Coupled Kernel Ensemble Regression

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ABSTRACT

In this paper, the concept of kernel ensemble regression scheme is enhanced considering the absorption of multiple kernel regressors into a unified ensemble regression framework simultaneously and coupled by minimizing total loss of ensembles in Reproducing kernel Hilbert Space. By this, one kernel regressor with more accurate fitting precession on data can automatically obtain bigger weight, which leads to a better overall ensemble performance. Comparing several single and ensemble regression methods such as Gradient Boosting, Support Vector Regression, Ridge Regression, Tree Regression and Random Forest with our proposed method, the experimental results of the proposed model indicates the highest performances in terms with regression and classification tasks using several UCI dataset.

Keywords

Ensemble regression, Multi-kernel learning, Kernel regression

1. INTRODUCTION

Regression is a technique from fundamental statistic useful for predicting outputs that are continuous. Regression techniques used for predicting data assimilation models have received a lot of active research hot spot in recent times, particularly in real-world applications [6]. Presently, regression is portrayed as one of the most fundamental big data statistical techniques utilized in solving issues of big data [5]. This help in predictions, in which both the sample size and the number of predictors are large for high-dimensional regressions. Additionally, it plays an important role in optimizing operations of complex systems due to its ability to forecast systems behaviors [14]. As a result, regression techniques have been adopted in wide application areas, including but not limited to data mining, computer vision and medical image analysis [18].

Furthermore, a lot of strategies have been adopted in the execution of regression processes with diverse schemes. These schemes are mainly divided into two categories: single regression models and ensemble regression models [11]. The single regression model can also be sub grouped into non-linear and linear methods, whilst linear regression, ridge regression and lasso regression among others are the representative examples of the non-linear method. For example, Santiago et al. [21] demonstrated the effectiveness of multivariate linear regression models towards their application in virtual screening and mechanistic interrogation. Hellton et al. [12] also proposed the use of ridge regression with cross-validation as a plug-in estimate. Mangalathu et al. [19] proposed a methodology to identify the relative impact of input variables and level of treatments needed in the estimation of seismic demand models and fragility curves using lasso regression.

On the other hand, linear methods, such as, kernel ridge regression and support vector regression (SVR) are widely known for their theoretical or experimental results. For example, Li et al. [17] proposed a kernel ridge regression with truncated Gaussian radial basis function kernel (KRR-TRBF) to train classifiers and further authenticates a current user as a legitimate user or an imposter. Cheng et al. [4] developed a full polynomial chaos expansion (PCE) metamodel based on an SVR technique using an orthogonal polynomials kernel function.

Besides, in the second broad category, ensemble regression model combines several decision trees to produce better predictive performance than utilizing a single decision tree. The main principle behind the ensemble model is that a group of weak learners come together to form a strong learner. This has yielded success in many real-world applications, such as decision tree regression, random forest regression and gradient boosting regression. For example, Hariharan et al. [9] explained how a random forest based model was used to estimate model parameters for modeling of a greenfield terrain by non-deposit samples.

The Introduction of the Reproducing Kernel Hilbert Space (RKHS)[22] into the structure of the linear regression methods, significantly contributes to a higher performance result as compared with the non-linear regression methods. Thus the discrete relationship among data samples are characterized better. However, the selection of parameters has great influence on the performance of a single kernel regression method. The selection of a suitable kernel with their parameters is therefore a key problem for kernel regression methods that must be greatly considered.

From the above discussions, we propose a novel coupled kernel ensemble regression(CokER) that takes advantages of both the ensemble method and linear learning method. In the proposed kernel ensemble regression method, base kernel regressors are obtained by varying kernel types and their parameters. Ensemble regressor is therefore obtained by combining base kernel regressors. The coupled kernel ensemble loss is then minimized in multiple Reproducing Kernel Hilbert Spaces (RKHSs). The proposed method optimizes each base kernel regressor in separate RKHSs and then couples them into one regression model in multiple RKHSs. This therefore overcomes the difficulty in the selection of kernel function and parameters which exist in single kernel methods. Similar but different from the previously proposed methods of coupled regression methods in the field of facial recognition [16], artificial neural network [15] and partial least square [8], the proposed method optimizes each base kernel regressor in separate RKHS and then couples them into one regression model in multiple RKHSs whiles the existing coupling methods try to combine multiple RKHSs into one unified space.

The main contributions of this paper are as follows:

- 1. In the proposed method, base kernel regressors are coupled and co-optimized in a coupled ensemble framework by minimizing loss in multiple RKHSs. This is done without multiple RKHSs being combined into one unified space.
- 2. The coupled ensemble idea can find appropriate kernel types and their parameters in a base kernel regressor through a pool of ensemble regression framework, which is different from the state-art-of-work.
- 3. Additional experiments on artificial data sets, UCI regression and classification data sets indicate that compared to other regression methods, for example, random forest and SVR, the proposed method has the advantages of effective performances in keeping lowest regression loss and highest classification accuracy

The rest of the paper is organized as follows: Section 2 introduces some related works with respect to the topic under discussion. Section 3 presents the proposed method. Experimental results are presented in section 4. Finally, section 5 concludes the paper.

2. RELATED WORK

Regression learning has been addressed in a lot of prior studies. In this section, two main categories of regression have been introduced. Thus, single regression model and ensemble regression model. The single regression model is also classified into two sub categories: linear and non-linear methods

2.1 Non-linear methods

Lasso is a non-linear regression method that involves correcting the total size of the regression coefficients. Lasso regression is a regularization technique that's useful for feature selection and to prevent over-fitting training data. It works by penalizing the sum of absolute value (L1 norm) of weights found by the regression. Wang et al. [24] proposed a lasso regression algorithm that employs variable selection for feature variables and further guide the trained predictor towards a generalization solution, thereby improving the accuracy and interpretability of the model. Experimental results showed superiority prediction of fuel consumption compared to existing methods. Zhang et al. [27] recently proposed a locally weighted ridge regression method to overcome the problem of online sensitivity identification using ordinary regression methods that are prone to large errors.

Linear regression which is also an example of the non-linear methods, attempts to model the relationship between two variables by fitting a linear equation to observed data. One variable is considered to be an explanatory variable, and the other is considered to be a dependent variable. Hirukawa et al.[13] demonstrated how the ordinary least square estimator of the linear regression model used matched samples of inconsistent and non-standard convergence rate to deal with earning data of missing observations to be imputed. Experimental results showed that the estimators had an indirect-inference interpretation and attained a parametric convergence rate when the number of matching variables is no greater than four.

2.2 Linear methods

The Introduction of the Reproducing Kernel Hilbert Space (RKHS) into the structure of the linear regression methods, significantly contributes to a higher performance result as compared with the non-linear regression methods. Thus the discrete relationship among data samples are characterized better. The nonlinear problem is transformed into a linear problem through the application of different mathematical kernel functions. Representative kernel methods are kernel ridge regression and SVR methods.

Kernel ridge regression (KRR) is an instance of a natural extension of ridge regression and combines ridge regression with kernel tricks. It thus learns a linear function in the space induced by the respective kernel and the data. For non-linear kernels, this corresponds to a non-linear function in the original space. Chang et al. [2] proposed a kernel ridge regression (DSKRR) method based on divide-and-conquer strategy that provides error analysis for distributed semi-supervised learning. Their results showed that the unlabeled data played an important role in reducing the distributed error and enlarging the number of data subsets in DSKRR.

Support Vector Regression methods are the natural extension of SVM, proposed by Drucker [7]. SVR method is identical to kernel ridge regression base on their model forms. Chen et al. [3] proposed a three-layer weighted fuzzy support vector regression (TL-WFSVR) model for understanding human intention, and it is based on the emotion-identification information in human-robot interaction. Experimental results showed that the proposed TLWFSVR model obtained higher intention understanding accuracy and less computational time than that of the comparative methods.

2.3 Ensemble regression model

Ensemble regression (ER) can combine individual regressors together and keep their performance better as compared to the single regression model. Tree regression method is used to predict the numerical outcomes of the dependent variables. Rathore et al. [20] presented a decision tree regression-based approach for the number of faults prediction in a given software module.

Furthermore, Gradient Boosting Decision Trees (GBDT) is an addictive ensemble regression model in decision trees. Wang et al. [25] proposed a new fusion method based on the LR algorithm and GBDT algorithm for mobile recommendation system. Their method is observed to achieve a good F1 score in a mobile recommendation scenario.

Among ensemble regression methods, random forest (RF) method is a useful machine learning technique which can be applied in both regression and classification problems. Hasan et al. [10] applied random forest for intrusion detection problems. The research indicated that random forest takes less time to train its classifier than SVM and also achieves more accurate results than SVM classifier. Wu et al. [26] used random forest regression approach to analyze the weekly analysis of influenza-like illness rate using one year period of factors. Experimental results showed that regression errors decreased from 5.04% to 4.35% in mean absolute percentage error (MAPE) and 2.85E-04 to 1.97E-04 in mean square error (MSE) for prediction of weekly ILI rate.

3. THE PROPOSED METHOD

This section introduces a new coupled kernel ensemble regression method, which can help to find suitable kernel types and parameters in base kernel regressors. Base kernel regressors are coupled and weighted to form the final ensemble regressor, by minimizing total loss in multiple Reproducing Kernel Hilbert Spaces (RKHSs). In the following subsection introduces the Reproducing Kernel Hilbert Space (RKHS) and then the proposed coupled kernel ensemble regression method is proposed in the next subsection.

3.1 Reproducing Kernel Hilbert Space

Reproducing Kernel Hilbert Space is a special Hilbert space associated with a kernel such that it reproduces (via an inner product) each function in the space. It has a wide range of applications in machine learning, such as SVR and Radial Basis Functions [23].

Given data $\{\{(x_i, y_i)\}_{i=1}^n \in R^p \times R^p\}$, where R^p is a *p*-dimensional real space. A kernel *k* provides a similarity measure between pairs of datapoints

$$k: R^p \times R^p \to R: (x_i, y_i) \leftrightarrow k(x_i, y_i)$$
(1)

The set of these mappings can be extended by including all possible finite combinations, adjoining the limits and constructing an inner product base on the chosen kernel

$$\varphi(x)(x') = \langle k(x, \bullet), k(x', \bullet) \rangle = k(x, x')$$
(2)

where k has a symmetric property that means $k(x_i, x) = k(x, x_i)$. Some suitable functions can be regarded as kernels: -The Polynomial kernel

$$k(x_i, x_j) = (ax_i^T x_j + b)^c$$
(3)

- The RBF kernel(Radial Basis Function)

$$k(x_{i}, x_{j}) = exp(-\frac{\|x_{i} - x_{j}\|}{\mu})$$
(4)

- The Gaussian kernel

$$k(x_i, x_j) = exp(-\frac{\|x_i - x_j\|^2}{2\sigma^2})$$
(5)

where $a, b, c, \mu, \sigma \in R$. Meanwhile, **K** denotes a Gram matrix which is obtained according to samples. It is a symmetric and semipositive definite matrix, which can be shown as follows:

$$\mathbf{K} = \begin{pmatrix} k(x_1, x_1) & k(x_1, x_2) & \cdots & k(x_1, x_N) \\ k(x_2, x_1) & k(x_2, x_2) & \cdots & k(x_2, x_N) \\ \vdots & \vdots & \ddots & \vdots \\ k(x_N, x_1) & k(x_N, x_2) & \cdots & k(x_N, x_N) \end{pmatrix}$$
(6)

The resulting RKHS has then the property that every evaluation operator and norm of any element in H_k is bounded. For a Mercer Kernel $\mathbf{K} : \chi \times \chi \longrightarrow R$, there is an associated RKHS H_k of the function $\chi \longrightarrow R$ with the corresponding norm $\| \|_k$. The standard framework estimates an unknown function by minimizing

$$f^* = argmin \sum_{i=1}^{N} \nu(x_i, y_i, f) + \iota \|f\|_k^2$$
(7)

Where ν is a kind of loss function, such as squared loss $(y_i - f(x_i))^2$ for RLS or hinge loss function max $[0, 1 - y_i f(x_i)]$ for SVM. $\iota ||f||_k^2$ is regarded as a smoothness conditions on possible solutions in the RKHS. In this case, it is proven that the optimal representer of f in Eq.7 can be defined as a finite sum around the observations:

$$f = \sum_{i=1}^{N} \alpha_i k(x_i, x) \tag{8}$$

Therefore, the problem is reduced to optimizing over the finite dimensional space or coefficients α_i , which is the algorithmic basis for SVM and other kernel methods.

3.2 Coupled kernel ensemble regression

The proposed method can combine multiple kernel regressors into a unified ensemble regression framework and the weight of each kernel regressor in this ensemble method is coupled by minimizing total loss of ensembles in Reproducing Kernel Hilbert Spaces. This gives an advantage of one kernel regressor having more accurate fitting precession on data and can, therefore, obtain bigger weight which leads to a better overall ensemble performance

Firstly, different kernels are obtained according to samples. Suppose a regression problem has a training set X with regression result $(X = \{(x_1, y_1), ...,$

 (x_N, y_N)) and a testing set X_t without regression result $(X_t = \{(x_1, ..., x_{N_t})\})$ where $x_n(x_n \in \mathbb{R}^d, n = 1, ..., N)$ expresses a training sample, y_n is the true regression result of x_n , and $x_m(x_m \in \mathbb{R}^d, m = 1, ..., N_t)$ expresses a testing sample. N is the number of training samples and N_t is the number of testing samples. The base kernel regression model is

$$\|K\alpha + b - y\|^2 + \lambda \alpha^T K \alpha \tag{9}$$

where K denotes a kernel matrix which can be obtained according to samples, α is a column vector related to the weight of every sample, b expresses bias term for the specific K.The proposed method aims to obtain the optimal co-regularized weight vector of base regressors. The term $||K\alpha + b - y||^2$ is the square loss for determining the performance of the base kernel regression model.

Unfortunately, since regression performance varies dramatically with the selection of both kernel functions and their parameters, it is also hard to obtain suitable Kernel functions and parameters which are commonly selected manually in practice. To overcome this problem, the proposed method can combine multiple kernel regressors into a unified ensemble regression framework without considering the selection of both kernel functions and their parameters in individual kernel regressors. L different kernels are used in the proposed framework and a new coupled kernel ensemble regression model is proposed:

$$\underset{w,\alpha_{i}}{\operatorname{arg\,min}} \frac{1}{2} \sum_{i=1}^{L} W_{i}(\|K_{i}\alpha_{i} + b_{i} - y\|^{2} + \lambda \alpha_{i}^{T} K_{i}\alpha_{i})$$

$$s.t. \quad 1^{T} W = 1$$
(10)

Where L is the number of kernels. Assuming that the number of training samples is N, and the number of testing samples is N_t and $W = [W_1, ..., W_L]^T$ denotes a weight vector of individual kernel regression model. K_i represents the different kernel matrix. K_i is the i-th base Gram matrix and the dimension of K_i is $N \times N$ for training dataset, $N_t \times N$ for testing dataset. α_i denotes a column vector related to the weight of every sample for each K_i . The dimension of α_i is $N \times 1$ for training dataset, $N_t \times 1$ for testing dataset, b_i is a column vector that has the same dimension as samples, and each value in the vector is equal to a specific K_i . y denotes the true output and its dimension is the same as samples. λ is the constriction parameter that smoothens the model.

We take the derivative of formula 9 with respect to α_i and obtain the following formula.

$$\alpha_i = (K_i + \lambda I)^{-1} y \tag{11}$$

where I is an identity matrix which has the same dimension as training K_i .

According to Formula 11, we can get

$$b_i = \frac{1}{N} \left(\sum_{t=1}^{N} y_t - \sum_{j=1}^{N} K_i(x_j, x_t) \alpha_{i,j} \right)$$
(12)

We considered W_i to be W_i^r (r represents the control parameter for the weights of multiple features) because linear programming attains its optimum solution at the extreme ends, i.e either $W_i = 0$ or $W_i = 1$. That means there will only be one kernel selected contrary to the proposed objective of exploring the rich complementation of multiple kernels. When r = 1, it is only one kernel that will be selected in the optimal result, which is undesirable, but if r > 1the outcome is based on multi-kernel balancing. r is a man-made value to obtain appropriate w. It can further be deduced as:

$$W_{i} = \frac{\left(\frac{1}{\zeta_{i}}\right)^{r-1}}{\sum_{i=1}^{L} \left(\frac{1}{\zeta_{i}}\right)^{r-1}}$$
(13)

Where $\zeta_i = ||K_i \alpha_i + b_i - y||^2 + \lambda \alpha_i^T K_i \alpha_i$ denotes the loss of each kernel. According to Eq, the optimal weight of the ensemble method can be obtained, where r is a parameter to obtain appropriate w. An ensemble regression model is obtain by combining the various base kernel models linearly. The proposed kernel ensemble regressor is built using the following formula 14

$$f(x_t) = \sum_{i=1}^{L} W_i(\sum_{j=1}^{N} K_i(x_j, x_t)\alpha_{i,j} + b_i)$$
(14)

4. EXPERIMENTAL RESULTS

In this section, all the experimental results under different settings are presented. For a fair comparison, each dataset is randomly split into 2/3 (training data) and 1/3 (testing data) and the regularization parameter is obtained by cross-validation method. In our experiments, five comparative methods (Gradient boosting, Tree Regression, Support Vector Regression, Ridge Regression and Random Forest) are selected as base models. Mean Square Error (MSE) and Mean Absolute Error (MAE) are selected as the criteria [1].

$$MAE = \frac{1}{N_t} \sum_{i=1}^{N_t} |f(x_i) - y_i|$$
(15)

$$MSE = \frac{1}{N_t} \sum_{i=1}^{N_t} \left(f(x_i) - y_i \right)^2$$
(16)

In the proposed method, a demonstration of how to combine the base kernel model of the ensemble is shown. A single polynomial kernel model in Eq.3 is applied as the basic model of the ensemble for different datasets.

There are three parameters (a, b and c) in this type of model and the different values of the parameters show different effects on the experimental results. Generally, we set $a \in \{1 * 1e - 6, 1 * 1e - 5, \dots, 1000\}$, $b \in \{1 * 1e - 6, 1 * 1e - 5, \dots, 1000\}$ and $c \in \{1, 2, 3, 4, 5\}$. For each dataset, we select the optimal parameters (a, b and c) and base kernels are obtained by 10-fold cross validation in experiments. The parameter L in Eq.10 denotes the number of base polynomial kernel models. The generalization ability of an ensemble regressor will be good if there are enough base models. However, excessive base models may consist of many worse base models and result in low classification accuracy. Therefore,

Table 1. : Descriptions of UCI dataset

Datasets	Samples	Attributes
Abalone	4177	8
RedWine	1599	11
WhiteWine	1030	8
Mg	1385	6
Space	3107	6

we take $L \in \{10, 20, 50, 100, 150\}$. In our experiments, we select 20 combinations among three parameters (a, b and c).

And the parameter in Eq.10 is the parameter that smoothens the base regressor. The parameter r in Eq.13 is the control parameter for the weights of multiple base models. In our experiments, we select values for λ and r as 0.1 and 2, respectively.

4.1 Dataset description

The selection of nine benchmark publicly available datasets for the evaluation of the performance of the proposed model is made. These datasets are from the UCI database repository, a detailed summary is presented in Table 1.

4.2 Experimental settings

We compared the effectiveness and robustness of the proposed novel Kernel Ensemble Regression with the conventional multikernel features. The performances of some single and ensemble regression preserving methods such as Ridge Regression, Random Forest and Support Vector Regression among others are conducted but with careful tuning of the parameters. All the datasets selected are applied to these methods.

4.3 Performance Evaluations and comparisons

This section discusses the general performance of the proposed coregularized kernel ensemble regression algorithm and all the comparative methods.

Table 2 shows the comparisons of the MSE mean of the CoKer, linear models and ensemble models. As shown in Table 2, the result of CoKER on Abalone dataset is very small as compared to the other comparative methods. This indicates that, CoKER produces a smaller Mean Square Error (MSE) of 3.599 with respect to the abalone datasets. The tree regression model performs poorly with the highest MSE of 4.491. Considering the Red Wine dataset, all the comparative methods and CoKER yielded a positive result with respect to MSE. Nevertheless, CoKER performs best with an MSE of about 0.15% compared to the comparative methods. CoKER leads with a value of 56.0335 which is 3.42% better. It is then followed by ridge regression, while tree regression comes in with the least performance. Tree regression method has the worst performance of 6.4709 for this dataset.CoKER yields better results than the other methods in all the datasets except the Bodyfat dataset. Finally, with a value of 0.022, CoKER leads the others on the Space dataset. From the MSE values presented in table 2, it demonstrates that for MSE values, CoKER proves beyond doubt to be the best method considering the comparative methods

Fig 1 shows another dissimilar view of MSE comparisons among the propose CoKER, single models and ensemble methods on UCI regression datasets. Fig 1(a) demonstrates MSE comparisons on the application of abalone dataset with CoKER having the lowest median of 3.599. The model with the next best median is the random forest. In Fig 1(b), the box plot indicates that for the RedWine

Table 2. : The average of MSE comparison of CoKER, single models and ensemble models

Datasets	CoKER	GB	TR	SVR	RR	RF
Abalone	3.59	3.91	4.49	4.31	4.18	4.00
RedWine	0.41	0.42	0.50	0.61	0.44	0.43
WhiteWine	0.49	0.51	0.53	0.68	0.51	0.49
Mg	0.013	0.016	0.022	0.0175	0.020	0.014
Space	0.022	0.024	0.032	0.039	0.024	0.023

Table 3. : The average of The MAE comparison of CoKER, single model and ensemble models

Datasets	CoKER	GB	TR	SVR	RR	RF
Abalone	0.124	0.134	1.026	1.170	0.174	4.002
RedWine	0.051	0.061	0.397	0.517	0.100	0.070
WhiteWine	0.101	0.104	0.108	0.121	0.152	0.107
Mg	0.0007	0.001	0.011	0.035	0.002	0.001
Space	0.021	0.028	0.038	0.054	0.035	0.046

Table 4. : Descriptions of UCI classification dataset

Dataset	Samples	Attribute
Diabetes	768	8
German	1000	20
LD	345	7
Abalone	4177	8
Dexter	2600	20000

dataset, CoKER again performs better than the other methods with a value of 0.410. Gradient boosting is the second best performer while SVR has the worst lower bound performance. Fig 1(a) and 1(c) shows a lot of flat shaped plot. This implies that, when the regression variance becomes smaller, the more stable the method, and the lower the median, the better the regression result of the method. This because most of the variance is very small, i.e., 10^{-4} and could not be shown in the tables.. Here, CoKER again performed better than the rest of the methods.

From the above discussion, it can conclude that the propose CoKER outperforms the comparative methods.

Table 3 presents the mean MAE comparisons among the propose CoKER, linear models and ensemble models. From the results, it can be seen that, when applied to the Abalone dataset, the CoKER attains the optimal result of 0.1245. Gradient boosting lags slightly behind by 0.01% with a value of 0.134. Random forest yields the worst result with a value of 4.002. When applied to WhiteWine, CoKER performs better than other methods by 1.17%. CoKER provides the best result with a value of 0.134 which is 35% better than the others. CoKER, when applied to Mg turns out to be the best performer with a value of 7.0000e-04 and gradient boosting being the worst performer with a value of 0.001. The Space dataset had the proposed method performing better than the others by 1.3%.

It could be seen from the results in table 3 that the propose CoKER has better MAE values compared with the prior studies on the various datasets. From the experimental results, it could be realized that, the propose CoKER outperforms the prior approaches in all experiments.

Figure 2 also gives a different view of MAE comparisons among the propose CoKER, single models and ensemble methods. From

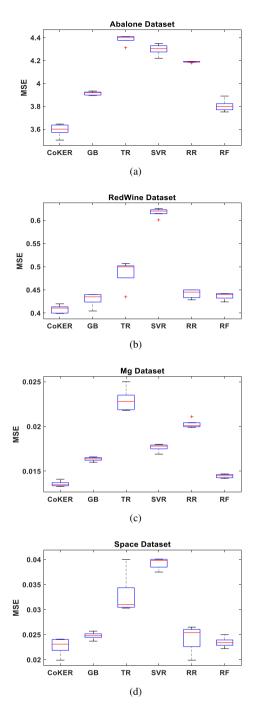


Fig. 1: Box Plot of the respective datasets for MSE: (a) Abalone (b) Redwine (c) Mg (d) space

Fig 2(a) we can see that, has a better MAE value for the Abalone dataset, with the upper bound value of 0.1245. For the WhiteWine dataset, Fig 2(b) shows that CoKER retains the best result with an upper bound value of 0.102. Gradient boosting follows closely behind with a value of 0.1042. A difference of 0.003. From Figure 2,most of the plots have flat shape, which means the smaller the regression variance, the more stable the method. And the lower the

Table 5. : The comparison of classification mean accuracies of CoKer comparative methods, (cls = classification)

Dataset	CoKER	RF	Ridge cls	LibSvm	GB	Tree cls
Diabetes	81.85	80.46	81.64	67.57	78.56	80.34
German	80.18	75.67	79.81	71.17	69.89	73.98
LD	61.73	31.30	61.73	61.73	60.78	61.34
Abalone	80.78	79.94	80.56	79.89	75.86	79.98
Dexter	68.83	67.98	68.56	65.78	63.89	68.67

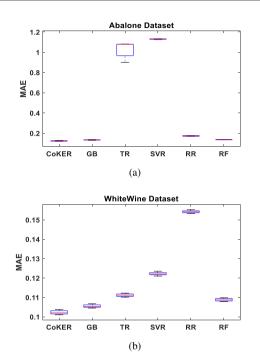


Fig. 2: Box Plot of the respective datasets for MAE: (a) Abalone (b) WhiteWine

median of the method in the figure, the better the regression result of the method. This because most of the variance is very small, i.e., 10^{-4} and did not show them in the tables.

From the above discussion, it can conclude that, the proposed CoKER demonstrates more effectiveness and superiority than the prior studies in regression accuracy.

4.4 Classification

Although all the models discussed in the previous section are intended for regression tasks, classification task is also experimented to further verify the stability of the proposed model.

4.5 Data description

The selection of nine benchmark publicly available datasets for the evaluation of the performance of the proposed model is made, which are Diabetes, German, Liver-disorders (LD), Abalone, and Dexter. A summary is presented in Table 4.

Table 5 did not present the variance of the experiment because they were very small, of about 10^{-4} . which means the smaller the vari-

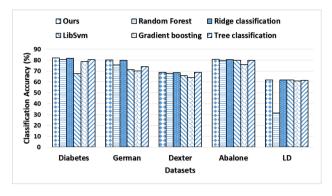


Fig. 3: Comparison of the mean classification accuracies of the various datasets across the five comparative methods

ance, the more stable the method and the lower the median of the method in the table, the better the classification result of the method Fig. 3 presents a comparison of the mean classification accuracies of all the methods across the five datasets. From the figure it can see that, the propose CoKER obtains the highest accuracy of 81.86% on the diabetes dataset, followed closely by Ridge Classification method with an accuracy of 81.66%. Random Forest, Tree Classification and Gradient boosting methods followed suit in that order with the LibSvm method being the worst in classification performance of about 67.58%.

On the German dataset, all the methods show similar performance maintaining their positions as in the Diabetes dataset. Also CoKER and Ridge classification method obtain a slight reduction in classification performance of less than 2%. Whilst Random Forest, Gradient boosting and Tree Classification methods all experience a great reduction of at least 5%.

Dexter dataset got all the methods performing below 70% accuracy, with CoKER leading with an accuracy of 68.83% which is a reduction of about 13% from the diabetes dataset. Random Forest, Ridge Classification and Tree Classification all experience a reduction of close to 13%. Gradient boosting obtains the greatest reduction of about 14% with LibSvm being the least reduced of about 2%.

Finally, on the LD dataset, CoKER obtains the highest accuracy together with two other methods: Ridge Classification and LibSvm. Interestingly, LibSvm which has been the worst performing in all the datasets became one of the best in LD dataset. Also Random Forest which has been performing well in classification in other datasets got the worst classification performance of 31.30%. Tree Classification on the other hand obtains 61.35% of classification accuracy being the second followed by Gradient Boosting with an accuracy of 60.78%.

Generally, CoKER obtains the highest classification accuracy across all the datasets, with the best coming from Diabetes, Abalone and German datasets in that order, followed by Dexter and lastly LD dataset which did not perform so well. It demonstrates a clear distinction between CoKER and the comparative methods on classification datasets as shown in fig.3. Hence, it can concluded that our propose CoKER obtains a better classification performance according to all the experimental results of classification performance.

4.6 Digits Recognition

This section we discusses the classification performance of the propose CoKER in recognition of handwritten digits using MNIST dataset. It contains 10,000 handwritten digit images for classifier

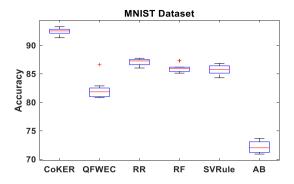


Fig. 4: Box plot of the mean classification accuracies of MNIST dataset across the five comparative methods

testing. Comparing CokER with five different methods namely Weighted Classifier Ensemble method based on Quadratic Forms (QFWEC), Ridge regression (RR), Random Forest (RF), Simple Vote Rule (SVRule) and Adaboost (AB).

Fig. 4 shows the classification accuracy of the propose CoKER and the comparative methods on the MNIST dataset. The proposed method outperforms the rest of the comparative methods. More significantly, AB achieves the lowest classification accuracy performance than the rest of the comparative methods which also perform a poorly as compared to the propose CoKER.

5. CONCLUSION

In this paper, we investigated the problem of how to combine a set of kernel regressors into a unified ensemble regression framework. The framework can simultaneously couple multiple kernel regressors by minimizing total loss of ensembles in Reproducing Kernel Hilbert Space. In this way, one kernel regressor with more accurate fitting precession on data, can obtain bigger weight, which leads to a better overall ensemble performance. Experimental results on several UCI datasets for regression and classification, compared with several single models and ensemble models such as Gradient Boosting (GB), Tree Regression (TR), Support Vector Regression (SVR), Ridge Regression (RR) and Random Forest (RF), illustrate that, the proposed method achieves best performances among the comparative methods.

6. REFERENCES

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