients. Given that, the dataset of synthetic records, which was used as the training set, included a total of 4389 records. To confirm the goodness of the training set, we compared the distributions of primary outcome and exposure variables with those obtained from the test set (Table S1 and Figures S3-S5).

#### 2.5. Statistical Analysis

All variables of the "real" dataset were described according to their type and skewness using descriptive statistics (frequencies and percentages [%] or median and interquartile range (IQR)). In an epidemiological and descriptive point of view, we compared these

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variables between dead and alive patients using the Mann-Whitney U test for quantitative variables and the Chi-Squared test and Chi-Squared for trend test for qualitative variables. We first used a logistic regression model to evaluate the association of SAPS II (continuous) with death. Next, we applied a logistic regression model, also including sex (dichotomous), patient's origin (categorical: Other ward/healthcare facility, community), type of ICU admission (categorical: Medical, surgical), non-surgical treatment for acute coronary disease (dichotomous), surgical intervention (dichotomous), presence of invasive devices at ICU admission (three dichotomous variables for urinary catheter, intubation and central venous catheter, respectively), trauma (dichotomous), impaired immunity (dichotomous), antibiotic therapy in 48 h before or after ICU admission (dichotomous). We also used Receiver Operating Characteristic (ROC) curves to assess the ability of the logistic regression models to accurately identify patients who dead from those who did not. Results were reported in terms of Area Under the Curve (AUC) and 95% Confidence Interval (95% CI). With respect to the model on SAPS II alone, we identified the best cut-off value which maximized the Youden Index. For the best cut-off value, sensitivity and specificity with their 95% CI were calculated. All tests were performed at a significance level  $\alpha = 0.05$  and statistical analysis was conducted using SPSS v.25 (IBM Corp., Armonk, NY, USA).

#### 2.6. Machine Learning Algorithm

We next compared the predictive performance of 7-day mortality between logistic regression model and a machine learning algorithm. Specifically, the algorithm combined SAPS II with the following variables collected at ICU admission: Sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of intubation, presence of urinary catheter, presence of central vascular catheter; trauma, impaired immunity, and antibiotic therapy in 48 h before or after ICU admission. For the current analysis, we chosen the supervised SVM algorithm as modelling tool, which can be used for classification—especially for binary classification—and regression problems. However, our dataset was not linearly separable, not allowing to satisfy all the constraints of SVM [44]. For this reason, we used a non-linear Kernel function (i.e., the Gaussian Kernel, also called as Radial basis function Kernel, RBF). Slack variables with penalty were also introduced to satisfy all the constraints in the minimization problem of SVM [44]. More details on the machine learning algorithm are reported in the Supplementary Materials. The SVM model was trained on the training set composed by synthetic records, and then tested on the test set made of "real" records. Since patients who developed HAIs during their ICU stay are generally at higher risk of death, we also tested the SVM model on those who did not acquire HAIs within seven days from ICU admission. We also assessed the predictive performance of a SVM model, which included all variables collected at ICU admission except of SAPS II. Results are reported in terms of AUC, accuracy, sensitivity, and specificity with their 95% CI. The analyses were performed using Python and Support Vector Classification (SVC) from Sklearn 0.22.1.

#### 3. Results

#### 3.1. Characteristics of the Dataset of "Real" Records

The current analysis included 3782 SPIN-UTI participants without missing data (60.2% males), surveyed from 2006 to 2019. In this subsample, the median age was 70.0 years (IQR = 20) and median SAPS II score at admission was 49 (IQR = 27). Overall, 70.9% came from other wards/hospitals and 56.9%, reported a medical type of ICU admission. In particular, 4.7% and 11.4% of patients reported trauma and/or impaired immunity, respectively. Patients who underwent antibiotic therapy, surgical intervention, or non-surgical treatment for acute coronary disease were 62.6%, 34.8%, and 9.0%, respectively. With respect to invasive devices, the presence of urinary catheter, intubation and central venous catheter was reported in 77.0%, 62.4%, and 40.5% patients, respectively. Table 1 compares characteristics of patients who died (n = 875; 23.1%) within seven day from ICU admission with those who were still alive (n = 2907; 76.9%). Specifically, patients who died were older,

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more likely men, and with a higher SAPS II than those who did not die. Moreover, they were also more likely to come from other ward/healthcare facility and to report a medical type of ICU admission than those alive. The first group also consisted more of patients who reported impaired immunity and less traumatic events. Instead, no differences were evident for surgical intervention, non-surgical treatment for acute coronary disease, antibiotic therapy on admission and presence of invasive devices at ICU admission.

<b>Table 1.</b> Characteristics of	patients with comp	olete data according	to their outcome status.

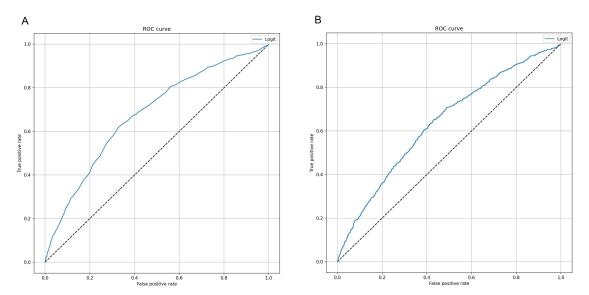
Characteristics	Patients ( <i>n</i> = 3782)	Died Patients (n = 875)	Alive Patients ( <i>n</i> = 2907)	<i>p</i> -Value
Age, years	70.0 (20.0)	74.0 (17.0)	69.0 (21.0)	< 0.001
Sex (% men)	60.2%	55.0%	61.7%	< 0.001
Patient's origin				
Other ward/healthcare facility	70.9%	70.1%	71.1%	< 0.001
Community	29.1%	29.9%	28.9%	
SAPS II at admission	49.0 (27.0)	59.0 (27.0)	46.0 (25.0)	< 0.001
Type of ICU admission				
Medical	56.9%	59.3%	56.1%	< 0.001
Surgical	43.1%	40.7%	43.9%	
Trauma	4.7%	2.4%	5.4%	< 0.001
Impaired immunity	11.4%	15.0%	10.3%	< 0.001
Non-surgical treatment for acute coronary disease	9.0%	10.2%	8.7%	0.174
Surgical intervention	34.8%	32.5%	35.5%	0.306
Antibiotic therapy in 48 h before or after ICU admission	62.6%	62.2%	62.8%	0.744
Presence of urinary catheter at ICU admission	77.0%	75.9%	77.4%	0.351
Presence of intubation at ICU admission	62.4%	61.7%	62.6%	0.646
Presence of central venous catheter at ICU admission	40.5%	38.5%	41.0%	0.182

<sup>\*</sup> Results are reported as median (interquartile range) for continuous variables, or percentage (%) for categorical variables. Statistical analyses were performed using the Mann–Whitney, the Chi-squared test, or the Chi-squared test for trend.

#### 3.2. Applying Logistic Regression Models to Predict the Risk of 7-Day Mortality

We first applied a logistic regression model on the dataset of "real" records, using SAPS II as the independent and 7-day mortality as the dependent variable. Accordingly, Figure 1A illustrates the accuracy of SAPS II for predicting the risk of 7-day mortality for all patients admitted in ICU. We noted that SAPS II was able to discriminate patients who died from those who did not, with AUC of 0.678 (95% CI = 0.657–0.700; p < 0.001) and accuracy of 69.3% (95% CI = 67.8–70.8%). The coordinates of the ROC curve are reported in Table S2. Specifically, the best cut-off value of SAPS II, which maximized the Youden index, was 54.5. The application of this value resulted in sensitivity of 61.9% (95% CI = 60.4–63.4%) and specificity of 67.1% (95% CI = 65.6–68.7%). We further applied a logistic regression model, which combined SAPS II with additional patients' characteristics collected at ICU admission. However, as indicated in Figure 1B, both AUC and accuracy of this model remained moderate (AUC = 0.637; 95% CI = 0.616–0.659; Accuracy = 65.2%; 95% CI = 63.7–66.7%). In line, sensitivity and specificity for death were 49.0% (95% CI = 47.5–50.5%), and 70.0% (95% CI = 68.5–71.5%), respectively.

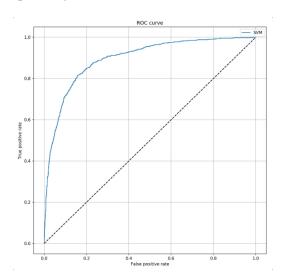
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**Figure 1.** Receiver Operating Characteristic (ROC) curves of logistic regression models to predict 7-day mortality. (**A**) This curve shows the predictive performance of a logistic regression model using Simplified Acute Physiology Score (SAPS II) alone. (**B**) This curve shows the predictive performance of a logistic regression model including sex, patient's origin, type of intensive care unit (ICU) admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission.

#### 3.3. The SVM Algorithm Improved the Prediction of Patients Who Died

Next, we aimed to develop a machine learning algorithm, which could improve the prediction of 7-day mortality in ICU. To do that, we used the SVM algorithm by combining SAPS II with other characteristics collected at ICU admission. Interestingly, the ROC curve of SVM predictive model (Figure 2) achieved an AUC of 0.896 (95% CI = 0.881–0.910; p < 0.001), with an accuracy of 83.5% (95% CI = 82.4–84.7%). In line, sensitivity and specificity were 81.0% (95% CI = 79.9–82.1%) and 84.0% (95% CI = 82.9–85.1%), respectively.



**Figure 2.** ROC curve of the Support Vector Machines (SVM) algorithm to predict 7-day mortality. This curve shows the predictive performance of the SVM algorithm including SAPS II, sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission. The parameters applied to the SVM algorithm were C = 2 and  $\gamma = 0.3$ .

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## 3.4. The SVM Algorithm Maintained Its Predictive Ability among Patients Who Did Not Develop HAIs

We also tested the predictive ability of the SVM classifier among patients who did not develop HAIs within 7 days from ICU admission. To do that, we removed 520 patients with at least one HAIs from the test set. Interestingly, the model did not depend on the onset of HAI, since both AUC (0.903; 95% Confidence Interval = 0.881-0.912; p < 0.001) and accuracy (83.8%; 95% CI = 82.6–85.0%) remained stable (Figure 3). In line, sensitivity and specificity were comparable to those obtained in the overall analysis (82.0%; 95% CI = 80.8–83.2%; and 84.0%; 95% CI = 82.8–85.2%, respectively).

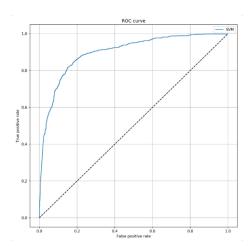
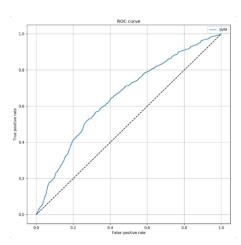


Figure 3. ROC curve of the SVM algorithm to predict 7-day mortality, by excluding infected patients. This curve shows the predictive performance of the SVM algorithm including SAPS II, sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission. The parameters applied to the SVM algorithm were C = 2 and  $\gamma = 0.4$ .

#### 3.5. The Predictivie Performance of the SVM Model by Removing SAPS II

The Shapley plot reported in Figure S6 shows the contribution of each predictors to the SVM model output. Since SAPS II was the predictor with the highest importance, we finally evaluated the predictive performance of the classifier after removing SAPS II. Interestingly, the SVM model without SAPS II led to an AUC of 0.653 (95% CI = 0.632–0.675; p < 0.001), with an accuracy of 68.4% (95% CI = 66.9–69.8%) on the test set (Figure 4). Accordingly, sensitivity and specificity decreased to 32.0% (95% CI = 30.5–33.5%) and 74.0% (95% CI = 72.5–75.5%), respectively.



**Figure 4.** ROC curve of SVM algorithm predicting 7-day mortality, by excluding SAPS II score. This curve shows the predictive performance of the SVM algorithm including sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission. The parameters applied to the SVM algorithm were C = 2 and  $\gamma = 0.1$ .

#### 4. Discussion

In past years, numerous early warning scores were developed and employed to monitor and predict patients' conditions and severity, as well as their adverse events in healthcare facilities [23–27]. Among these, SAPS II still represents one of the most widely used tool to estimate patients' risk of death and other adverse outcomes [28–37]. In view of these considerations, we first aimed to evaluate the accuracy of SAPS II alone to identify patients who died within seven days from their admission in ICUs, using a large dataset from the SPIN-UTI project. Although AUC obtained was statistically significant, the low accuracy of nearly 69% discouraged the routinely application of SAPS II to achieve this purpose.

Applying novel predictive algorithms, however, could be important to ameliorate patient's safety and management in clinical practice, especially in the ICU setting. Thus, we hypothesized that combining SAPS II with other variables collected at ICU admission could improve the prediction of 7-day mortality [56–60]. Indeed, it is now well-established that machine learning algorithms could overcome the limitations of traditional existing tools, also allowing early prediction of mortality [9,23–27,40–42,56,61]. To do that, we developed a SVM model, which combined SAPS II with the following patients' characteristics at ICU admission: Sex, patient's origin, type of ICU admission, non-surgical treatment for acute coronary disease, surgical intervention, presence of invasive devices at ICU admission, trauma, impaired immunity, antibiotic therapy in 48 h before or after ICU admission and presence of infection in seven days of ICU stay. The model exhibited an AUC of 0.90 with an accuracy of 83.5% on the test set. Interestingly, its predictive performance was higher than SAPS II alone and even than a logistic regression model including additional patients' characteristics collected at ICU admission. We also demonstrated that the performance for predicting 7-day mortality was also similar in only patients who did not acquired HAIs during their hospitalization. However, other early warning scores—for example the Acute Physiology and Chronic Health Evaluation II (APACHE II) and the Mortality Probability Model (MPM)—can assess disease severity at ICU admission and predict the risk of death, with results that were similar to SAPS II [32,62]. Thus, it will be interesting to validate our model in countries where other early warning scores are most commonly used.

Overall, our findings underlined the potentially crucial role of machine learning algorithms in many public health issues, providing clinicians with better diagnostic tools and improving medical care in the next future. The promising benefits of applying machine learning on healthcare quality rely on the opportunity of making prevention and diagnosis as early as possible, in a context of precision medicine applicable to all set-

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tings. We recognize that findings from machine learning are frequently considered as a sort of "black box" for clinicians [63,64]. However, in our study, we aimed to use only predictors that might be related to severity and mortality of ICU patients on the basis of previous experience and expert opinion. This has certainly resulted in a more interpretable algorithm for clinicians and public health professionals [63,64]. Indeed, applying supervised machine learning algorithms could support medical professionals in parameter optimization, clustering and classification problems [21,64–70]. Accordingly, these algorithms—if properly applied—could overcome limitations of existing traditional early warning scores [9,23–27,40–42,56,61]. For instance, a previous study developed a machine learning algorithm based on vital signs of ICU patients, such as heart and respiration rate, oxygen saturation and blood pressure. In particular, the algorithm was able to predict mortality in ICU with an accuracy of 91.6% [71]. Our findings—together with those from other research groups—lay the foundations to develop automated and real-time tools able to identify patients who need more attention because of their high risk of death.

Our model had several strengths, including the better ability of predicting 7-day mortality if compared with an early warning score as SAPS II. Moreover, our model was trained and tested on large datasets obtained through patient-based surveillance, structured and standardized according to ECDC protocol. This should allow validation and comparison with other European countries. On the other hand, however, we cannot completely exclude historical bias due to a 14-year period of data collection. Beyond that, there were other considerations to keep in mind when interpreting our results. The first one was that our findings confirmed the importance of developing and validating early warning scores to predict the risk of death and other adverse outcomes in ICUs and other wards. Indeed, although we used several variables collected at ICU admission, the removal of SAPS II from the model significantly reduced the predictive performance. The second consideration was that machine learning requires a lot of variables and records, which are not always available in each healthcare settings. Although we used variables that can be easily collectible at ICU admission (e.g., patients' demographic, origin, and type of admission, medical history, and disease severity), a lot of patients had both structural missing and missing at random data. While the first type of missing data could be easily managed by improving their collection in the next SPIN-UTI editions (e.g., making them mandatory), those that miss at random will continue to exist. This still remains a common issue encountered when analyzing real-world data. For this reason, we cannot completely exclude potential bias related to the high proportion of missing data. To partially manage missing data, we adopted a dual approach to generate synthetic records from those incomplete. Indeed, we created a dataset of synthetic records that was used as the training set for our machine learning algorithm. However, while it remains preferable using real data to train the algorithms, the comparison between training and test sets showed no significant differences. Finally, we recognize that our model was based on non-temporal variables related to patients' characteristics as predictors and 7-day mortality as a sort of surrogate outcome for identifying patients who needed more attention in ICU. Thus, it will be our task to consider clinical outcomes prior to death and a time-series approach (e.g., survival analysis) for improving our model.

#### 5. Conclusions

With these considerations in mind, to the best of our knowledge, our study is the first employing the SVM algorithm to discriminate patients who died within seven days from their ICU admission from those who did not. The model showed good predictive performance, even though improvable. For this reason, further studies should be encouraged to develop and validate risk prediction models, which could help to predict adverse outcome as early as possible, and to improve patient care globally.

**Supplementary Materials:** The following are available online at https://www.mdpi.com/2077-038 3/10/5/992/s1, Figure S1: Matrix of missing values; Figure S2: Selection of records with complete data satisfying inclusion criteria; Figure S3: Comparison of Age (A) and SAPS II score (B) distributions between Training and Test sets, Figure S4: Comparison of dichotomous variables between Training

and Test sets, Figure S5: Comparison of categorical variables between training and test datasets; Figure S6: Shapley plot showing the contribution of each predictor to the SVM model output; Table S1: Composition of training and test datasets; Table S2: Coordinates of the ROC curve of logistic regression model with SAPS II alone; Supplementary Methods for Data Imputation and Support Vector Machine model.

**Author Contributions:** Conceptualization: M.B., A.M. and A.A.; Methodology: M.B., A.M., G.F., P.M.R., G.G., and I.M.; Formal analysis and investigation: A.M., G.F., and P.M.R.; Writing—original draft preparation: A.M., G.F., G.G., M.B., and P.M.R.; Writing—review and editing: All the Authors; Supervision: A.A. All authors have read and agreed to the published version of the manuscript.

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**Institutional Review Board Statement:** The study was approved by the Ethics Committee "Catania 1", Catania, Italy (protocol numbers 111/2018/PO and 295/2019/EMPO).

**Informed Consent Statement:** Patient consent was waived due to the specific study design, data collection, anonymization procedure, and reporting of aggregate data.

**Data Availability Statement:** The data presented in this study are available on request from the corresponding author.

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Aida Bianco, Maria Pavia (Azienda Ospedaliera Universitaria "Mater Domini", Catanzaro); Marcello Pasculli, Cesare Vittori (Azienda Ospedaliera Universitaria Senese, Siena); Giovanni Battista Orsi (Azienda Ospedaliera Sant'Andrea, Roma); Cristina Arrigoni (IRCCS Policlinico San Matteo di Pavia); Maria Patrizia Olori (Stabilimento Ospedaliero "C. e G. Mazzoni", Ascoli Piceno); Massimo Antonelli, Patrizia Laurenti, Riccardo Maviglia (Fondazione Policlinico Gemelli, Roma); Franco Ingala, Carmela Conte, Salvatore Russo, Laura Condorelli (ASP 8, Siracusa); Patrizia Farruggia (Presidio Ospedaliero Unico Aziendale dell'Azienda USL di Bologna); Cristina Maria Luisa (Ente Ospedaliero Ospedali Galliera di Genova); Italia Galassi (Presidio Ospedaliero "SS. Filippo e Nicola" di Avezzano).

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