Machine Learning model for EEG eye state classification

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Abstract

In this work is presented a Machine Learning (ML) model that, given an EEG signal, can predict eye states (i.e. eyes opened or closed) in a time window from 0.1 to 1 second with an f1 score ranging from 0.9823 to 0.8208, and an avarage f1 score of 0.9055. The few predictions where the metric is less than 0.99, match with the transition from eyes opened to closed or viceversa. The findings in this study will contribute to an ideal solution to the problem of EEG based eye state classification.

1 Introduction

If we consider just epilepsy diagnoses, approximately 30% of the patients EEG is misread and consequently misdiagnosed because of human mistakes [1, 2, 3]. This percentage is certainly higher if not only epilepsy, but every EEG diagnosable diseases are considered. Luckily, AI architectures applied to our daily life are rapidly increasing, in fact EEG eye state classification has been successfully applied in the areas of infant sleep-waking state identification [4], driving drowsiness detection [5], epileptic seizure detection [6], bipolar mood disorder classification (BMD) [7], and ADHD patients (Attention Deficit Hyperactivity Disorder) [7].

The fast and powerful cross-validation methods that we rely on in ML, such as *traintest splits* and *k-fold cross validation*, do not work in the case of time series data. This is because they ignore the temporal components inherent in the problem [8, 9]. The work [10] which this project takes inspiration from, may presents the same methodological flaw in how this time series is evaluated. By not respecting the temporal order of instances when evaluating models, it allows them to use "future" information to make the prediction.

What characterizes this work is the use of *walking forward validation*, which is proven to be the most effective cross validation methodology in time series classifiers [11]. Therefore, I developed a walking forward function which divides the dataset into chronologically ordered parts and, in each run, all data available before the part to predict is used as the training set, while the part to predict is used as test-set. Progressively, in each run, the next part is added and taken into account.

A second function is build (sliding_window) which for the whole dataframe, after selecting a time-step, shapes a training, validation and test-set according to the amount of data in that specific time-step. This function is then integrated into the first one (walking_forward_5) so that the model can understand when the eyes are opened or closed in a selectable time window from 0.1 to 1 second.

2 Method

2.1 Data

All information used in this work comes from one persistent EEG recording¹ collected with a 14 channels headset² and each record was labelled manually. The eye state was recognized by camera during the EEG continuous measurement and added later physically to the record in the wake of examining the feature outlines; "1" means an eye-shut and "0" the eye-open state. The record has a duration of 117 seconds with a sampling rate of 128 Hz which imposes a minimum threshold on the recognition rate of algorithm. Consequently there's a total of 14976 measurements (128*117) and 14 features in the data-set, where each of them is the information acquired by one single sensor. The fifteenth channel represents the output.

¹The database corpus was collected from UCI Machine Learning Repository

²Emotiv EPOC Headset

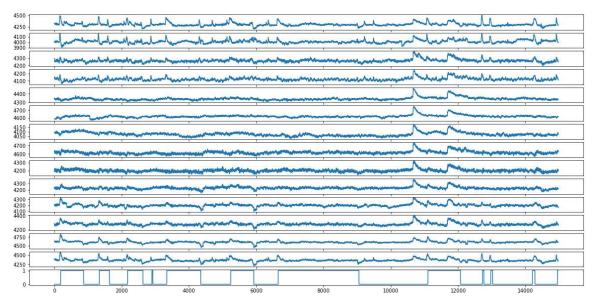


FIGURE 1: Visualization of whole data set

2.2 Model

For this work it was chosen the random forests (RF) algorithm, developed nearly 15 years ago [12], which creates many trees on the subset of the data and combines the output of all the trees reducing overfitting problem in decision trees and the variance, improving therefore the accuracy. It was chosen the RF because it is known indeed, to be one of the best ready-to-go algorithms available for classification, having incredible success across a variety of learning disciplines. For instance, it have fared well in machine learning competitions involving biology such as Ishwaran and colleagues work [13]. Moreover, random forests work poorly any time that one variable is much more important than all the others, which is not the case of this work [14].

2.3 Cross Validation

Temporal components must be considered when analysing time-series, and to do that data can be reshaped so that the value at the previous time-step is used to predict the value at the next time-step, maintaining the chronological order of the data. The sliding_window function shown in Fig.2, do that using as paramaters the dataset, the number of data (array_length) corresponding to the specific time-step (e.g. if 1sec is equal to 128 data, 0.5s is equal to 64, etc.), and the total amount of data in the dataframe. After that, it will model.fit and model.predict a training set and a validation set of length equal to array_length, adding the f1 score to the list f1_tot. This process is looped until it covers the whole dataframe. At the end, the function will return the f1_tot mean.

```
def sliding window(dtset,array length,num dati nel dtframe):
model = RandomForestClassifier(n estimators=45)
array_start = 0
f1_tot = []
for x in range(array length, num dati nel dtframe, array length):
  xtrn_no_output = dtset.drop(['output'],axis=1)
  xtrn = xtrn_no_output[(array_start):(x)]
  ytrn = dtset.iloc[(array_start):(x),-1]
  xval = xtrn no output[(x):((x)+array length)]
  yval = dtset.iloc[(x):((x)+array_length),-1]
  model.fit(xtrn,ytrn)
  prev = model.predict(xval)
  f1_value = f1_score(yval,prev,zero_division=1)
  f1_tot.append(f1_value)
  array_start = x
f1_mean = np.mean(f1_tot)
return f1 mean
```

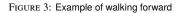
FIGURE 2: sliding_window code

2.3.1 Walking Forward validation

Since non-time-series techniques randomly sample the datasets to create test and training sets, they do not preserve such differences which exist in practice. As said by Falessi et al. "Thus, walk-forward is not only more accurate than 10-fold cross-validation and bootstrap but is also the only one that respects the property of the data, i.e., it preserves the order of data. [...] Therefore, the accuracy measured by non-time-series techniques is poorly realistic of any classifiers' practical adoptions" [11].

During walk-forward, the dataset is divided into parts which are then chronologically ordered and, in each run, all data available before the part to predict is used as the training set, and the part to predict is used as test-set. Afterwards, the model f1_score is computed as the average among runs. For instance, Fig.3 illustrates the walk-forward technique; the parts used for training are in orange, the ones used for testing are in gray, the ones not used are in white. Fig.3 describes a dataset related to a hypothetical project of five parts, and four runs. In the first run, the first part is used as training, and the second as testing, in the second run the first two parts are used as training and the second as testing, and so on. The accuracy is averaged among the four runs.

Run	Part				
	1	2	3	4	5
1					
2					
3					
4					



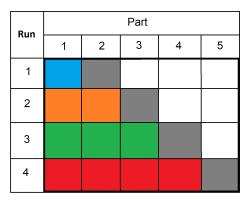


FIGURE 4: Corresponding walking forward

The dataset of this work was also splitted into 5 parts like the example (Fig.4). The walking_forward_5 function includes and loops the sliding_window function four times (Fig.5).

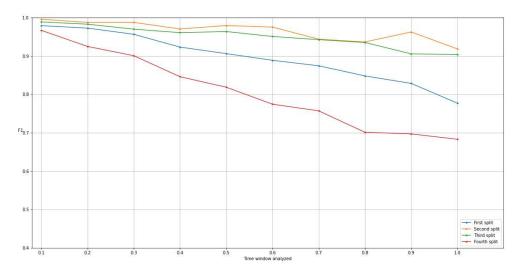
```
def walking_forward_5(dtset,datas_in_onefifth):
   Y_datasVAL = []
   Y_datasTEST = []
   for times in range(1,5,1):
       TrnVal_set = dtset[:(times*datas_in_onefifth)]
       Test_set = dtset[(times*datas_in_onefifth):((times+1)*datas_in_onefifth)]
       for time in range(12,132,12):
           accuracy_per_tenth_s = sliding_window(TrnVal_set,time,(times*datas_in_onefifth))
           Y_datasVAL.append(accuracy_per_tenth_s)
       for time in range(12,132,12):
           accuracy_per_tenth_s = sliding_window(Test_set,time,datas_in_onefifth)
           Y_datasTEST.append(accuracy_per_tenth_s)
   f1_T = np.mean(Y_datasTEST)
   f1_V = np.mean(Y_datasVAL)
   return f1_V, f1_T
```

FIGURE 5: walking_forward_5 code

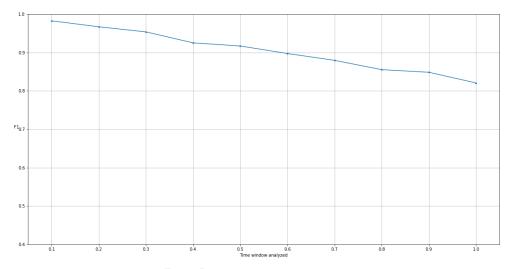
3 Conclusion and Discussion

3.1 Results

The results obtained in each split are showed in Fig.6. The dataset of this work was splitted into 5 parts and a comparison can be made with the w-f example (Fig.4). A second graph where the avarage f1_score for each time-step is avarege is illustrated in Fig.7. At the end, the overall f1_score of the model was measured averaging data from walking_forward_5 function from 0.1s to 1s for all four runs, with a final score of 0.9057.



 $F_{\mbox{\scriptsize IGURE }6:} \mbox{ Walking forward results at different time windows analyzed}$



 $F_{IGURE} \ 7: \ Avarage \ f1_score \ per \ time-steps$

Nevertheless, a further analysis was made to understand where the predictions were inaccurate: Fig.8 showed that these match precisely the moments of transition from eyes opened to eyes closed and viceversa.

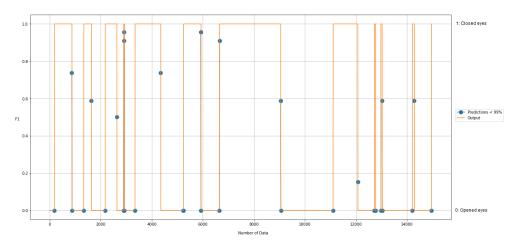


FIGURE 8: Eye-state transition overlapping with f1_score less than 0.99

3.2 Conclusions

The main problem with the present study is that it involves only a single subject which raises the question whether results are generalizable. Another issue concerns the channels quantity on the headset: less sensors would reduce production cost of required EEG devices and also speed up instance-based classification, but would also provide less data.

It is also necessary to consider that the headset used in this work, would not be available in a hospital setting (where is used most of the time) because the minimum amount of channels that must be used are 16 and not 14. In fact, according to LICE³ (Italian League Against Epilepsy) "Sixteen simultaneous recording channels are currently considered the minimum number required to show the areas that produce the most normal and abnormal EEG patterns". Furthermore, in the 10-20 International System, which is used as the standard model for daily EEG, the minimum channels required are even 19 [15].

In the future, a number of different algorithms could be empirically tried in order to be certain that Random Forest is actually the most suitable algorithm for this task.

³"Standard Electroencephalography and Activation tests"

References

- [1] Benbadis, S. R.; Kaplan, P. W.; "The Dangers of Over-Reading an EEG, Journal of Clinical Neurophysiology", July 2019 Volume 36 Issue 4 p 249 doi: 10.1097/WNP.000000000000598
- [2] Benbadis SR. ""Just like EKGs!" Should EEGs undergo a confirmatory interpretation by a clinical neurophysiologist?", Neurology. 2013; 80(1 Suppl 1):S47-S51.
- [3] Selim R Benbadis (2010);"The tragedy of over-read EEGs and wrong diagnoses of epilepsy, Expert Review of Neurotherapeutics", 10:3, 343-346, DOI: 10.1586/ern.09.157
- [4] P. A. Estévez, C. M. Held, C. A. Holzmann et al., "Polysomnographic pattern recognition for automated classification of sleep-waking states in infants", Medical and Biological Engineering and Computing, vol. 40, no. 1, pp. 105–113, 2002.
- [5] M. V. M. Yeo, X. Li, K. Shen, and E. P. V. Wilder-Smith, "Can SVM be used for automatic EEG detection of drowsiness during car driving?", Safety Science, vol. 47, no. 1, pp. 115–124, 2009.
- [6] K. Polat and S. Güneş, "Classification of epileptiform EEG using a hybrid system based on decision tree classifier and fast Fourier transform", Applied Mathematics and Computation, vol. 187, no. 2, pp. 1017–1026, 2007.
- [7] K. Sadatnezhad, R. Boostani, and A. Ghanizadeh, "Classification of BMD and ADHD patients using their EEG signals", Expert Systems with Applications, vol. 38, no. 3, pp. 1956–1963, 2011.
- [8] D. Roberts, V. Bahn, S. Ciuti, M. Boyce, J. Elith, G.G. Arroita, S. Hauenstein, J.J. Lahoz-Monfort, B. Schroder, W. Thuiller, D.I. Warton, B.A. Wintle, F. Hartig "Cross-validation strategies for data with temporal, spatial, hierarchical, or phylogenetic structure"
- [9] "Evaluating time series forecasting models. An empirical study on performance estimation methods" V. Cerqueira, L. Torgo, and I. Mozeti, 2019.
- [10] "A First Step towards Eye State Prediction Using EEG" Rösler et Suendermann, 2013.
- [11] Falessi, D., Huang, J., Narayana, L. et al. "On the need of preserving order of data when validating within-project defect classifiers", Empir Software Eng 25, 4805–4830 (2020). https://doi.org/10.1007/s10664-020-09868-x
- [12] Breiman L."Random forests", Mach Learn. 2001;45:5–32.

- [13] Ishwaran H, Kogalur UB, Lauer MS. "Random survival forests", Ann Appl Stat. 2008;2:841–860.
- [14] Strobl, C., Boulesteix, AL., Zeileis, A. et al. "Bias in random forest variable importance measures: Illustrations, sources and a solution". BMC Bioinformatics 8, 25 (2007). https://doi.org/10.1186/1471-2105-8-25
- [15] "American Electroencephalographic Society Guidelines for Standard Electrode Position Nomenclature", Journal of Clinical Neurophysiology, April 1991