

A survey on common biostatistics tools in neuroscience: Machine learning and Bayesian modeling

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Abstract. Machine learning was characterized by building models and finding correlations between data features, while logistic regression, decision trees, support vector machines (SVM), random forest (RF) and neural networks were recognized as common ML approaches. Bayesian modeling model uncertainty, which can estimate the features from the dataset directly instead of from sampling distribution. Their roles were extremely useful for the detection and progression for diseases in neuroscience. This review summarize different approaches in various diseases, hoping to introduce the potential roles of biostatistics tools in neuroscience.

Keywords: Neuroscience, Biostatistics, Bayesian Modelingy.

1. Introduction

Within the years pasted, the methodologies of biostatistics tools have been used to improve the detection and outcomes of neuroscience. Machine learning (ML) is characterized by using the tools from computer sciences, building models from large datasets, and finding correlations between data features. Based on input-output labels, the most common ML approach, supervised machine learning methods can be used for making decisions or predictions. Classification models are evaluated by F1-score, accuracy, precision, and recall, while regression models are evaluated by mean squared error (MSE) and R-squared. Logistic regression, decision trees, support vector machines (SVM), random forest (RF) and neural networks are the common approaches for the classification models, while linear regression, Lasso regression, ridge regression, decision trees (for regression), and neural networks (for regression) are characterized as useful ways for the regression model.

Bayesian modelling (BM) has been considered as an approach to model uncertainty, which can estimate the features from the dataset directly instead of from sampling distribution. Describing probability distributions, BM treat the features as random variables, which can represent the dependency relationships between variables and handle the uncertainty and randomness between different variables.

The tools of ML and BM have been used in the field of neuroscience, including Alzheimer's disease (AD)(1), Parkinson's disease(PD)(2), cerebrovascular diseases(3), and epilepsy(4). The application of BM can also provide useful approach to the diagnosis and progression of diseases in neuroscience. Thus, the effectiveness of biostatistics tools, ML and BM for instance, to improve the accuracy of detection for diseases in neuroscience will be examined, as well as the optimal choices for clinical decision making and patient outcome.

2. AD

Mild cognitive impairment (MCI) has characterized as the prodromal stage of AD dementia. The deposition of amyloid β ($A\beta$) is regarded as the highly risk of conversion of MCI to AD dementia, however, the deposition of pathological biomarker has mainly based on the positron emission tomography (PET) and cerebrospinal fluid (CSF), which is low availability due to high costs, safety issues, and invasive approaches. Therefore, by integrating and analyzing large amounts of data from different resources, including clinical features, and neuroimaging,

ML can help improve the detection of MCI and the progression of MCI to AD. Recent study found that model(including age, gender, education, diabetes, hypertension, and apolipoprotein E genotype, and neuropsychological test) showed good accuracy(area under operating characteristic curve[AUC]

0.837) in cross validation, and fair accuracy(AUC 0.765) in external validation using RF approach for the detection of amyloid positivity in MCI (5). Logistic regression was useful to predict the existence of pathological biomarkers from clinical risk factors and MRI variables, while the good accuracy was shown from amyloid (AUC 0.79), tau(AUC 0.73) and neurodegeneration(AUC 0.86)(6).

ML also exhibit good accuracy for the prediction of the progression of cognitive impairment. Logistic regression also found that model(including age, instrumental activities of daily living, marital status, and baseline cognitive function) showed fair accuracy (AUC 0.814) to predict cognitive impairment for 3-year by logistic regression(7).Another study also focused on other AD-specific antibody and out-performed accuracy was identified by SVM(AUC 0.94)(8).

Voice signal was also identified as an automated approach for the detection and progression of MCI and AD dementia, while ML was helpful to confirm their potential diagnostic value. The change of prosody was useful for the detection of MCI while principal component analysis was also used to combine the features with the similar character. The discrimination accuracy was 0.90 for training data and 0.65 for validation data measuring by logistic regression(9). The acoustic variables were also able to predict the amyloid status by various ML approaches (AUC 0.79), outperformed the model based on conventional neuropsychological test (AUC 0.66)(10). The joint fusion of acoustic and linguistic parameters were able to achieve AUC with 92.53 for training dataset and 93.89 for test dataset, indicating that speech-processing with able to act as an screening tools for patients with cognitive impairment(11).

BM was also recognized as potential useful approaches for the dection and progression of MCI and AD dementia. Recent study also found that participants with high blood pressure are more likely to exhibit high-risk of pathological biomarkers(12). Another study also found that glial activation of glial activation measuring by baseline PET imaging is a good way to predict the progression of AD(13). The effects of memory and executive performance in amyloid negative participants were also identified by BM(14).

3. PD

ML and BM were also useful for the detection and prediction of PD. Various studies focused on the potential diagnostic value of voice signal using ML approaches. Neural network showed that speech data was helpful to differentiate PD patients from other neurodegenerative diseases with the AUC of 0.93(15). Both ML and deep learning were useful for the dection of PD patients from voice signal(AUC 0.74 and 0.80 respectively)(16). The variables from voice and facial signals showed the good accuracy of the detection for PD using muliple ML approaches, while the RF and logic regression approaches were recognized as the best classifiers(AUC 0.85 and 0.84)(17).

The wearable device was also able to help the dection of frezzing-of-gait from PD patients, while the ML showed a good acurracy measuring by SVM(sensitivity 83%,specificity 80%)(18). Another study also showed that the kinematic data from wearable device was able to detect the subtype from PD with the F1 score of 79.6%(19). RF and decision trees were also helpful from the prediction of the contraction pattern of tremor from PD patients with the AUC of 0.92(20).MLwas trained using accelerometry data to achieve best test performance to distinguish clinically-diagnosed PD, while the results was outperformed compared with genetics, lifestyle and blood biochemistry(21).The fall risk can be predicted by gait parameters deviced from the results in the real-word(21).

BM was also recognized as a useful approach for the ML approaches in the detection and prediction of PD. Degeneration of posterior basal forebrain was associated with cognitive impairment of PD measuring by BM(22). Another study also focused on the roles of locus coeruleus by 7T MRI and the results showed the the vatiability of response inhibition significantly associated with the variability of locus coeruleus integrity measruing by BM(23).BM approaches also used to predict the progression of clinical symptom, while higher systolic blood pressure variability was recognized as a predictor for the progression of cognitive impairment(24). Measuring by BM, another study also found that the change of executive function was correlated with the change of walking performance

and the effect of early motor rehabilitation was also limited by the dysfunction of execution and attention(25).

4. Cerebrovascular disease

Multiple ML approaches were used to evaluate the prediction of the risk of post-stroke depression and anxiety. The results showed that several factors, stroke incidence and history of antidepressants for instance, were able to predict the risk of post-stroke depression and anxiety. Another study also found that clinical risk factor, including diabetes, hypertension, and hyperlipidemia, are able to predict large-artery atherosclerosis with AUC value of 0.93 measuring by logic regression(26). There also exist some studies focusing on the prediction of post-stroke cognitive impairment. Cortical infarcts, atrophy of medial lobe, the severity of initial stroke and the history of stroke were recognized as the predictors for cognitive impairment after stroke with the AUC value of 0.792 by neural network(27). Multiple ML approaches were also helpful to identify the onset time of wake-up stroke from MRI, while the best AUC was 0.895 from SVM(28).

ML was also used to predict the progression of intracerebral hemorrhage after intravenous thrombolysis for stroke. The clinical factors and history of treatment were used as the variables to develop models while the highest AUC was 0.87 for logistic regression(29). The level of consciousness, vital sign, sudden headache, and speech abnormalities were the useful approach to accurately predict the need for surgery with the AUC of 0.802 by extreme Gradient Boosting(30). Combined with clinical and laboratory factors, ten features were able to predict the possibility of death after hemorrhagic transformation with the AUC of 0.85(31). Other study also focused on the potential variables in CT to predict the expansion of perihematoma edema expansion for patients with intracranial hemorrhage with the combined AUC of 0.840(32).

BM was also useful in the prediction and detection of cerebrovascular disease. Glycemic gap, measuring the acute glycemic excursion, was helpful to predict the risk of stroke recurrence by bayesian logistic regression(33). BM was also helpful to estimate the prevalence of other symptom. The prevalence of apraxia of speech was evaluate for chronic aphasia after stroke by BM model(34). For the prediction of the adverse events from drug usage, BM was also recognized as a helpful approach. Several clinical risk factors, sex, anaemia included, were recognized as a model to predict the major bleeding event from the useage of novel oral atrial anticoagulants with atrial fibrillation participants(35).

5. epilepsy

ML approaches were also used for the detection and progression of epilepsy. Deep learning was able to diagnose temporal lobe epilepsy with high accurate based on T2-weighted imaging(T2WI) and fluid-attenuated inversion recovery(FLAIR)(36). Other study also found that centrottemporal spike-waves was able to detect epilepsy with a extremely high level of sensivity(99.8%) and specificity(98.4%) by neural network(37).Other study also focuses on the potential roles of using the electroencephalography to predict impaired consciousness and the best classifier achieved 100% predictive value(38).Neural network was also used to identify focal cortical dysplasias with the sensivity of 85%(39).

6. Conclusion

Recently, various studies focused on the potential roles of ML, BM in particular, for the detection and progression of neurodegenerative diseases. ML was able to identify various model to predict the progression of MCI to AD dementia, and to detect the stage of MCI using other automated device, voice signal for instance. Wearable device and voice signal were also able for the detection of PD and their subtype. For cerebrovascular diseases, ML and BM were helpful to construct model to

predict the risk of progression. ML were also able to develop model to detect special type of epilepsy and predict impaired consciousness.

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