**RESEARCH ARTICLE** 



## Surface water sodium (Na<sup>+</sup>) concentration prediction using hybrid weighted exponential regression model with gradient-based optimization

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## Abstract

Undeniably, there is a link between water resources and people's lives and, consequently, economic development, which makes them vital in health and the environment. Proper water quality forecasting time series has a crucial role in giving on-time warnings for water pollution and supporting the decision-making of water resource management. The principal aim of this study is to develop a novel and cutting-edge ensemble data intelligence model named the weighted exponential regression and hybridized by gradient-based optimization (WER-GBO). Indeed, this is to reach more meticulous sodium (Na<sup>+</sup>) prediction monthly at Maroon River in the southwest of Iran. This developed model has advantages over other previous methodologies thanks to the following merits: (i) it can improve the performance and ability by mixing the outputs of four distinct data intelligence (DI) models, i.e., adaptive neuro-fuzzy inference system (ANFIS), least square support vector regression (LSSVM), Bayesian linear regression (BLR), and response surface regression (RSR); (ii) the proposed model can employ a Cauchy weighted function combined with an exponential-based regression model being optimized by GBO algorithm. To evaluate the performance of these models, diverse statistical indices and graphical assessment including error distributions, box plots, scatter-plots with confidence bounds and Taylor diagrams were conducted. According to obtained statistical metrics and verified validation procedures, the proposed WER-GBO resulted in promising accuracy compared to other models. Furthermore, the outcomes revealed the WER-GBO (R = 0.9712, RMSE = 0.639, and KGE = 0.948) reached more accurate and reliable results than other methods such as the ANFIS, LSSVM, BLR, and RSR for Na prediction in this study. Hence, the WER-GBO model can be considered a constructive technique to forecast the water quality parameters.

Keywords Water quality  $\cdot$  Wavelet transform  $\cdot$  Weighted exponential regression  $\cdot$  Gradient-based optimization  $\cdot$  Bayesian linear regression

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#### Introduction

One of the serious problems attracting much attention recently is the issue of water environment pollution (Uddin et al. 2021). Owing to the importance of management and prediction of water quality (WQ) in a real-life water environment, researchers have developed numerous parameters (physical, biological, and chemical) that serve as the yardstick for the evaluation of water pollution (Vasistha and Ganguly 2020). The role of such parameters is to capture the balance between the rate of atmospheric exchange, oxygen consumption processes (such as nitrification, chemical oxidation, and aerobic respiration), and oxygen production processes (such as photosynthesis). Hence, there is a need to come up with a method for the accurate prediction of water quality (Haghiabi et al. 2018), because an accurate water quality prediction has a profound impact on the water management (Tao et al. 2019). Being that most of the available indicators of water quality normally vary with time, decision-makers can only rely on accurate water quality prediction in making accurate water environment management decisions that will ensure the maintenance of water quality values within an acceptable limit (Tiyasha et al. 2021a, b).

Data intelligence (DI) models have found increasing application in environmental engineering (Chen et al. 2008; Yetilmezsoy et al. 2011; Chakraborty et al. 2019; Hou et al. 2020; Al-Sulttani et al. 2021; Khan et al. 2021; Song et al. 2021). They have specifically been applied in DI-driven techniques in climate, morphology, hydrology, geology, and water chemical-related problems (Maier et al. 2014; Nourani et al. 2014; Yaseen et al. 2019; Li et al. 2021a). DI techniques have been successfully applied in the development of different learning models in the field of environmental monitoring, especially in the monitoring of hydrological parameters (Feng et al. 2021; Tiyasha et al. 2021a, b). In the study by (Fan et al. 2018), authors reported the real-time application of an intelligent water regimen monitoring system that could improve working efficiency. The performance of artificial neural network (ANN) integrated with wavelet transformation (WT) was tested for prediction the electrical conductivity (EC) by (Ravansalar and Rajaee 2015). A hybrid approach that relies on training an ANN model utilizing the multi-objective genetic algorithm (MOGA) was developed by Chatterjee et al. (2017) for improved WQ prediction accuracy (Chatterjee et al. 2017). The suggested model's performance was compared to that of the ANN-GA, the improved ANN model with particle swarm optimization method (ANN-PSO), and the SVM, and the findings revealed that the proposed model outperformed the benchmarking techniques. A hybrid model termed genetic algorithm (GA) and LSSVM (GA-LSSVM) was

developed by Bozorg-haddad et al. (2016) to predict water quality characteristics (Bozorg-Haddad et al. 2017). They conducted a comparison between the suggested model and genetic programming (GP). Na +, K +, Mg +, SO-24 and Cl dissolved solids (TDS) were modeled using a GA-LSSVR algorithm and a GP technique in the Sefidrood River in Iran. The findings show that the GA-LSSVR algorithm outperforms the GP algorithm when it comes to predicting water quality metrics. In another research, EC was predicted using coupled wavelet extreme learning machine (W-ELM) (Barzegar et al. 2018). Abobakr Yahya et al. (2019) revealed that an effective model for the estimation of WQ in the Ungauged River Catchment under Dual Scenarios had been developed (Abobakr Yahya et al. 2019). With modest prediction errors, their model could accurately forecast the WQ. Total dissolved solids were predicted using coupled multigene genetic programming (MGGP) with wavelet data pre-processing (Jamei et al. 2020). For the hybrid RNNs-DS model, Li et al. (2021a, b) merged recurrent neural networks (RNNs) with modified Dempster/Shafer (D-S) evidence theory to produce an RNNs-DS model that was both accurate and efficient (Li et al. 2021b). An array of ensemble machine learning models, comprising quantile regression forest (QRF), random forest (RF), radial support vector machine (SVM), stochastically gradient boosting (SGB), and gradient boosting machines (GBM), were used by Al-Sulttani et al. (2021) (Al-Sulttani et al. 2021) to estimate the amount of biochemical oxygen demand (BOD) in the Euphrates River, Iraq. Two separate feature extraction approaches, genetic algorithm (GA) and principal components analysis (PCA), were used to create integrative models. This investigation shows that the PCA-QRF model identified in research outperformed the solo methods and the GAintegrated models in this study. To estimate the BOD in the Haihe River Basin in China, Song et al. (2021) (Song et al. 2021) developed a novel hybrid model, the improved sparrow search algorithm (ISSA), which combined Cauchy mutation and opposition-based learning (OBL) with the long short-term memory (LSTM). The findings reveal that the addressing model outperforms the competitor models in terms of prediction accuracy.

One of the non-renewable resources affecting human life greatly is surface water owing to its importance to human existence (Asadollah et al. 2020; Tiyasha et al. 2021a, b). However, the rapid increase in economic development has negatively impacted the quality of surface water in some developing countries, and the sustained deterioration in surface water quality increases water stress and water demand, with consequential threats to human health in such regions (Tabari et al. 2011; Uprety et al. 2020; Wu et al. 2020; Dai et al. 2021; van Vliet et al. 2021). Hence, precise and timely monitoring and prediction of water quality are greatly important to avert threats to human life. Although several review articles on the prediction of water quality parameters using DI models have been published (Chen et al. 2020; Rajaee et al. 2020; Giri 2021; Tiyasha et al. 2021a, b), some limitations of the existing models as reported in the literature gives rise to the development of new DI models using different water quality parameters to ensure prediction accuracy improvement. The environmental protection agencies must take certain rapid measures when alerted on impending deterioration in water quality. However, these agencies mostly rely on old models for water quality prediction, and the low precision of these old water quality prediction models may limit their future application; the availability of few training data in the literature may also limit the precision of these learning models even though there are still uncertainties regarding the relationship between the amount of training data and the performance of surface water quality prediction models.

The performance of DI models in water quality prediction is not usually only a function of the applied models and the amount of dataset, but also it is determined by the selected water parameters for the training of the learning models (Tiyasha et al. 2020; Abba et al. 2020). Hence, the most important water parameters for the training of the learning models must be identified and selected without sacrificing the predictive performance of the model; this could significantly improve the prediction efficiency and reduce the overall prediction cost (Yaseen et al. 2018; Tian et al. 2022). Nevertheless, there are still studies on the evaluation of the performance of learning models in surface water quality prediction to identify the most important water parameters from large training datasets.

The majority of DI models have achieved significant results as each model has its advantages and disadvantages. In order to deal with the concerns of each model or the advantages of all models, ensemble models are attaining more attention (Jamei et al. 2021b; Pandey et al. 2021). The main goal of this research is to develop ensemble models using artificial intelligence (AI) (Ahmadianfar et al. 2021, 2022) techniques to forecast sodium (Na) in the Tange-Takab station, in the south of Iran. The proposed model is based on a weighted exponential regression optimized by gradient-based optimization (WER-GBO). To the best of the authors' knowledge, there are no published studies demonstrating the development of the WER-GBO model in the field of modeling, estimating, and predicting. The goal was obtained in three main steps: (1) selection of the best combination of input variables through the best subset analysis, (2) using four DI models (i.e., adaptive neuro-fuzzy

inference system (ANFIS), least square support vector machine (LSSVM), Bayesian linear regression (BLR), and response surface regression (RSR)) to predict the *N*a and finally development of an ensemble model (i.e., WER-GBO) to boost the performance of the single DI models in forecasting the Na.

The current research objectives are (i) to develop a robust and accurate predictive model based on the hybridization weighted exponential regression model using gradient-based optimization for monthly records of sodium (Na) at Maroon River in the southwest of Iran and four DI models (i.e., ANFIS, LSSVM, RSR, and BLR), (ii) to introduce the data preprocessing and best input variables selection, (iii) to compare results by using the standalone DI models, wavelet-based DI models, and the proposed ensemble model, and (iv) finally, to describe the conclusion.

## Methodology

#### Weighted exponential regression (WER)

WER is a weighted regression model based on the exponential function. The proposed WER uses a Cauchy function as a weighting function to increase the accuracy of the regression model's results. In addition, to show the ability of the WER model, predicting the Na is also modeled by a linear model called WLR. The WER and WLR models for the Na prediction are formulated as follows,

$$y_{WER} = w_1 X_1^{\beta_1} + w_2 X_1^{\beta_2} + w_3 X_1^{\beta_3} + w_4 X_1^{\beta_4}$$
(1)

$$y_{WLR} = w_1 X_1 + w_2 X_2 + w_3 X_3 + w_4 X_4$$
(2)

where  $y_{WER}$  and  $y_{WLR}$  are the outputs of WER and WLR models.  $\beta_1$ ,  $\beta_2$ ,  $\beta_3$ , and  $\beta_4$  are the tuning parameters of WER model.  $w_1$ ,  $w_2$ ,  $w_3$ , and  $w_4$  are the weighting functions based on the Cauchy function, which is formulated as,

$$w = \frac{a}{\left(b + c \cdot \left\|\left|x_i - x_j\right|\right\|^2\right)}$$
(3)

The gradient-based optimization algorithm is developed to extract optimal amounts of main unknown parameters of these two models based on measured data to predict the Na.

Accordingly, the following stages are implemented to derive optimal amounts of unknown coefficients of two models:

(1) All datasets consisting of input and output parameters (i.e., lag-time of discharge (Q) and Na) are collected.

(2) The collected datasets from 1980 to 2016 are utilized in the GBO to determine the best values of tuning parameters of two models. The main criterion to choose the best values for parameters of two models is the minimum objective functions, defined as,

Minimize 
$$g(x) = \sum_{i=1}^{N} (Na_{M,i} - Na_{DI,i})^2$$
 (4)

where  $Na_{M,i}$  and  $Na_{DI,i}$  are the measured and DI models, N denotes the number of the dataset.

(3) Optimal amounts of the main parameters of each model are selected based on stage (2).

## **Gradient-based optimization (GBO)**

GBO is a math-based optimization algorithm developed based on Newton's method (Özban 2004) to solve complex optimization problems (Ahmadianfar et al. 2020a). The method consists of two operators: (1) gradient search rule (GSR) and (2) local escaping operator (LEO). The main stages of GBO are explained in the following parts.

### Initialization

GBO is initialized by creating a set of random solutions (population) with the size of N and D dimension, which is expressed as,

$$x_m = LS + rand \times (US - LS) \tag{5}$$

where *LS* and *US* are the lower and upper limits of the problem space, and *rand* has a random value in [0, 1].

#### Gradient search rule (GSR)

GSR is a search mechanism in the GBO to find the promising regions in the solution space. The mechanism is acquired by Newton's method (Özban 2004). The GSR is expressed as,

$$GSR = randn.R_1.\frac{2\Delta x \times x_m}{(dp_m - dq_m + \epsilon)}$$
(6)

in which

$$dp_m = rand.(\frac{\left[u_{m+1} + x_m\right]}{2} + rand.\Delta x) \tag{7}$$

$$dq_m = rand.(\frac{\left[u_{m+1} + x_m\right]}{2} - rand.\Delta x)$$
(8)

$$u_{m+1} = x_m - randn. \frac{2\Delta x. x_m}{(x_w - x_b + \epsilon)} + GD$$
(9)

where *randn* has a random value with a normal distribution.  $\epsilon$  has a small value in [0, 0.1]. $x_w$  and  $x_b$  are the worst and best positions.  $R_1$  denotes a weighting factor. *GM* means the gradient direction to improve the exploitation. The *GD* is expressed as,

$$GD = rand.R_2 \times (x_b - x_m^{it}) \tag{10}$$

In Eq. (6),  $\Delta x$  is a differential vector, which is expressed as,

$$\Delta x = rand.|S| \tag{11}$$

$$S = \frac{(x_b - x_{a1}^{it}) + \mu}{2} \tag{11-1}$$

$$\mu = 2.rand. \left[ \left| \frac{x_{a1}^{it} + x_{a2}^{it} + x_{a3}^{it} + x_{a4}^{it}}{4} - x_m^{it} \right| \right]$$
(11-2)

where a1,a2,a3, and  $a4(a1 \neq a2 \neq a3 \neq a4 \neq m)$  are the four different integers values, randomly chosen from [1, N], and S is the step size.

#### Calculate the new solutions

To generate the new solution  $(Z1_m^{it})$  in the GBO, the GSR and GD are used by the following equations:

$$Z1_m^{it} = x_m^{it} - GSR + GD \tag{12}$$

$$Z1_m^{it} = x_m^{it} - randn.R_1.\frac{2\Delta x.x_m^{it}}{\left(dp_m - dq_m + \epsilon\right)} + GD$$
(13)

in which

$$GD = rand.R_2.(x_b - x_m^{it}) \tag{13-1}$$

where  $R_1$  and  $R_2$  are two adaptive weighting factors. The leading role of these factors is to make a proper equilibrium between the exploration and exploitation, which are formulated as,

$$R_1 = 2.rand.\omega - \omega \tag{14}$$

$$R_2 = 2.rand.\omega - \omega \tag{15}$$

$$\omega = \left| \alpha \times \sin\left(\frac{3\pi}{2} + \sin\left(\alpha \times \frac{3\pi}{2}\right)\right) \right|$$
(15-1)

$$\alpha = \alpha_{min} + \left(\alpha_{max} - \alpha_{min}\right) \cdot \left(1 - \left(\frac{it}{Maxit}\right)^3\right)^2$$
(15-2)

where *it* denotes the number of iteration, *Maxit* is the maximum number of iterations,  $\alpha_{min}$  and  $\alpha_{max}$  are equal to 0.2 and 1.2, respectively.

To increase the exploitation in the GBO, another solution (16) is formulated as follows:

$$Z2_{m}^{it} = x_{b} - randn.R_{1}.\frac{2\Delta x.x_{m}^{it}}{\left(dp_{m}^{it} - dq_{m}^{it} + \epsilon\right)} + rand.R_{2}.(x_{a1}^{it} - x_{a2}^{it})$$
(16)

Finally, based on the solutions  $Z1_m^{it}$  and  $Z2_m^{it}$ , the new solution  $(x_{new})$  is calculated as,

$$x_{new} = r_1 \cdot \left( r_2 \cdot V 1_l^k + (1 - r_2) \cdot V 2_l^k \right) + (1 - r_1) \cdot Z 3_m^{it}$$
(17)

*if rand* < 0.5

in which

$$Z3_m^{it} = x_m^{it} - w_1 \times (Z2_m^{it} - Z1_m^{it})$$
(17-1)

where  $r_1$  and  $r_2$  have two random values in [0, 1].

#### Local escaping operator (LEO)

The GBO uses an effective operator to avoid the local solutions called the LEO. The new solution  $(x_{new})$  is created based on two random solutions  $(x_{a1}^{it}, \text{ and } x_{a2}^{it})$  and the solutions  $Z1_m^{it}$  and  $Z2_m^{it}$ , which is expressed as,

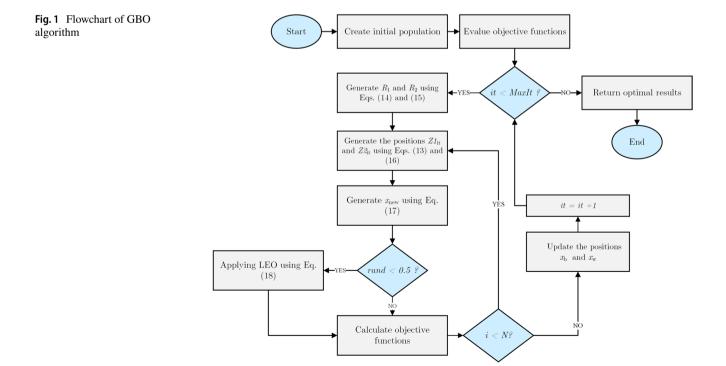
where  $\beta_1$  and  $\beta_2$  have two random values in [-1,1], and  $c_1$ ,  $c_2$ , and  $c_3$  have random values, which are defined as:

$$x_{new} = x_m^{it} + \beta_1 \cdot (c_1 \cdot x_{bs} - c_2 \cdot x_{rand}^{it}) + \beta_2 \cdot R_1 \cdot (c_3 \cdot (Z2_m^{it} - Z1_m^{it})) + \beta_2 \cdot (x_{a1}^{it} - x_{a2}^{it}))/2$$
  
else (18)

$$x_{new} = x_{bs} + \beta_1 \cdot \left(c_1 \cdot x_{bs} - c_2 \cdot x_{rand}^{it}\right) + \beta_2 \cdot R_1 \cdot \left(c_3 \cdot \left(Z2_m^{it} - Z1_m^{it}\right)\right) + \beta_2 \cdot \left(x_{a1}^{it} - x_{a2}^{it}\right)\right)/2$$

End

End



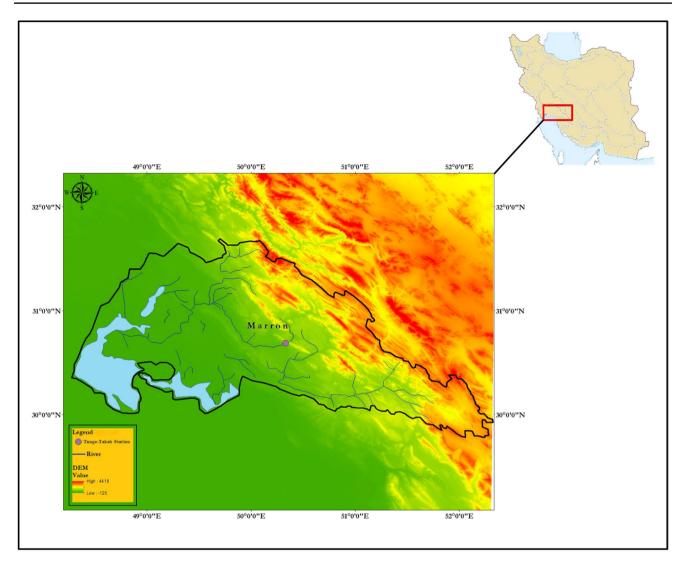


Fig. 2 Location of Tabge-Takab station

$$c_1 = \begin{cases} 2.rand & \text{if } \tau_1 < 0.5\\ 1 & \text{otherwise} \end{cases}$$
(18-1)

$$c_2 = \begin{cases} rand \ if \tau_1 < 0.5 \\ 1 \ otherwise \end{cases}$$
(18-2)

$$c_3 = \begin{cases} rand \ if \tau_1 < 0.5\\ 1 \ otherwise \end{cases}$$
(18-3)

where  $\tau_1$  has a random value in [0, 1].

The solution  $x_{rand}^{it}$  in Eq. (18) is formulated as,

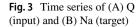
$$x_{rand}^{it} = \begin{cases} x_{rand} & \text{if } \tau_2 < 0.5\\ x_{a5}^{it} & \text{otherwise} \end{cases}$$
(18-4)

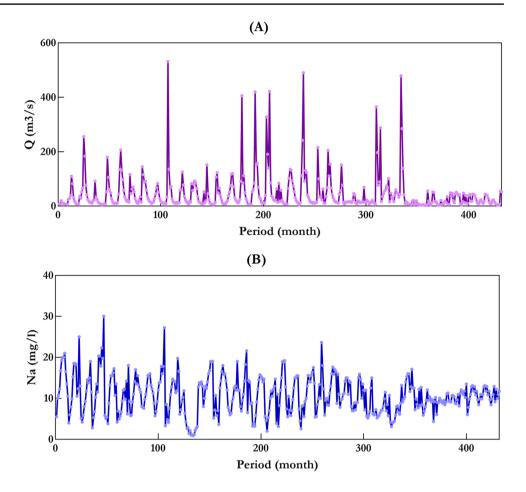
$$x_{rand} = LS + rand.(US - LS) \tag{18-5}$$

where  $x_{rand}$  is a new random solution,  $x_{a5}^{it}$  is a random selected solution ( $a5 \in [1, 2, ..., N]$ ), and  $\tau_2$  has a random value in [0, 1]. The GBO's flowchart is displayed in Fig. 1.

#### **ANFIS model**

There are a large variety of supervised learning algorithms, among which the hybrid learning algorithm is regarded for prediction problems, primarily because the hybrid learning algorithm is widely utilized. ANFIS, a hybrid model, originates from integrating ANN and fuzzy systems, employing ANN learning method to gain fuzzy if-then rules with sufficient membership functions (Jang 1993). It can learn things through vague data and bring along the result. On the other hand, ANFIS uses memory and self-learning neural





network's capabilities efficiently and creates a more constant training process (Huang et al. 2017).

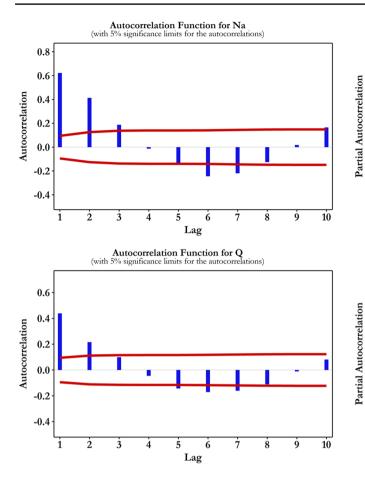
Overall, ANFIS is, in fact, made by five distinct layers (Orouji et al. 2013). To put it simply, the input layer is a previous parameter and the other layers, namely rule-based along with three constant factors and a consequent factor, as well an output layer. Regarding the first layer, it can convert the input into a degree in the range of 0 and 1 or fuzzification, being named premise parameter. Indeed, it can be an activation function along with membership function including trapezoidal or generalized Bell, gauss and triangular (Mohammadi et al. 2016). Furthermore, the second layer can calculate the incoming signals to neurons through the product operator, while the third layer can bring the signals to a stable condition. However, the fourth layer is about fuzzification. The last layer can finally abstract the weighted output proportions.

### Linear Regression (LR)

The linearity can be divided into two descriptions; linear to variable and linear to parameter. That is, modeling is considered a linear model on condition that the model is linear to parameter (Gujarati et al. 2012). Thus, the main criterion for the linear model is linear model to the parameter even though the model is linear to the variable or not to the variable. Regarding multiple linear regression models, their objective is to assess the relationship of two or several independent variables with dependent ones, which is widely used. Making correlation or relationship between dependent and independent variables as well as the relationship between independent ones with each independent linear variable is essential before starting the process of linear regression modeling. Besides, it is important to test the possibility of whether the relationship between the independent and dependent variables could utilize Spearman or Pearson correlation test (Hauke and Kossowski 2011), although there are various testing procedures to examine the linearity test, namely Reset test, White test and Terasvirta test (Teräsvirta et al. 1993). Provided that one dependent variable (y) of m independent variables  $(x_1, x_2, ..., x_m)$  is available, the models of multiple linear regression can be reformulated as:

$$y = a_0 + a_1 x_1 + a_2 x_2 + \dots + a_m x_m + \epsilon$$
(19)

$$y = aX + \epsilon \tag{20}$$



Partial Autocorrelation Function for Na (with 5% significance limits for the partial autocorrelations)

7

6

Lag

0.8

0.6

0.4

0.2

0.0 -0.2

-0.4

0.6

0.4

0.2

0.0

-0.2

-0.4

1

ż

3

4

Fig. 4 ACF and PACF graphs of Q and Na datasets

where y denotes the dependent variable, a is the coefficients of the LR model, and  $\epsilon$  denotes the error vector.

$$Y = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix} X = \begin{bmatrix} 1 & x_{11} & \dots & x_{1m} \\ 1 & x_{21} & \dots & x_{2m} \\ \vdots & \vdots & \vdots & \vdots \\ 1 & x_{n1} & \dots & x_{nm} \end{bmatrix} \boldsymbol{a} = \begin{bmatrix} a_0 \\ a_1 \\ \vdots \\ a_m \end{bmatrix} \boldsymbol{\epsilon} = \begin{bmatrix} \boldsymbol{\epsilon}_1 \\ \boldsymbol{\epsilon}_2 \\ \vdots \\ \boldsymbol{\epsilon}_n \end{bmatrix}$$
(21)

To reach the regression model, the *a* parameter can be calculated by the following formula (Draper and Smith 1998):

$$\widehat{a} = \left(X^T X\right)^{-1} X^T Y \tag{22}$$

#### **Bayesian Linear Regression (BLR)**

Regression modeling with parameter calculation technique is Bayesian linear regression (BLR), wherein the Bayesian method (Box and Tiao 2011) is used. This procedure lies on likelihood, prior and posterior distribution, in which parameters are calculated based on posterior distribution multiplying the prior one with likelihood distribution. Additionally, there is an OLS (i.e., ordinary least square) calculation strategy for linear regression model, with it working by normal distributed error hypothesis, at  $\epsilon \sim N(0, \delta^2)$ . The variables (*YIX*,  $a, \delta^2$ ) are highly likely to be distributed normally since they are similar to the error. Therefore, pdf or probability density function for variables (*YIX*,  $a, \delta^2$ ) ~  $N(Xa, \delta^2)$ can be defined as (Mutiarani et al. 2012):

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$$prob(Y \mid X, a, \delta^{2}) = \frac{1}{\sqrt{2\pi\delta^{2}}} \exp\left[-\frac{1}{2\delta^{2}}(Y - Xa)^{T}(Y - Xa)\right]$$
(23)

According to the pdf, the likelihood of the mentioned variables is formulated in the following:

$$prob(Y \mid X, a, \delta^{2}) = \prod_{j=1}^{m} \frac{1}{\sqrt{2\pi\delta^{2}}} \exp\left[-\frac{1}{2\delta^{2}}(Y - Xa)^{T}(Y - Xa)\right]$$
(24)

$$prob(Y \mid X, a, \delta^2) = (\delta^2)^{-\frac{m}{2}} \exp\left[-\frac{1}{2\delta^2}(Y - Xa)^T(Y - Xa)\right]$$
(25)

Hence, many prior distributions are available to be employed in the Bayesian procedure of linear regression model, among which the distribution of prior conjugate

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Mode	Statistic	Q	Na
Training	No. of observations	302	302
	Minimum	4.900	0.900
	Maximum	532.000	30.000
	Median	23.550	11.450
	Mean	51.575	11.276
	SD	71.707	5.039
	Skewness (S)	3.646	0.262
	Kurtosis (K)	17.070	0.064
Testing	No. of observations	130	130
	Minimum	3.060	3.120
	Maximum	479.000	17.100
	Median	17.900	9.660
	Mean	37.099	9.556
	SD	2.733	64.282
	Skewness (S)	0.042	4.648
	Kurtosis (K)	-0.095	24.643
AC	R <sub>1</sub>	0.439	0.622
	R <sub>2</sub>	0.216	0.413
	R <sub>3</sub>	0.099	0.188
	$R_4$	-0.045	-0.013

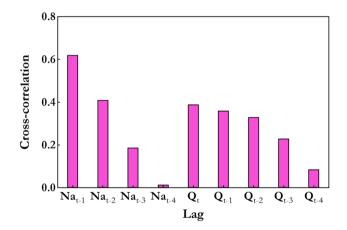


Fig. 5 Cross correlation values between input and target

(Rubio and Genton 2016) can be one of the probable methods. By considering the manner of iteration in marginal posterior, the calculation of regression model's parameters through Bayesian procedure can be done. Also, posterior distribution can be estimated by multiplying the likelihood as well the prior distribution function (Mutiarani et al. 2012).

$$p(a, \sigma^{2} | Y, X) \propto p(Y | X, a, \sigma^{2})p(\sigma^{2})p(a | \sigma^{2}) = (\delta^{2})^{-\frac{1}{2}}$$

$$\exp\left[-\frac{1}{2\delta^{2}}(Y - Xa)^{T}(Y - Xa)\right] \cdot (\delta^{2})^{-\left(\frac{\mu}{2}+1\right)} \cdot (\delta^{2})^{-\left(\frac{\mu}{2}+1\right)} \cdot (26)$$

$$\exp\left(-\frac{us^{2}}{2\delta^{2}}\right) \cdot (\sigma^{2})^{-k/2} \exp\left[-\frac{1}{2a^{2}}(a - \mu)^{T}\Lambda(a - \mu)\right]$$

where k is the number of regression coefficients,  $(\delta^2)^{-(\frac{u}{2}+1)} \exp(-\frac{us^2}{2\delta^2})$  denotes the inverse *Gamma(a,b)*, where a = u/2 and  $b = us^2$ .  $\mu$  is  $[a_0, a_1, \dots, a_m]^T$ .

MCMC or Markov Chain Monte Carlo (Geyer 1992) algorithm is employed to reach the calculation of the regression model's parameters in the Bayesian procedure. Indeed, Gibbs sampling is one of the most well-known approaches in MCMC. In the following, the process of iteration for parameters calculation is done till the burn-in circumstances are completed.

## LSSVM model

Vapnik (1995) presented initially the support vector machine or (SVM) algorithm based on statistical learning theory (Vapnik et al. 1995). This algorithm indeed focuses on formulating the training process in modeling by employing quadratic programming. Likewise, an improved SVM called the least squares SVM (LSSVM) was introduced by (Suykens and Vandewalle 1999) to minimize the computational time of the SVM algorithm process. Take into account a training set of *M* datasets and the input and output data  $x_m$ and *y*, respectively. The LSSVM-based models in feature space for regression matters are described as (Samui and Kothari 2011):

$$y = Q^T \mathcal{O}(x) + c \tag{27}$$

Table 2	BRS to optimally select
the inpu	t combination of Na for
DI mod	els

	Combinations	$\mathbb{R}^2\%$	$R^{2-Ajd}\%$	C <sub>p</sub>	PC
1	Na <sub>t-1</sub>	0.386	0.384	90.192	0.617
2	Q <sub>t</sub> /Na <sub>t-1</sub>	0.486	0.484	7.610	0.519
3	$Q_t / Na_{t-2} / Na_{t-1}$	0.494	0.490	3.053	0.513
4	$Q_{t} / Na_{t-3} / Na_{t-2} / Na_{t-1}$	0.496	0.492	2.991	0.513
5	Q <sub>t-2</sub> /Q <sub>t</sub> /Na <sub>t-3</sub> /Na <sub>t-2</sub> /Na <sub>t-1</sub>	0.497	0.491	4.394	0.515
6	$Q_{t-3} / Q_{t-2} / Q_t / Na_{t-3} / Na_{t-2} / Na_{t-1}$	0.497	0.490	6.165	0.517
7	$Q_{t\text{-}3}  / Q_{t\text{-}2}  / Q_{t\text{-}1} /  Q_t  / \text{Na}_{t\text{-}3}  / \text{Na}_{t\text{-}2}  / \text{Na}_{t\text{-}1}$	0.497	0.489	8.000	0.519

400

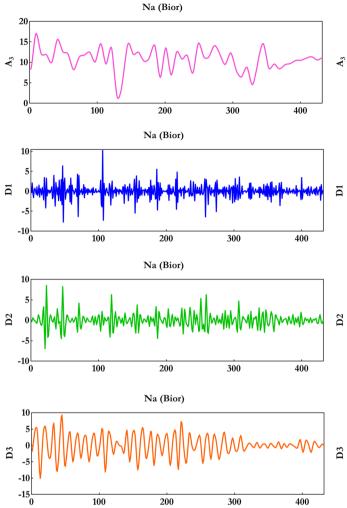


Fig. 6 Decomposition of Q and Na time series for Bior6.8 and Dmey

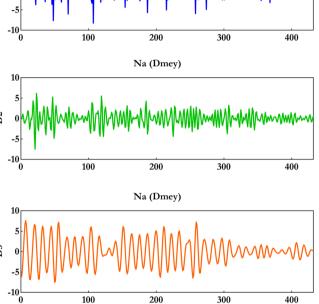
in which  $\emptyset(x)$  is considered an adaptable weight vector, *c* denotes the scalar threshold regarded as bias. *Q* defines the mapping function moving the input data into a larger dimensional feature space. In the following, the given optimization problem is mathematically written for function calculation:

$$\begin{aligned} \text{Minimize} &: \frac{1}{2} \mathcal{Q}^T \mathcal{Q} + \frac{1}{2} \lambda \sum_{t=1}^M v_t^2 \text{ Subjected to } : y \\ &= \mathcal{Q}^T \bigotimes(x) + c + v_t, t = 1, \dots, M \end{aligned}$$
(28)

wherein *v* describes the error variable while  $\lambda$  being the regulative constant. Eventually, the LSSVM can be achieved by making a solution for the given optimization problems:

$$f(x) = \sum_{i=1}^{M} a_i Kr(x, x_i) + c$$
(29)

where  $Kr(x, x_i)$  is considered the kernel function. In this research, radial basis function (RBF) as a kernel function is broadly utilized to support the previous research (Baesens



Na (Dmey)

200

Na (Dmey)

300

20

15

10

5

 $\overset{0}{\overset{}_{0}}{}^{\scriptscriptstyle L}$ 

10

100

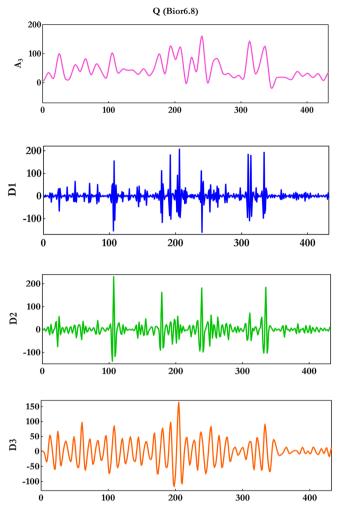
et al. 2000), in which diverse datasets compared the polynomial kernels and RBF. More significantly, the first kernel seems to be more appropriate than the second. RBF can be described by:

$$K(x, x_i) = \exp\left(-\frac{\|x - x_i\|^2}{2\sigma^2}\right)$$
(30)

where  $\sigma$  denotes the tuning parameters of the kernel function.

## **Response surface regression (RSR)**

In order to find a relationship between a number of input variables determined by  $(x_1, x_2, ..., x_m)$  and a response Y, RSR (Gunst 1996; Bezerra et al. 2008) containing a set of approaches is employed. Overall, this relationship is estimated by the polynomial model as:



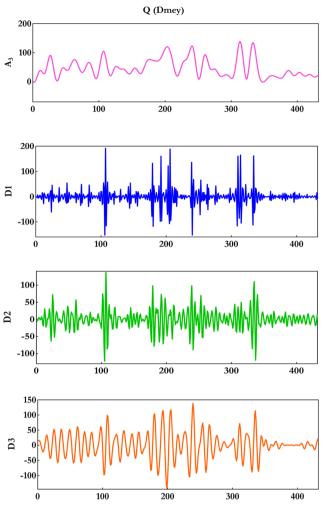


Fig.6 (continued)

$$y = f(X)a + \epsilon \tag{31}$$

in which  $X = (x_1, x_2, ..., x_m)$ , f(X) indicates the vector function of *K* containing the powers and cross-products of  $x_1, x_2, ..., x_m$  to a distinct degree detected by  $d(\ge 1)$ . Besides,  $\epsilon$  defines a random experimental error. *a* describes the vector of *k* stable parameters. In RSM, usually, two major conventional models are utilized, including the first-degree (d = 1)and second-degree model (d = 2). In this study, the seconddegree model RSM is employed, wherein

$$y = a_0 + \sum_{j=1}^{m} a_j x_j + \sum_k \sum_{j>k} a_{ki} x_k x_j + \sum_{j=1}^{m} a_k x_j^2 + \epsilon$$
(32)

## **Evaluation criteria**

In this study, seven well-known statistical metrics are used to assess the efficiency of DI models in forecasting the Na, including the correlation

coefficient (R), mean absolute percentage error (MAPE), root mean square error (RMSE), Willmott's agreement Index ( $I_A$ ), mean absolute error (MAE), relative absolute error (RAE), and Kling–Gupta efficiency (KGE) (Gupta et al.), which are formulated as,

$$RMSE = \left(\frac{1}{N}\sum_{i=1}^{N} \left(Na_{M,i} - Na_{DI,i}\right)^{2}\right)^{0.5}$$
(33)

$$R = \frac{\sum_{i=1}^{N} \left( Na_{DI,i} - \overline{Na_{DI}} \right) \cdot \left( Na_{M,i} - \overline{Na_{M}} \right)}{\sqrt{\sum_{i=1}^{N} \left( Na_{DI,i} - \overline{Na_{DI}} \right)^2 \sum_{k=1}^{N} \left( Na_{M,i} - \overline{Na_{M}} \right)^2}}$$
(34)

$$RAE = \frac{\sum_{i=1}^{N} |Na_{DI,i} - Na_{M,i}|}{\sum_{i=1}^{N} |Na_{DI,i} - \overline{Na_{M}}|}$$
(35)

$$MAE = \left(\frac{1}{N}\right) \sum_{i=1}^{N} \left| Na_{DI,i} - Na_{M,i} \right|$$
(36)

$$MAPE = \left(\frac{100}{N}\right) \sum_{i=1}^{N} \left| \frac{Na_{DI,i} - Na_{M,i}}{Na_{M,i}} \right|$$
(37)

$$IA = 1 - \frac{\sum_{i=1}^{N} (Na_{DI,i} - Na_{M,i})^2}{\sum_{i=1}^{N} \left( \left| (Na_{DI,i} - Na_{DI}) \right| + \left| \left( Na_{M,i} - \overline{Na_M} \right) \right| \right)^2, 0 < IA \le 1$$
(38)

$$KGE = 1 - \sqrt{(R-1)^2 + (SD_{DI}/SD_M - 1)^2 + (\overline{R_{DI}}/\overline{R_M} - 1)^2}$$
(39)

where *N* is the total number of datasets,  $Na_{M,i}$  denotes the measured dataset of *N*a, whilst  $Na_{DI,i}$  expresses the predicted Na by DI models.  $\overline{Na_{DI}}$  and  $\overline{Na_{M}}$  denote the average amounts of the predicted Na and measured dataset correspondingly.  $SD_{DI}$  and  $SD_{M}$  are the standard deviation of the predicted and measured datasets.

In this paper, the performance index (PI), a multi-index criterion, was employed to convert the stated seven metrics into a single metric. This assists the user to facilitate decision making in selecting the best DI model. The proposed PI is formulated as,

$$PI = \frac{1}{7} \left( \frac{R_{min}}{R} + \frac{RMSE}{RMSE_{max}} + \frac{MAE}{MAE_{max}} + \frac{RAE}{RAE_{max}} + \frac{MAEP}{MAEP_{max}} + \frac{KGE_{min}}{KGE} + \frac{IA_{min}}{IA} \right)$$
(40)

where  $MAEP_{max}$ ,  $MAE_{max}$ ,  $RMSE_{max}$  and  $MAE_{max}$  express the maximum amounts of MAEP, MAE, RMSE and MAEcomputed by DI models, whereas  $IA_{min}$ ,  $R_{min}$  and  $KGE_{min}$ indicate the minimum amounts of IA, R and KGE, which are calculated by DI models.

This study uses the Taylor diagram (Taylor 2001), a graphical metric, to clearly show the higher efficiency of the proposed model in comparison with other models. To implement this metric, three statistical metrics are used to plot this graph, such as standard deviation (SD), centered root mean square error (cRMSE), and (R). The performance of DI models is specified by geometrical distance to desired point (target point) in a polar space displayed in the graph (Taylor 2001). The cRMSE is expressed as,

$$cRMSE^2 = SD_M^2 + SD_{DI}^2 - 2.R.SD_M.SD_{DI}$$

$$\tag{41}$$

where  $SD_M$  and  $SD_{DI}$  denote the standard deviation of the measured and predicted amounts by DI models correspondingly.

#### Case study and data preprocessing

## Study area

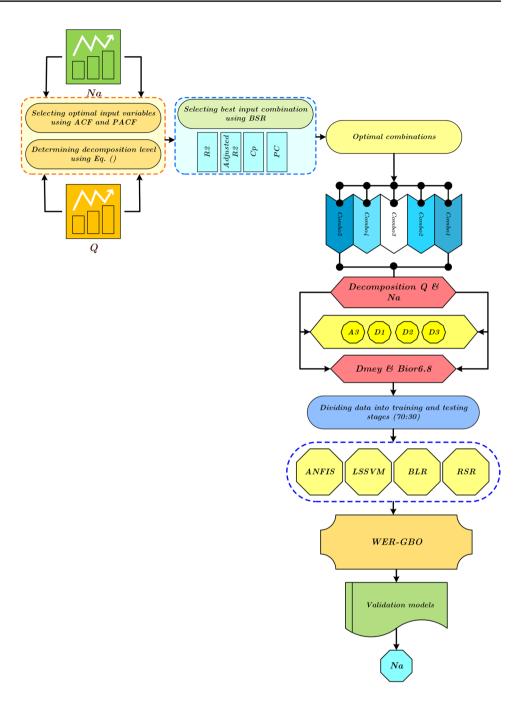
This research lies on water quality prediction based on Na prediction estimated at the Tange-Takab station in Maroun River, Khuzestan province, in Iran with Longitude 50° 20' 02", Latitude 30° 41' 09", and 280 heights from mean sea level. This study is conducted by considering the main water quality parameters including Sodium (Na) and discharge (Q) on a monthly basis. Since the area is comprised of the drainage area of 6,824 km<sup>2</sup> and roughly 310 km long, this river has an integral role in various purposes such as irrigation, drinking water as well recreational activities in Iran, in particular local dwellers in this province. Figure 2 illustrates the locality of this station (Fig. 3).

#### Pre-processing and selecting the best combination

In this part, the principal combinations of the gathered water quality datasets during the 36 years, from 1980 to 2016 on a monthly basis, are addressed in order to forecast the Na through DI models. These data consist of training and testing datasets, being 70 and 30 per cent or 302 and 130 months of the whole dataset correspondingly. The time series of Q as independent variables are displayed in Fig. 4A. To add to it, the Na time series as a target in testing and training intervals are shown in Fig. 4B. Table 1 provides the information on the factor's categorizations, namely kurtosis (K), average (AVG), skewness (S), maximum (MAX), minimum (MIN), autocorrelation coefficients (AC), and standard deviation (SD) for testing, training and other data points. According to Table 1, the amounts of K and S for Na in the training and testing dataset stand at [0.064, 24.643] and [0.262, 4.648] ranges. Nonetheless, K range ([15.4, 48.81]) and S range ([3.469, 6.49]) for the O indicate that the time series' distribution has significant disparity from regular distribution.

One of the most significant steps in the prediction of Na parameter through DI models is to choose the most appropriate input variables' combination. Simply put, the time series' lags have a profound impact on this step (Malik et al. 2019; Jamei et al. 2020). Three statistical methods, namely cross correlation (CC), auto-correlation function (ACF) and partial auto correlation function (PACF), are employed to determine the input variables' input combination (Barzegar et al. 2016).

As illustrated in Fig. 4, the calculation process of efficient input parameters is done by employing PACF and ACF. Apparently, the lagged input variables 1-month and 2-month's AFC has a profound impact on  $Q_t$ , original input datasets. By contrast, the ACF of 1-month, 2-month, and 3-month lagged input variable  $Na_t$  is more efficient compared to the lagged times  $Q_{t-3}, Q_{t-4}, \ldots, ;Na_{t-4}, \ldots$  The 1-month lagged signal can be set for  $Na_t$  and  $Q_t$  with regard to PCFA. **Fig. 7** Flowchart of DI models for predicting Na



As can be observed in Fig. 5, the input signal at the present time,  $Q_t$ ,  $Q_{t-1}$ ,  $Q_{t-2}$ ,  $Q_{t-3}$ , and  $Na_{t-1}$ ,  $Na_{t-2}$ ,  $Na_{t-3}$ , timelagged signals for  $Na_t$  have a large proportion of correlation, bringing along an influential predictive model in comparison with  $Na_{t-4}$  and  $Q_{t-4}$ . The comparison of cross-correlation and target signals ( $Na_t$ ) indicates the  $Na_{t-1}$  and  $Na_{t-2}$  affect significantly  $Na_t$  parameter prediction in target thanks to high correlation coefficients, accounting for by 0.62 and 0.41  $Na_{t-1}$ ,  $Na_{t-2}$  correspondingly. Summing up, the assessment of ACF, CC, and PACF proves that there is a suitable range for lags, as lagged t to t - 4 for  $Na_t$  and t - 1 to t - 3 months in the present month prediction of  $Na_t$ .

The best subset regression, BSR, was addressed in this research to specify the most appropriate input patterns from probable and current patterns. In this regard, a wide range of various factors such as Mallows ( $C_p$ ) (Gilmour 1996), R<sup>2</sup>, Amemiya prediction criterion (PC) (Claeskens and Hjort 2008) and adjusted R are employed to detect optimal input pattern for water quality targets. The PC and  $C_p$  are formulated as (Kobayashi and Sakata 1990):

Table 3Comparison of thestatistic metrics achieved byfour DI models to forecast the	Model	Criteria	Combination	1			
			Combo 1	Combo 2	Combo 3	Combo 4	Combo 5
Na	ANFIS	R	0.610	0.595	0.578	0.564	0.471
		RMSE	3.113	2.990	2.981	2.991	3.272
		MAE	2.378	2.264	2.222	2.198	2.529
		RAE	1.106	1.053	1.033	1.022	1.176
		MAPE	30.459	29.007	29.005	28.141	31.665
		$I_A$	0.716	0.722	0.706	0.708	0.664
		KGE	0.559	0.559	0.542	0.535	0.442
		PI	0.898	0.875	0.879	0.877	1.000
	LSSVM	R	0.567	0.568	0.578	0.567	0.574
		RMSE	2.973	2.926	2.997	2.935	2.897
		MAE	2.240	2.195	2.258	2.189	2.166
		RAE	1.041	1.020	1.050	1.018	1.007
		MAPE	29.236	28.720	29.572	28.468	28.154
		I <sub>A</sub>	0.703	0.705	0.702	0.710	0.716
		KGE	0.531	0.531	0.536	0.535	0.544
		PI	0.995	0.984	0.996	0.980	0.969
	BLR	R	0.540	0.543	0.536	0.535	0.535
		RMSE	2.799	2.781	2.849	2.854	2.851
		MAE	2.139	2.129	2.202	2.206	2.205
		RAE	0.995	0.990	1.024	1.026	1.025
		MAPE	29.183	29.114	30.099	30.154	30.135
		I <sub>A</sub>	0.710	0.711	0.702	0.701	0.702
		KGE	0.526	0.526	0.516	0.515	0.516
		PI	0.978	0.975	0.999	1.000	0.999
	RSR	R	0.572	0.577	0.568	0.559	0.570
		RMSE	2.958	2.704	2.837	3.082	2.889
		MAE	2.290	2.023	2.115	2.336	2.157
		RAE	1.065	0.941	0.983	1.086	1.003
		MAPE	30.034	27.086	28.095	30.645	28.725
		I <sub>A</sub>	0.711	0.733	0.723	0.705	0.719
		KGE	0.542	0.557	0.546	0.525	0.545
		PI	0.977	0.909	0.938	1.000	0.949

Boldface indicates the best prediction results

$$C_p = \frac{RSS_k}{MSE_m} + 2k - N, m > k \tag{42}$$

$$PC = ((n+k)/(n-k))(1 - (R^2))$$
(43)

wherein  $RSS_k$  indicates the squares' residual sum,  $MSE_m$ is considered mean squared error, N denotes the historical number of datasets and k defines the number of predictors. Table 2 depicts the classification of the outcomes of BSR analysis  $Na_t$ , and also the analysis of  $Na_t$  was assessed for five distinct suitable combinations. These combinations are the optimum input data in predictive models according to the most appropriate outcomes, including PC ([0.513, 0.517]),  $\mathbb{R}^2$  ([0.492, 0.497]),  $C_p$  ([2.991, 8.00]). Indeed, this method cannot alone bring along the certainty of the most appropriate input combinations, which in turn a more authentic assessment in the most suitable and possible input combinations is necessary. That is, five input combinations are considered to improve the DI models being specified by grey highlight in Table 2.

## **Results and discussion**

#### Model development

To predict the  $Na_t$  in the wavelet-complementary and standalone frameworks, several DI models, namely LSSVM,

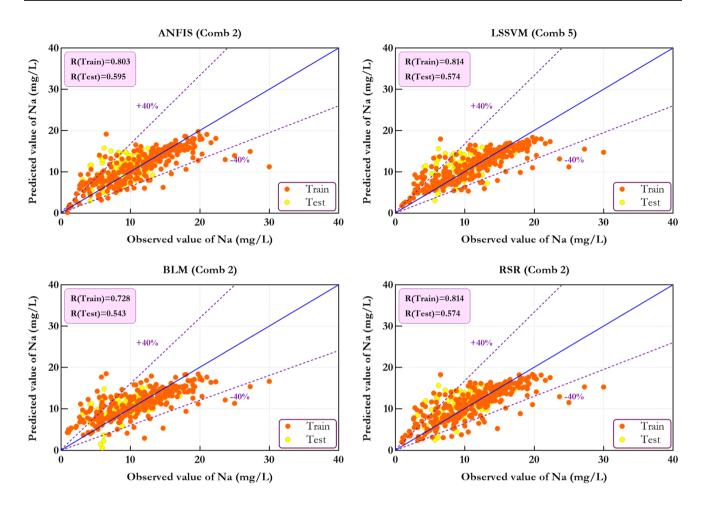


Fig. 8 Comparison of scatter graphs of predicted and measured amounts of Na applying the standalone DI models

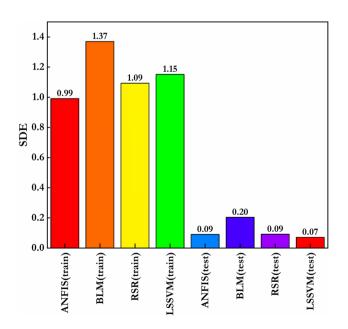


Fig. 9 SDE values of all models for training and testing stages

RSR, BLR, and ANFIS, are utilized in this stage, after which the obtained predicted amounts from these models are considered for inputs in a meticulous model, named weighted exponential regression. This model was in fact integrated with gradient-based optimization (WER-GBO) in order to predict the  $Na_t$ . This means it employs the principal benefit of these four models to reach more exact Na, prediction. More significantly, the essential WER parameters in the proposed model are optimized by the algorithm of GBO. To add to it, using various cutting-edge machine learning methods, namely LSSVM, RSR, ANFIS, and BLR, played an integral role in proving the ability of WER-GBO in forecasting, which eventually brought about striking novelty in this study. The LSSVM acquired their setting parameters during the process of the trial-test procedure, in which the values of parameters are  $\lambda = 8.80$  and  $\gamma = 2.00$ .

## Wavelet-DI models

To boost the certainty and accuracy of DI models, wavelet can be a beneficial tool, mainly because wavelet decomposes

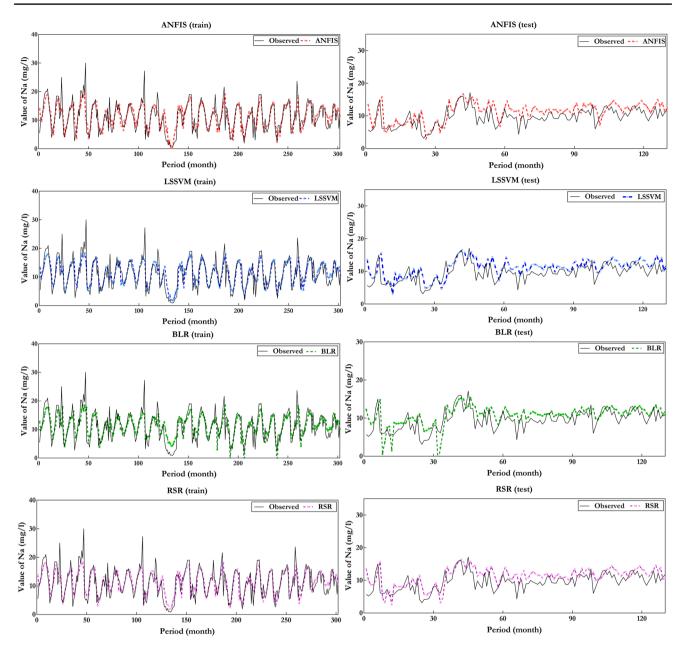


Fig. 10 Variations of the predicted and measured Na for the best performance of DI models

the inputs into two classifications, namely high-frequency (approximate) dataset and low-frequency (details) dataset. One of the most well-known wavelets in hydrological modeling, called discrete wavelet transform (DWT), is employed in this research. Two of most commonly used mother wavelets are discrete Meyer (Damey) and Biorthogonal 6.8 (Bior6.8) (Ahmadianfar et al. 2020b; Jamei et al. 2021a). In fact, these wavelets have been constructive ones in water quality predictive models. In other words, these mentioned wavelets support compressed form while being advantageous in creating time localization (Nourani et al. 2014; Freire et al. 2019; Ahmadianfar et al. 2020b; Jamei et al. 2021a). The mother wavelet (i.e., Bior6.8 and Dmey) was utilized in this study to take apart in the time series. The optimal decomposition grade (1) of wavelet transform for the water quality time series was defined as below (Barzegar et al. 2018):

$$nW = int[\log(M)] \tag{44}$$

wherein M is considered the number of datasets, standing at 432. The proportion of disintegration grade will account for 3.

Table 4Comparison of thestatistic metrics achievedby ANFIS models for allcombinations

Table 5 Comparison of the

statistic metrics achieved by BLR models for all

combinations

Model	Criteria	riteria Combination						
		Combo 1	Combo 2	Combo 3	Combo 4	Combo 5		
Wavelet-Dmey	R	0.931	0.937	0.964	0.966	0.967		
	RMSE	1.065	1.016	0.729	0.699	0.699		
	MAE	0.833	0.745	0.561	0.530	0.526		
	RAE	0.387	0.346	0.261	0.246	0.245		
	MAPE	10.143	9.550	7.050	6.344	6.230		
	$I_A$	0.963	0.966	0.981	0.983	0.982		
	KGE	0.912	0.916	0.952	0.951	0.929		
	PI	1.000	0.953	0.805	0.780	0.781		
Wavelet-Bior	R	0.896	0.902	0.926	0.930	0.905		
	RMSE	1.242	1.200	1.028	1.002	1.168		
	MAE	0.904	0.816	0.767	0.770	0.867		
	RAE	0.420	0.380	0.357	0.358	0.403		
	MAPE	11.537	10.208	9.346	9.233	10.341		
	$I_A$	0.938	0.947	0.960	0.962	0.946		
	KGE	0.801	0.894	0.892	0.885	0.844		
	PI	1.000	0.934	0.882	0.879	0.955		

Boldface indicates the best prediction results

Model Criteria Combination Combo 1 Combo 2 Combo 3 Combo 4 Combo 5 Wavelet-Demy R 0.945 0.966 0.966 0.966 0.968 0.899 RMSE 0.705 0.703 0.704 0.682 MAE 0.714 0.560 0.558 0.544 0.512 RAE 0.253 0.238 0.332 0.260 0.259 MAPE 8.712 7.153 7.058 6.597 6.015 0.971 0.982 0.983  $I_A$ 0.982 0.983 KGE 0.924 0.955 0.955 0.947 0.931 ΡI 1.000 0.872 0.858 0.834 0.870 0.913 0.931 0.925 Wavelet-Bior R 0.929 0.932 0.996 RMSE 1.117 1.008 1.035 0.990 MAE 0.826 0.740 0.751 0.796 0.767 RAE 0.384 0.344 0.349 0.370 0.356 MAPE 9.890 9.024 9.478 8.999 9.010 0.962  $I_A$ 0.953 0.963 0.960 0.963 0.904 0.898 0.890 0.879 KGE 0.885 0.934 ΡI 0.999 0.940 0.968 0.946

Boldface indicates the best prediction results

In the following stage, disintegrated time series being constructive in this process were gathered (e.g.,  $Q = A_3 + \sum_{k=1}^{3} D_k$ ) and set as inputs in order for complementary DI models with regard to the input combinations of *N*a. Besides, the As, approximations, and Ds, details of signals of *N*a and Q simulation, are illustrated in Fig. 6. In Fig. 7, DI models' flowchart for *N*a prediction parameters is drawn.

## Investigation of the standalone DI model abilities

In this step, four DI models, ANFIS, LSSVM, BLR, and RSR, were assessed in terms of performance and ability on the five diverse input parameters' combinations. The outcomes are given in Table. 3, in which the fundamental variety of DI models' forecasting ability in test data for combinations in  $Na_t$  prediction is provided. According to

by LSSVM models for all combinations

Model	Criteria	Combinatio	n				
		Combo 1	Combo 2	Combo 3	Combo 4	Combo 5	
Wavelet-Demy	R	0.936	0.962	0.958	0.954	0.952	
	RMSE	1.008	0.783	0.808	0.846	0.874	
	MAE	0.781	0.604	0.620	0.646	0.660	
	RAE	0.363	0.281	0.288	0.300	0.307	
	MAPE	9.926	7.682	7.790	7.945	8.118	
	I <sub>A</sub>	0.960	0.977	0.976	0.973	0.971	
	KGE	0.850	0.902	0.915	0.901	0.891	
	PI	1.000	0.856	0.866	0.886	0.900	
Wavelet-Bior	R	0.898	0.929	0.922	0.924	0.906	
	RMSE	1.259	1.053	1.098	1.089	1.221	
	MAE	0.960	0.799	0.823	0.827	0.931	
	RAE	0.446	0.371	0.382	0.384	0.433	
	MAPE	12.387	10.069	10.235	10.149	11.746	
	I <sub>A</sub>	0.937	0.957	0.953	0.954	0.940	
	KGE	0.814	0.858	0.851	0.857	0.808	
	PI	0.999	0.886	0.903	0.901	0.978	

Boldface indicates the best prediction results

Table 7Comparison of thestatistic metrics achievedby RSR models for allcombinations

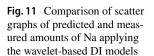
Model	Criteria	Combination				
		Combo 1	Combo 2	Combo 3	Combo 4	Combo 5
Wavelet-Demy	R	0.940	0.961	0.952	0.948	0.932
	RMSE	0.970	0.749	0.847	0.866	1.023
	MAE	0.763	0.590	0.648	0.627	0.753
	RAE	0.355	0.274	0.301	0.292	0.350
	MAPE	9.251	7.595	8.064	7.595	9.210
	IA	0.965	0.980	0.975	0.973	0.964
	KGE	0.901	0.952	0.946	0.941	0.928
	PI	0.991	0.857	0.903	0.892	0.991
Wavelet-Bior	R	0.902	0.921	0.927	0.814	0.878
	RMSE	1.177	1.069	1.027	1.705	1.331
	MAE	0.861	0.808	0.789	1.019	0.943
	RAE	0.400	0.376	0.367	0.474	0.438
	MAPE	10.208	10.173	9.669	12.778	11.333
	I <sub>A</sub>	0.947	0.958	0.961	0.899	0.935
	KGE	0.863	0.904	0.902	0.808	0.870
	PI	0.852	0.818	0.802	1.000	0.905

Boldface indicates the best prediction results

Table 3, the performance and ability of the ANFIS model in forecasting the  $Na_t$  in the testing process proves that Combo 2 (R=0.595, RMSE=2.990, MAPE=29.007, KGE=0.559, and PI=0.875) stands out compared to the other combinations. By considering five combinations, the best combination of inputs in BLR and RSR was detected. The most appropriate combinations for both are Combo 2 (RSR: (R=0.577, RMSE=2.704, MAPE=27.086, KGE=0.557,

and PI = 0.909), and BLR: (R = 0.543, RMSE = 2.781, MAPE = 29.114, KGE = 0.526, and PI = 0.975)). Turning to LSSVM model, the best combination is considered Combo 4 (R = 0.574, RMSE = 2.897, MAPE = 28.154, KGE = 0.544, and PI = 0.969)).

Different models, including ANFIS, LSSVM, BLR, and RSR, are utilized to determine the prediction outcomes of 6 water quality indicators demonstrated in



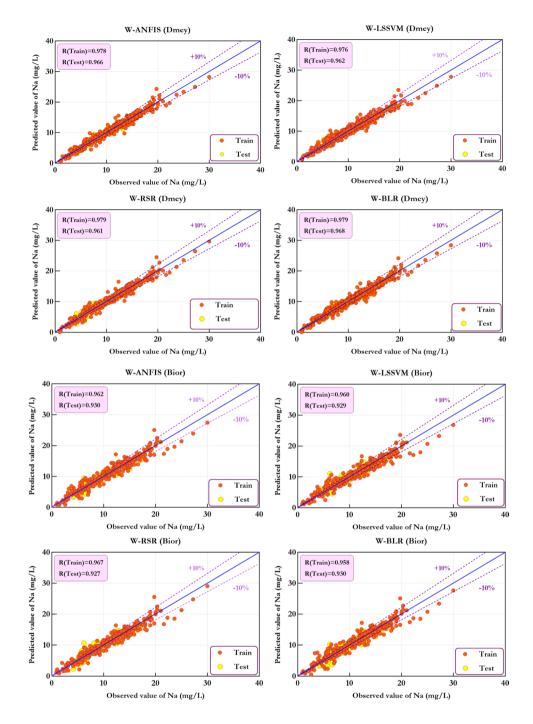


Fig. 8. The X-axis defines the measured value, and the Y-axis depicts the predicted value. More specifically, the ideal forecasting outcomes will be distributed on both sides of the line or Y = X. In turn, it is apparent that the error is conforming to the Gaussian distribution rule. In other words, the location of points based on the line has a direct relation with the amount of error. Thus, providing that the points are adjacent to the Y = X line, the error is minor. On the other side, by considering the

forecasting of 4 indicators, the spots of DI models are out of the Y = X line. As can be observed in Fig. 8, the distribution of prediction outcomes for many models is on the equal side of Y = X, which results in significant deviation and some under-fitting or over-fitting problems. Additionally, the error value stands at  $\pm 40\%$  based on predicted proportions obtained by two distinct DI models. As a consequence, five standalone DI models are not able to predict the Na accurately.

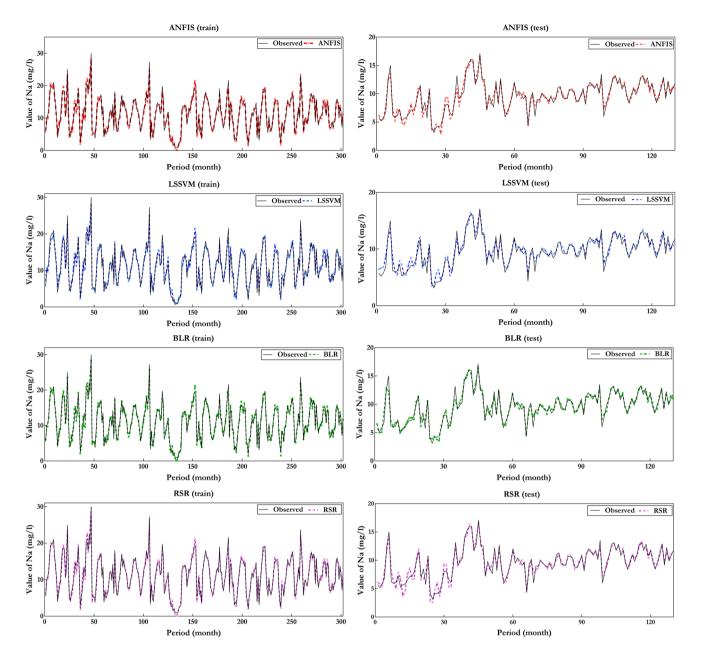


Fig. 12 Variations of the predicted and measured Na for the best performance of W-DI models

Since stability is a vital matter in prediction, the standard deviation error (SDE) can be an influential factor to investigate the models' prediction stability. In Fig. 9, the SDEs of forecasting the different water quality indicators are drawn. More importantly, the SDEs of ANFIS reached the first and second-best place in training and testing stages, accounting for 0.99 and 0.09 correspondingly. Moreover, results reveal the SDEs of LSSVM, standing at 0.07, outperform in the testing step. The efficiency of ANFIS, LSSVM, BLR, and RSR models are addressed using time series diagrams based on the measured and calculated proportion of *N*a within training and testing stages, which is provided in Fig. 10. As a result, all DI models cannot forecast the Na correctly.

# Evaluate the performance of wavelet-based DI models

The W-ANFIS, W-LSSVM, W-BLR, and W-RSR models were improved with the purpose of enhancing the accuracy and certainty of standalone DI models (i.e., ANFIS, LSSVM,

Mode	Criteria	WER-GBO	WLR-GBO
Train	R	0.9813	0.9811
	RMSE	0.9671	0.9741
	MAE	0.7028	0.7125
	RAE	0.1711	0.1735
	MAPE	7.7907	8.2642
	I <sub>A</sub>	0.9905	0.9904
	KGE	0.9731	0.9730
	PI	0.987	1.000
Test	R	0.9721	0.9712
	RMSE	0.6399	0.6525
	MAE	0.4976	0.5030
	RAE	0.2314	0.2338
	MAPE	5.8858	6.0126
	I <sub>A</sub>	0.9854	0.9847
	KGE	0.9486	0.9398
	PI	0.990	1.000

 Table 8 Comparison of statistical metrics of WER-GBO and WLR-GBO models

Boldface indicates the best prediction results

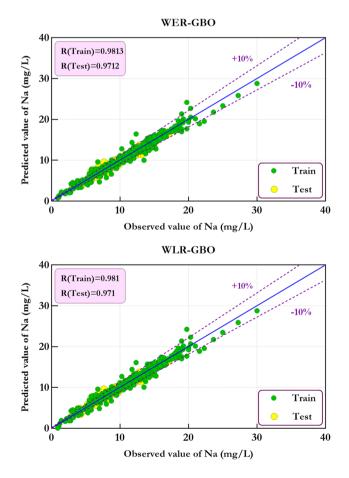


Fig. 13 Comparison of scatter graphs of predicted and measured amounts of Na applying the WER-GBO and WLR-GBO models

BLR, and RSR. Two mother wavelets (i.e., Bior6.8 and Dmey) are used in the decomposition process of time series for *N*a. The assessment of W-DI models' ability with various mother wavelets is implemented by considering five distinct combinations of input variables. The parameter settings of LSSVM-Bior6.8 ( $\lambda = 271.73$  and  $\gamma = 34.24$ ) and LSSVM-Dmey ( $\lambda = 111$  and  $\gamma = 19.4$ ) are calculated based on trial-and-error.

Table 4 provides the data on the forecasting accuracy and certainty of W-ANFIS models considering all combinations and two mother wavelets. As shown, the most suitable combination for the W-ANFIS model with mother wavelets Dmey and Bior6.8 can be Comb 4 (R=0.966, RMSE=0.699, MAPE=6.344, KGE=0.951, and PI=0.780) and also for  $Na_t$  prediction in testing step can be Comb 4 (R=0.930, RMSE=1.002, MAPE=9.233, KGE=0.885, and PI=0.879). The ultimate results prove Dmey outperforms the Bior6.8 for W-ANFIS, which stems from its more accurate and exact performance than Bior6.8 mother wavelet (Table 5).

In the W-BLR model, Comb 4 is considered the best combination being equal to all mother wavelets, and the related data are provided in Table. Moreover, the final results of different mother wavelets for the most appropriate combination stand out at W-BRL-Dmey (Combo5: R=0.968, RMSE=0.682, MAPE MAPE=6.015, KGE=0.931, and PI=0.834), and W- BLR -Bior6.8 (Combo2: R=0.931, RMSE=0.996, MAPE=9.024, KGE=0.904, and PI=0.934). Accordingly, the outcomes confirm that the most suitable mother wavelet is admittedly Dmey in W-BLR, mainly because it reached higher certainty and accuracy than other models.

Regarding W-LSSVM, as can be seen in Table 6, Combo 2 is considered the winner for two mother wavelets. Besides, the estimated outcomes for Bior6.8 and Dmey mother wavelets for the most appropriate combination stand at: W-LSSVM -Dmey (R=0.962, RMSE=783, MAPE=7.682, KGE=0.902, and PI=0.856), and W-LSSVM-Bior6.8 (R=0.929, RMSE=1.053, MAPE=10.069, KGE=0.858, and PI=0.886). Eventually, the best mother wavelet for the W- LSSVM is considered Bior6.8, which results in the highest certainty in comparison with Dmey.

Table 7 reports the W-RSR model's results with Bior6.8 and mother wavelets Dmey, with it indicating Comb 2 (R=0.961, RMSE=0.749, MAPE=7.595, KGE=0.952, and PI=0.857) and Comb 3 (R=0.927, RMSE=1.027, MAPE=9.669, KGE=0.902, and PI=0.802) as the best combinations for Dmey and Bior6.8, respectively. Therefore, Dmey considered the best mother wavelet that makes powerful performance and more accuracy than Bior6.8.

In Fig. 11, the scatter plot of forecasted values versus experimental ones is demonstrated. The main criterion to confirm the accuracy of the proposed ANFIS model in prediction ability can be the high density of points in the vicinity of the Y=Xline. According to this figure, there is a significant coincidence between experimental values and predicted ones. This means the proposed ANFIS model creates acceptable and reliable

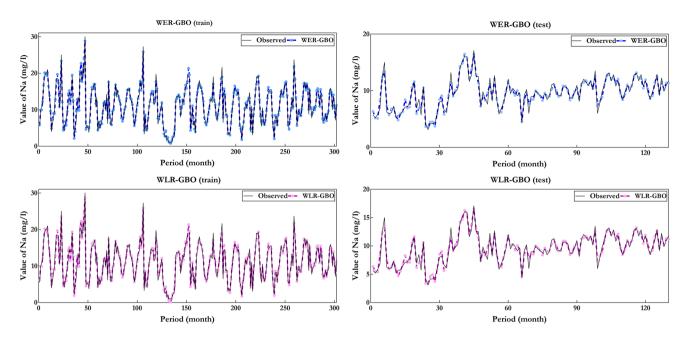


Fig. 14 Variations of the predicted and measured Na for the best performance of W-DI models

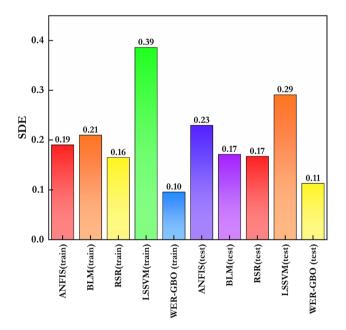


Fig. 15 SDE values of all models for training and testing stages

predictions. The time series plots, related to predicted and experienced *N*a in the testing and training phases, are depicted in Fig. 12. According to this figure, all predictive models reach considerable performance, and the predictive values are equivalent to the estimated Na.

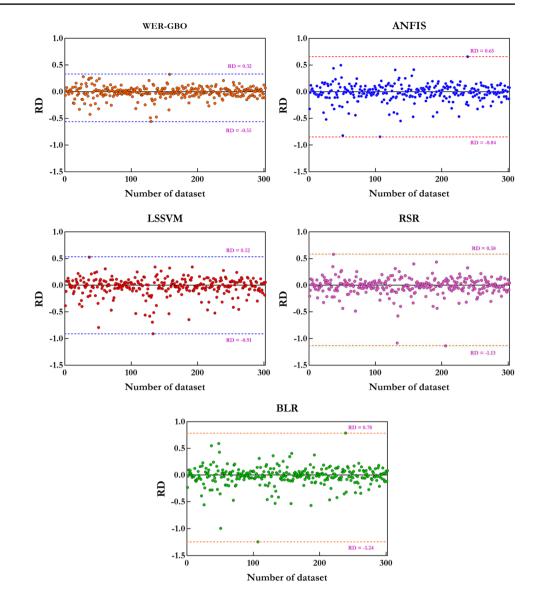
## Weighted exponential regression with gradient-based optimization

In this research, two new regression-based models, known as weighted linear regression (WLR) and weighted exponential regression (WER) models and optimized by the GBO algorithm for Na prediction, are introduced. These mentioned models are considered ensemble ones to utilize the most exact outcomes of ANFIS, LSSVM, BLR, and RSR models as inputs in order to forecast Na. To put it simply, the input parameters for WLR-GBO and WER-GBO models are in fact the predicted values gained by W-ANFIS-C4, W-LSSVM-C2, W-BLR-C5, and W-RSR-C2. Table 8 provides the data on statistical factors acquired by WLR-GBO and WER-GBO models for testing and training steps. Table 8 reports the WER-GBO outperforms according to the majority of important criteria for testing (R=0.972, RMSE=0.639, MAPE=5.885, KGE=0.948, and PI = 0.990) and training (R = 0.9813, RMSE = 0.9671, MAPE=7.7907, KGE=0.9731, and PI=0.987) phases. Thus, it can be concluded that the WER-GBO model is able to reach more meticulous and accurate Na prediction compared to the WLR-GBO model. By drawing the scatter plots of these two models (Fig. 13), it reveals that they can predict Na meticulously because the calculated values in both models are found near the 45° straight line of the scatter plots.

In Fig. 14, the time series plots of predicted and observed values of the WER-GBO and WLR-GBO are illustrated, with the figure showing a close agreement between predicted and measured values. Overall, by comparing the computational intelligence strategies, it can be gained that both models, WER-GBO

forecasting Na

Fig. 16 RD of all DI models in



and WLR-GBO, are concomitant with sufficient performance for both testing and training steps.

### Comparison of the performance of all DI models

The analysis of the model is conducted to detect the most efficient model. That is, DI models, namely ANFIS, LSSVM, BLR, RSR, and WER-GBO, are employed for Na prediction. From what has been gained by DI models, the WER-GBO is able to bring about more accurate prediction amongst other models during the training and testing steps. In addition, Fig. 15 displays the prediction outcomes' SDEs for diverse water quality predictors. In this regard, the SDEs in the WER-GBO model for training and testing steps stand out at 0.10 and 0.11, gaining the first rank. Indeed, it can be ratification for suitable stability of the WER-GBO model in Na prediction. Relative deviation (RD) analyzes the error to address the accuracy of the proposed model. The distribution of RD,  $RD = \frac{Na_M - Na_{DI}}{Na_M}$ , relative deviation, for predictive models during the testing stages was shown. In Fig. 16, it is apparent that the compression of error distribution in the WER-GBO model is far more than in other models. Besides, by constituting the relative deviation range of methods, it can be concluded WER-GBO model includes the minimum deviation range  $(-0.55 \le \text{RD} \le 0.32)$ , which leads to better performance than ANFIS  $(-0.84 \le \text{RD} \le 0.65)$ , LSSVM  $(-0.91 \le \text{RD} \le 0.52)$ , RSR  $(-1.13 \le \text{RD} \le 0.58)$ , and BLR  $(-1.24 \le \text{Er} \le 0.78)$  models.

The standard deviation and correlation coefficient (R), according to the Taylor diagram, are used to scrutinize the models' ability and performance, which elaborates the efficiency of models (Jamei et al. 2020). As can be seen in the diagram,

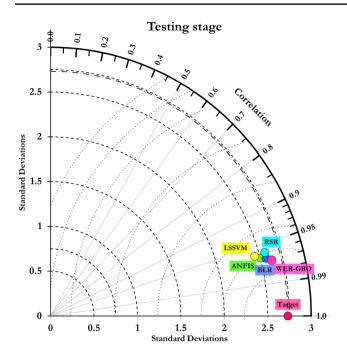


Fig. 17 Taylor diagram of all DI models for forecasting Na

by considering the standard deviation and *R*, there is a more compelling and tangible relationship between experienced and predicted Na. In Fig. 17, the Taylor diagram is depicted, while connecting the present monthly Na with all DI models. Hence, the best performance in the forecasting of Na is achieved by the WER-GBO model amongst other models while being the nearest model to the target.

## Conclusion

Prediction of water quality parameters using authoritative and precise models has a profound impact on saving time and cost. The main objective of this research is to develop two state-ofthe-art data intelligence methods based on the weighted exponential regression hybrid with GBO (WER-GBO) and the weighted linear regression hybrid with GBO (WLR-GBO) to forecast Na parameter. Indeed, these mentioned models utilize exponential and linear regression relationships with the Cauchy function as a weighted function to raise forecasting accuracy. Furthermore, the role of the GBO algorithm is to optimize the fundamental parameters in the WER model. In this regard, in the first step, in which the ANFIS, LSSVM, BLR, and RSR models are utilized, the proposed model is performed to reach the Na prediction. Consequently, the input variables in the WER-GBO model are the predicted values of Na gained by the mentioned DI models. In this research, the aim of developing a new ensemble model is to boost forecasting accuracy.

Thus, the ACF, PACF, and CC in this study are employed to identify the Na lags' number. Likewise, the measured time series of Na as well as Q on a monthly basis during the over 36 years in the Maroon River, located in southern Iran, are used in this study. The most appropriate subset regression employed in this process detects the input variable combinations' number. Because the relationship of Na and time series is nonlinear and sophisticated, the wavelet transform is utilized with three decomposition grades to maximize the prediction certainty in ANFIS, LSSVM, BLR, and RSR.

Monthly *Na* prediction is implemented in surface water by wavelet-DI models, using Dmey and Bior6.8 as two mother wavelets. Having integrated with ANFIS, LSSVM, BLR, and RSR models to forecast *Na*, Dmey confirmed the best and outstanding progress in the simulation accuracy grade, while Bior6.8 reached a suitable performance. The W-BLR-Dmey in Combo 5 gained striking ability in monthly *Na* prediction (R=0.968, RMSE=0.682, and KGE=0.931), followed by W-ANFIS-Dmey model in Combo 4 W-RSR-Dmey in Combo 2, and W-LSSVM-Dmey in Combo 2. Overall, the assessment of all DI-based models confirms that the supplementary model of W-BLR can accurately forecast the Na parameter.

The predicted values obtained by the mentioned DI models are applied for the input variables in WER-GBO and WLR-GBO models. From what has been gained as these models' outcomes, it can be concluded that WER-GBO outperforms WLR-GBO according to results of testing step as R = 0.9813, RMSE = 0.967, and KGE = 0.973 for training stage and R=0.9721, RMSE=0.9639, and KGE=0.948. Moreover, by considering SDE and RD as statistical errors, the proposed WER-GBO has the advantage over ANFIS, LSSVM, BLR, and RSR models thanks to reaching more meticulous and accurate Na prediction. Based on the Taylor diagram, the predicted value of Na via WER-GBO has more similar features such as standard deviation and correlation to the measured Na than other models. Hence, the ensemble WER-GBO-based method, gathering the benefits of all complementary techniques, can be a breakthrough in predicting water quality parameters, particularly surface water.

Authors contributions Iman Ahmadianfar, Seyedehelham Shirvani-Hosseini, Arvin Samadi-Koucheksaraee, and Zaher Mundher Yaseen contributed to data curation, formal analysis, methodology, investigation, visualization, writing—original draft, review and editing, draft preparation, resources, and software.

Data availability Data will be provided upon request from authors.

#### Declarations

**Ethical approval** The manuscript is conducted within the ethical manner advised by the Environmental Science and Pollution Research.

**Consent to participate** Not applicable

**Consent to publish** The research is scientifically consent to be published.

Competing interests The authors declare no conflict of interest.

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