# Boosting framework via clinical monitoring data to predict the depth of anesthesia

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#### Abstract.

BACKGROUND: Prediction of the depth of anesthesia is a difficult job in the biomedical field.

**OBJECTIVE:** This study aimed to build a boosting-based prediction model to predict the depth of anesthesia based on four clinical monitoring data.

**METHODS:** Boosting is a framework algorithm that is used to train a series of weak learners into strong learners by assigning different weights according to their classification accuracy. The input of the boosting-based prediction model included four types of clinical monitoring data: electromyography, end-tidal carbon dioxide partial pressure, remifentanil dosage, and flow rate. The output was the depth of anesthesia.

**RESULTS:** The boosting framework model built in this study achieved higher prediction accuracy and a lower discrete degree in predicting the depth of anesthesia compared with the DT-, KNN-, and SVM-based models.

**CONCLUSIONS:** The boosting framework was used to set up a prediction model to predict the depth of anesthesia based on four clinical monitoring data. In the experiments, the boosting framework model of this study achieved higher prediction accuracy and a lower discrete degree. This model will be useful in predicting the depth of anesthesia.

Keywords: Boosting framework, depth of anesthesia, prediction model

# 1. Introduction

Prediction of the depth of anesthesia is a difficult job in the biomedical field. The depth of anesthesia is related to the life safety and rehabilitation of patients in operation [1]. Shallow anesthesia can lead to intraoperative awareness, and deep anesthesia can make it difficult to meet the needs of the operation. Therefore, a prediction model to predict the depth of anesthesia needs to be proposed.

Many factors affect the depth of anesthesia. Electromyogram (EMG) records the bioelectrical imaging of muscle fibers. It is important to evaluate human activity in the anesthesia system. The monitoring of end-tidal carbon dioxide partial pressure (ETCO2) reflects lung ventilation and pulmonary blood flow. Ventilation is adjusted according to ETCO2 measurements during anesthesia. Remifentanil is a common anesthetic, and the dosage and flow rate of remifentanil also affect the depth of anesthesia. In this study, EMG, ETCO2, remifentanil dosage, and flow rate were selected as the main factors affecting the depth of

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anesthesia. A prediction model was proposed to predict the depth of anesthesia. Supervised learning was applied to the field of anesthesia to mine data or build a model to predict the depth of anesthesia [2].

Decision tree (DT) [3] was used to study the anesthesia depth index [4]. Support vector machine (SVM) [5] methods were used to monitor the level of hypnosis based on a series of striking features [6]. The K-nearest neighbor (KNN) [7] was used in clinical practice to classify states of anesthesia as awake, mild, moderate, and deep [8]. However, these methods are sensitive to parameters and easy to overfit, and have low precision. Therefore, these methods might not meet the requirements in specific scenarios. Boosting is an ensemble learning algorithm that combines a set of basic classifiers with different weights to become a strong classifier [9]. It is not easy to overfit. Boosting is highly suitable for building a prediction model for the depth of anesthesia.

In this study, the boosting framework was used to build a prediction model to predict the depth of anesthesia based on four clinical monitoring data. Boosting is a framework algorithm that usually trains a series of weak learners into strong learners by giving different weights according to their classification accuracy. The input of the proposed prediction model included four clinical monitoring data EMG, ETCO2, remifentanil dosage, and flow rate. The output was the depth of anesthesia. DT, SVM, and KNN also were used to build the proposed prediction models for the depth of anesthesia. The results of this study showed that the boosting framework model had a lower mean-squared error (MSE), mean absolute error (MAE), and a higher R-square (R2) compared with those of the DT-, KNN-, and SVM-based models.

The contributions of this study are as follows:

- 1) EMG, ETCO2, remifentanil dosage, and flow rate were selected as indicators to predict the depth of anesthesia more accurately.
- 2) The core of this study was that a boosting-based prediction model to predict the depth of anesthesia was built based on four clinical monitoring data.
- 3) Several popular machine learning algorithms, such as DT, KNN, and SVM, were used to build prediction models to compare performances in predicting the depth of anesthesia.
- 4) In the experiments, the boosting framework model of this study achieved higher prediction accuracy and a lower discrete degree in predicting the depth of anesthesia compared with the DT-, KNN-, and SVM-based models.

The structure of this study was as follows: Section 2 presents several popular prediction modeling methods; Section 3 presents the proposed methods; Section 4 presents the experimental design, the experimental results, and the discussions; and Section 5 presents the conclusion.

## 2. Method

# 2.1. Decision tree

DT is a tree structure in which each internal node represents a test on an attribute, each branch represents a test output, and each leaf node represents a category. In machine learning, DT is a predictive model. It represents a mapping between object attributes and object values. Entropy represents how chaotic the system is.

The expression for entropy is defined by:

$$H(x) = -\sum p_i * \log p_i \tag{1}$$

where p is the probability. As p approaches 1 (high probability), H approaches 0 (low entropy).

### 2.2. K-nearest neighbors

According to KNNs, every sample can be represented by its closest k neighbors. The idea of this method is very simple and intuitive: a sample belongs to a category if most of the k similar samples in the feature space (i.e., the closest neighbors in the feature space) belong to that category. This method only determines the classification of the samples according to the category of the nearest one or several samples.

Distance can be measured by several ways; one of the most commonly used is the Euclidean distance, expressed as:

$$D(x,y) = \sqrt{(x_1 - y_1)^2 + (x_2 - y_2)^2 + \dots (x_n - y_n)^2} = \sqrt{\sum_{i=1}^n (x_i - y_i)^2}$$
(2)

# 2.3. Support vector machine

For regression problem, given the training data  $D = \{(x_1, y_1), (x_2, y_2), \dots, (x_m, y_m)\}$  a regression model can be built to make the predicted and accurate values as close as possible.

For the sample (x, y), the traditional regression model usually calculates the loss directly based on the difference between the model output f(x) and the real output y. The loss is 0 if and only if f(x) is the same as y. On the contrary, SVM assumes that one can tolerate at most  $\varepsilon$  errors between f(x) and y. The loss is calculated only if the absolute value of the difference between f(x) and y is greater than  $\varepsilon$ .

The learned regression model of SVM is defined as:

$$f(x) = \omega \wedge T * x + b \tag{3}$$

where  $\omega$  is the normal vector of the decision surface and determines the direction of the decision surface, and b determines the position of the decision surface.

## 3. Proposed methods

A framework algorithm called boosting was designed to obtain a subset of samples by operating on the sample set; subsequently, a series of base classifiers were trained on the subset with a weak classification algorithm over time by tweaking error measures. These models were trained sequentially, and not in parallel. Weak learners were those with an error rate of slightly less than 50%. A series of trained weak learners were combined, and each weak learner  $f_i(x)$  had a corresponding weight  $\alpha_i$ .

Let  $\omega_t(n) = \frac{1}{N}$ , where N is the number of training samples and M is the selected number of iterations. For i in M:

a. Select  $f_{i}(x)$ , the weak learner, such that  $\varepsilon_{i}$  is minimized

$$f(x) = \omega \wedge T * x + b \tag{4}$$

b. Compute the weight of the learner chosen.

c. Update the weights of the training examples  $\omega_{t+1}$  and return to

The boosting framework is a method for combining weak learner  $f_i(x)$  to form a strong classifier F(x).

$$F(x) = sign\left(\sum_{i=1}^{T} \alpha_i f_i(x)\right)$$
(5)



Fig. 1. Relationship of different parameters with MSE.

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	DT	SVM	KNN	GBR	ABR
MSE	2.8873	2.8120	0.3218	0.0891	0.0622
MAE	1.2112	1.1907	0.3779	0.1717	0.1607
$R^2$	0.0478	0.0205	0.6956	0.9219	0.9421

MSE, Mean-squared error of the prediction; MAE, mean absolute error of the prediction;  $R^2$ , squared correlation coefficient of the prediction.

Two typical representatives in boosting framework were the gradient boosting regressor (GBR) [10] and AdaBoost regressor (ABR) [11]. ABR used fractional error points to identify problems and improve the model by adjusting the weight of fractional error points. The samples misclassified by the previous basic classifier could be strengthened, and the weighted samples could be used to train the next basic classifier again. At the same time, a new weak classifier was added in each round until a predetermined small error rate or a predetermined maximum number of iterations was reached. Different from ABR, GBR selected the gradient descent direction during iteration to ensure the best final result. If the loss function declined continuously, it indicated the model was constantly improving, and the best way was to make the loss function decline in its gradient direction.

# 4. Experiments

## 4.1. Data set

The data set in the experiments came from the public data set of Seoul National University Hospital, Seoul, Republic of Korea (https://vitaldb.net/dataset/?query=viewer). The public data set is a comprehensive data set of surgical patients composed of intraoperative bio-signals and clinical information. The bio-signal data included in the public data set were high-quality data such as 500-Hz waveform signals and numeric values at intervals of 1–7 s. The acquisition and release of the data were approved by the Institutional Review Board of Seoul National University Hospital (H-1408-101-605). The study was also registered at clinicaltrials.gov (NCT02914444).

The data were obtained from several patients who underwent routine or emergency noncardiac surgery. The average age of the patients was 47.7 years. The data set included EMG, ETCO2, remifentanil dosage, flow rate, and bispectral index (BIS) indicators for these patients. The sampling interval was 1 s. A total of 11,986 sets of data were available in total. Some data were randomly excluded so that the BIS was as evenly distributed as possible. Then, the data were normalized. The data set underwent fivefold cross-validation to improve the generalization ability of the model. The data were divided into five equal portions, one for each experiment, and the rest for training. The experiment was averaged five times. The data set was grouped into training sets and validation sets [12] following the ratio of 4:1. The training set was used to establish the prediction model of the depth of anesthesia. The validation set was used to validate some of the characteristics of the prediction model.

# 4.2. Experimental design

BIS is an integrated anesthesia depth index based on statistical theory by analyzing some characteristic parameters [13]. In recent years, BIS is widely used in the monitoring of the depth of anesthesia [14].

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Observing markers include BIS, EMG, ETCO2, remifentanil dosage, and flow rate. EMG, ETCO2, remifentanil dosage, and flow rate have a great influence on the depth of anesthesia. BIS directly reflects the depth of anesthesia. The BIS value ranges from 0 to 100. A large BIS value indicates a deep degree of anesthesia, and vice versa. Therefore, the input of the depth of anesthesia prediction model was EMG, ETCO2, remifentanil dosage, and flow rate in this study. The output of the depth of anesthesia prediction model was BIS.

In this study, DT, KNN, SVM, and the proposed method (boosting framework) were used to build prediction models to compare the performances in predicting the depth of anesthesia. GBR and ABR were used for setting up the prediction model of anesthesia. Different parameters in algorithms affect the results to a certain extent, such as the max depth of DT, GBR, and ABR, kernal type of SVM, and neighbors of KNN. Therefore, the effect of the prediction models under different parameters was explored.

# 4.3. Results and discussion

In the experiments, the effect of the algorithms was explored under different parameters. A range of parameters from 1 to 50 was tested such as the max depth value of DT, ABR and GBR, and the number of KNN neighbors. Besides, the effect of SVM under four different kernel types, namely linear, rbf, poly, and sigmoid, was also tested. Figure 1 shows that the fluctuation of MSE value of GBR and ABR was smaller than other methods, indicating that the boosting framework method was less affected by parameters. The MSE value of GBR and ABR eventually leveled off at 0.11 and 0.06. Figure 1 also shows the MSE values of the algorithms under different parameters. The MSE value of GBR and ABR (with the max depth value of 15 and 31) was the best, which was far lower than that of other algorithms. Similarly, the optimal max depth of DT was 48, the optimal number of neighbors of KNN was 5, and the optimal kernel of SVM was "linear." The results of the experiments are recorded in Table 1. The indicators included were MSE, MAE, and  $R^2$ .

Table 1 shows the prediction characteristics of these four prediction methods under optimal parameters. The MSE and MAE values of the boosting framework method were lower than those of the other three methods. The  $R^2$  value of the boosting framework method was higher than the other three methods. These results showed that the boosting framework method had higher prediction ability and lower dispersion degree. Figure 2 shows the comparison of predicted and actual values of the four methods.

Compared with ECG and other clinical indicators commonly used in previous studies [1], we selected four clinical indicators EMG, ETCO2, remifentanil dosage, and flow rate that were more closely associated with BIS to predict the depth of anesthesia. Besides, in previous studies on the depth of anesthesia, several popular machine learning algorithms such as DT, KNN, and SVM, were used to build prediction models [4,6,8]. In this paper, a boosting-based prediction model to predict the depth of anesthesia was built based on four clinical monitoring data. The experimental results show that compared with the DT-, SVM-, and KNN-based models in previous studies, the boosting framework prediction method had lower MSE and MAE values and higher R2 values. That means the boosting framework method has a smaller variation range under different parameters. That means the boosting framework method was not easily affected by parameters.

In summary, the results in the table and figure demonstrated that the boosting framework prediction method had higher prediction accuracy and was not easily affected by parameters. That means the boosting framework method could be used to monitor the depth of anesthesia.



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Fig. 2. Comparison of predicted and actual values of DT, SVM, KNN, and the boosting framework.

## 5. Conclusion

In this study, the boosting framework was used to set up a prediction model to predict the depth of anesthesia based on four clinical monitoring data. The boosting framework method had the following advantages: First, it was more accurate and realized the depth of anesthesia prediction successfully. Second, it was less affected by parameters compared with the DT-, KNN-, and SVM-based models. Third, it had higher prediction accuracy and lower discrete degree compared with those of the DT-, KNN-, and SVM-based models. Therefore, the boosting framework method can be adopted to predict the depth of anesthesia.

## **Conflict of interest**

None to report.

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