Feature Selection for Brain-Computer Interface Using Nearest Neighbor Information

Yung-Kyun Noh
Department of Computer Science
KAIST
Daejeon, Korea
yungkyun.noh@gmail.com

Byoung-Kyong Min

Department of Brain and Cognitive Engineering
Korea University
Seoul, Korea
min bk@korea.ac.kr

Abstract—We consider the feature selection problem for a braincomputer interface (BCI). A BCI collects data from sensors, and the d ata are discriminated u sing in formation in a dimensional space. We show how re levant f eatures in a high dimensional s pace can b e selected u sing a s imple ne arest neighbor m ethod f or esti mating an i nformation-theoretic measure, J ensen-Shannon d ivergence. Conventional nonparametric estimation using nearest neighbors already works very well for the f eature s election problem and outperforms many ot her methods. In this paper, we show how this nearest neighbor method can be further exploited by properly trimming the non-informative di rection f or a di stance ca lculation, a nd estimate the Jensen-Shannon divergence more a ccurately. Through experiments with synthetic data, we sho who w the proposed meth od outperforms a conventional near est neighbor method as well as other feature selection methods with a large margin.

Keywords-component; featu re selection; J ensen-Shannon divergence; information theory; nearest neighbor

I. Introduction

Communication using a brain-computer int erface (BCI) faces many co mputational issues in e xtracting r elevant information from se nsor s ignals and i n trans forming the information i nto a co de for communication. In gen eral, the entire process d epends on a data-drive n an alysist hrough learning, and many computational problems arise because of the high dimensionality of the data compared with the amount of data and restricted computational resources. Once a proper method is used for feature selection in the first stage, we can take advantages of various perspectives such as the accuracy, speed of communication, learning time, and memory.

In machine I earning c ommunity, many infor mationtheoretic ap proaches ha ve been investigated for feature selection. For di scriminating different si gnals in f eature selection, it is important not to lose i mportant in formation for discrimination, wh ile o ther irr elevant information can be ignored. In addition, sel ecting red undant f eatures is I ess preferred even when the features contain re levant in formation individually. A simple *t*-test and S NR (si gnal-to-noise-ratio) have previously been widely used as statistical criteria to select individual ge nes [1,2,3]. The top features identified by these statistics are then c oncatenated t ogether to form the relevant feature set for analysis. Un fortunately, this approach does not consider any dependency or causality between the identifed features. To obtain the set of features that are collectively the most informative, multivariate methods are necessary. The statistical criterion commonly discussed in this approach is the Jensen-Shannon divergence for two-class classification, which is defined as the mutual information between the data $\mathbf{x} \in \Re^d$ and labels $y \in \{1,2\}$:

$$J_{JS}(X;y) = -\sum_{y=1}^{2} \int p(\mathbf{x},y) \log \frac{p(\mathbf{x})p(y)}{p(\mathbf{x},y)} d\mathbf{x}.$$
 (1)

Here, $p(\mathbf{x}, y)$, $p(\mathbf{x})$, and p(y) are the joint density function, the marginalized density of the data, and the probability of the labels. By selecting d number of sets which maximizes this Jensen-Shannon divergence, we can use a set of the most discriminative features from the perspective of the Shannon information.

In this paper, we explain a nearest neighbor method for estimating t he Jensen-Shannon divergence, and how this method is significantly improved using generative information. By generative information, we mean a method for measuring the distance for the n earest-neighbor's election obtained from the class-conditional densities using g enerative models. F or feature s election, we cho ose a set o f can didate features using forward's election. After we calc ulate t he J ensen-Shannon divergence using the nearest neighbor information along with the generative information, we choose the features one-by-one by recursively selecting additional features that increases the Jensen-Shannon diver gence the most. When comparing the proposed method wit ha conventional method using only nearest nei ghbor i nformation, as well as other w ell-known feature selection methods, the proposed method is shown to be superior in the selection of the r elevant features. In a ddition, once the rele vant features are selected through the proposed selected features yield higher classi fication method, the accuracies.

The rest of this paper is organized as follows. We briefly introduce o ur method for es timating t he J ensen-Shannon divergence in Section II and our feature se lection method in Section III. In Section IV, we present experiments comparing the c lassification acc uracies and the numbers of re levant

features between the proposed and other conventional methods. Finally, we provide some concluding remarks in Section V.

II. JENSEN-SHANNON DIVERGENCE ESTIMATION USING NEAREST NEIGHBOR ESTIMATION

Nearest ne ighbor of one datum is an other datum with the shortest distance between the two. As an example, a method for estimating KL-divergence and Re nyi α -divergence was proposed [6]. Si milarly, we introduce a Jensen-Shannon divergence estimator using the nearest neighbors.

A. Conventional Nearest Neighbor Estimation

A nearest-neighbor est imator f or a J ensen-Shannon estimator can be designed using a Monte-Carlo summation of the nearest neighbor information:

$$\widehat{J}_{JS}(X;y) = \frac{1}{N} \left(\sum_{i; \mathbf{x}_i \in C_1} \log \frac{u_1(\mathbf{x}_i)}{u(\mathbf{x}_i)} + \sum_{i; \mathbf{x}_i \in C_2} \log \frac{u_2(\mathbf{x}_i)}{u(\mathbf{x}_i)} \right)$$
(2)

where $u(\mathbf{x})$, $u_1(\mathbf{x})$, and $u_2(\mathbf{x})$ contain the n earest neighbor information, and each term is defined as $u(\mathbf{x}) = (N-1)l(\mathbf{x})^d$ with $l(\mathbf{x})$, the distance to the n earest neighbor of $\mathbf{x} \in \Re^d$; $u_1(\mathbf{x}) = (N_1 - 1)l_1(\mathbf{x})^d$ with $l_1(\mathbf{x})$, the distance to the nearest neighbor with in class 1; and $u_2(\mathbf{x}) = (N_2 - 1)l_2(\mathbf{x})^d$ with $l_2(\mathbf{x})$, the distance to the nearest neighbor within class 2. Here, N, N_1, N_2 are the number of total data, the number of data belonging to class 1, and the number of data belonging to class 2, respectively, and C_1 and C_2 are the sets of class 1 and class 2.

The asy mptotic con vergence of the is estimator can be proven using the convergence proofs in previous research for a KL-divergence estimator and Renyi α -divergence estimator [4, 6]; however, the proof is not presented in this paper.

B. Metric Learning Using Generative Information

The estimation method described in the previous section is not accurate in a high-dimensional estimation problem because of the i naccuracy of the distance. Inste ad of using the Euclidean distance, metric learning using the Mahalanobis-type distance was proposed in [5]:

$$d(\mathbf{x}_1, \mathbf{x}_2) = \sqrt{(\mathbf{x}_1 - \mathbf{x}_2)^T A(\mathbf{x}_1 - \mathbf{x}_2)},\tag{3}$$

where the positive matrix A is lear ned using a continuous twice-differentiable generative model such as a Gaussian. We obtain the finite sampling bias of Eq. (2) up to the second order, and find the curvature dependency of this bias. We can appropriately control the curvature to minimize the bias. The details of minimizing the bias through metric control can be found in [5].

III. FEATURE SELECTION USING JENSEN-SHANNON DIVERGENCE

The feature's election can be performed by finding a predefined number of features having maximum Jensen-Shannon divergence. However, the combinatorial estimation is often formidable, and a forward selection method is often used. The forward selection in crementally add snew features showing the greatest increase of Jensen-Shannon divergence. For example, the ith feature is selected using the estimated divergence using previously selected i-1 features and new candidate features:

$$\mathbf{x}_i = \arg\max_{\mathbf{x}_k \notin \{\mathbf{x}_1, \dots, \mathbf{x}_{i-1}\}} \widehat{J}_{JS}(\mathbf{x}_1, \dots, \mathbf{x}_{i-1}, \mathbf{x}_k; y)$$

IV. EXPERIMENTS

In this section, we show how the forward selection of the features can be improved using the proposed nearest neighbor Jensen-Shannon estimation with the appropriate metric control. We ge nerated t wo sets of 1000-dimensional Gauss ian data points using the random mean and covariance. The two sets use the same mean and covariance, but for the second set, the mean and covariance of t he first th irty di mensions ar e slightly perturbed. W hen w e c lassify two sets o f data, th e discriminative in formation is only within the first thirty dimensions. In this co nfiguration, conv entional clas sifiers barely succeed in capturing the discriminative information with around 1000 data points. We performed feature selection using our pro posed method with the Jense n-Shannon diver gence estimated usi ng the nearest n eighbor m ethod exploiting the Gaussian g enerative model (JSGNN); Jensen-Shannon divergence us ing t he near est n eighbor method (JSNN); a recently well-known feature selection method, mIMR; and an individual selection of features using the t-score.

In Fig. 1, we used four different configurations of two-class Gaussians by increasing the mean difference between the two sets while using the same covariance configuration. In a high-dimensional space, even when the mean difference is small, there is enough discriminative information that is consistently captured through the proposed feature selection. When the mean difference is small, the conventional mIMR and t-score do not choose informative features (the first 3 0 dimensionalities), while JS GNN and JS NN choose many informative features, and show high classification accuracies. Moreover, JSGNN outperforms JSNN in all cases with a large margin. Herein, we used twenty chosen features.

As sho wn in Fig. 2 , JS GNN can choose in formative features better than o ther methods. The conventional mIMR method bare ly cap tures the relevant features when the mean difference is small, despite the fact that mIMR also considers the Jensen-Shannon divergence. JS GNN consistently chooses around nine of the twenty chosen features. By adopting only generative information, the nearest neighbor estimation method has become significantly useful in the feature selection problem.

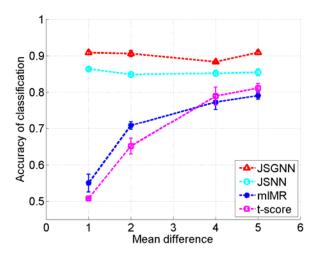


Figure 1. Accuracy of classification with selected features



In th is wo rk, we i ntroduced a novel feature se lection method o btained from the nearest neighbor estimation of an information-theoretic measure, Jensen-Shannon diver gence. We showed that generative information can improve the simple nearest neighbor method significantly using an appropriate metric for a reduction of the estimation bias.

An estimation of Jensen-Shannon divergence is inherently a multivariate approach considering the maximum accumulation of information excl uding redundant i nformation. However, a conventional approximation of J ensen-Shannon divergence such as mIMR does not perform well under all situations owing to an inaccurate estimation. The proposed method can be used in many BCI applications by estimating the correct Jensen-Shannon divergence.

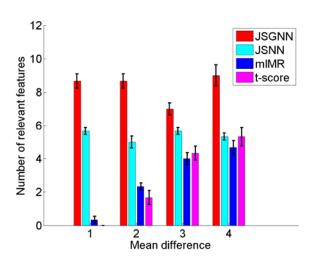


Figure 2. Number of relevant features selected

ACKNOWLEDGMENT

This work was supported by the Basic Science Res earch Program (grant number 2012R1A1A1038358), which is funded by the Ministry of Education, Science, and Technology through the National Research Foundation of Korea.

REFERENCES

- [1] Golub, T. R., Slonim, D. K., Ta mayo, P., Huard, C., Gaasenbeek, M., Mesirov, J. P., Coller, H., Loh, M. L., Downing, J. R., Caligiuri, M. A., Bloomf eld, C. D., & Lander, E. S. (1999). Molecular classification of cancer: class disc overy and class prediction by gene expression monitoring. Science (New York, N.Y.), 286, 531–537.
- [2] Haury, A.-C., Gestraud, P., & Vert, J.-P. (2011). The inf uence of feature selection methods on accuracy, stability and interpretability of molecular signatures. *PLoS ONE* 6(12): e28210.
- [3] Lai, C., Reinders, M., Veer, L. V., & Wessels, L. (2006). A comparison of univariate and multivariate gene selection techniques for classif cation of cancer datasets. *BMC Bioinformatics*, 7.
- [4] Leonenko, N., Pronzato, L., & Sa vani, V. (2008). A class of R'enyi information est imators for multidimensional den sities. *Annals of Statistics*, 36, 2153–2182.
- [5] Noh, Y.-K., Zhang, B.-T., & Lee, D. D. (2010). Generative local metric learning for ne arest neighbor c lassif cation. In Adv ances in Ne ural Information Processing Systems 23, 1822–1830.
- [6] Poczos, B., & S chneider, J. (2011). On the estimation of alphadivergences. Proceedings of the International Conference on Artificial Intelligence and Statistics. 609–617.