



Modeling of an activated sludge process for effluent prediction—a comparative study using ANFIS and GLM regression

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Abstract In this paper, nonlinear system identification of the activated sludge process in an industrial wastewater treatment plant was completed using adaptive neuro-fuzzy inference system (ANFIS) and generalized linear model (GLM) regression. Predictive models of the effluent chemical and 5-day biochemical oxygen demands were developed from measured past inputs and outputs. From a set of candidates, least absolute shrinkage and selection operator (LASSO), and a fuzzy brute-force search were utilized in selecting the best combination of regressors for the GLMs and ANFIS models respectively. Root mean square error (RMSE) and Pearson's correlation coefficient (*R*-value) served as metrics in assessing the predicting performance of the models. Contrasted with the GLM predictions, the obtained modeling results show that the ANFIS models provide better predictions of the studied effluent variables. The results of the empirical search for the dominant regressors indicate the models have an enormous potential in the estimation of the time lag before a

desired effluent quality can be realized, and preempting process disturbances. Hence, the models can be used in developing a software tool that will facilitate the effective management of the treatment operation.

Keywords Wastewater treatment process modeling · Predictive models · ANFIS · Fuzzy exhaustive search · GLM regression · LASSO regularization

Introduction

Tightening environmental constraints have made it increasingly important for chemical process plants to be operated efficiently and in an environmentally friendly manner. Although several strategies have been put in place to combat the menace of water pollution; wastewater treatment still remains a major global challenge for process industries, as the performance of any wastewater treatment plant (WWTP) is affected by a distinct combination of some physical, chemical, and biological factors (Belanche et al. 2000; Mjalli et al. 2007; Singh et al. 2010).

Important water quality parameters frequently used to assess the performance of WWTPs include 5-day biochemical oxygen demand (BOD), chemical oxygen demand (COD), and total suspended solids (TSS). Frequent monitoring of these parameters helps plant operators preempt process disturbances and, hence, maintain environmental balance. This task often involves laborious and expensive laboratory analyses (Dupuit et al. 2007). Therefore, the determination of wastewater

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treatment indices using predictive models could help in a safe and cost-effective management of the treatment process (Belanche et al. 1999; Mjalli et al. 2007; Singh et al. 2010; Nadiri et al. 2018).

Wastewater treatment processes are highly nonlinear: involving strongly coupled physical, chemical, and biological activities. Hence, predictive models developed from mechanistic approaches are usually dimensionally complex, and the solutions are intractable, computationally intensive, and time-consuming (Mjalli et al. 2007; Araromi et al. 2014; Nadiri et al. 2018). Also, several assumptions often applied to simplify the resulting model introduce some degree of uncertainty which significantly affects the precision (Gontarski et al. 2000; Gernaey et al. 2004).

Contrariwise, soft computing techniques such as artificial neural network (ANN) and adaptive neuro-fuzzy inference system (ANFIS) have been found to be efficient in the formulation of high-performing models of complex nonlinear processes from historical data (El-Din and Smith 2002; Hamed et al. 2004; Mjalli et al. 2007). Singh et al. (2010) applied linear and nonlinear techniques to model a biological WWTP based on the effluent BOD and COD. The techniques are: partial least squares regression (PLSR), multivariate polynomial regression (MPR), and ANN. The authors concluded the nonlinear modeling techniques (MPR and ANN) provided better predictions.

Nasr et al. (2012) also developed an ANN model to predict the performance of a biological WWTP in Egypt using BOD, COD, and TSS as the characteristic parameters. The authors indicated the model sufficiently describes the nonlinear behavior of the WWTP and can be used as a valuable assessment tool by plant operators and management. In Mjalli et al. (2007), ANN models for the prediction of the effluent BOD, COD, and TSS of a local WWTP were formulated. The model was incorporated into a computer program to help operators monitor the WWTP performance and ensure compliance with discharge regulations. Pai et al. (2009) compared the performance of ANN and ANFIS in the prediction of the effluent suspended solids and COD of a hospital WWTP. It was reported that ANFIS outperforms ANN in the prediction exercises.

Though ANN is a robust tool in modeling highly nonlinear processes, it has some undesirable demerits. They are (1) the choice of network structure

and the problem of poor generalizations associated with its incorrectness; (2) the knowledge of the model is stored in synaptic weights and cannot be easily interpreted; and (3) large amount of data is needed to generate an adequate model. However, recent studies have proved that ANFIS overcomes these limitations, but the choice of model inputs still poses a challenge in ANFIS modeling (Passino and Yurkovich 1998; Hamed et al. 2004; Pai et al. 2009; Suh et al. 2009; Pai et al. 2011; Nasr et al. 2012).

Generalized linear model (GLM) regression is another peculiar class of nonlinear modeling techniques which has found applications in wastewater process modeling (Dürrenmatt and Gujer 2012). It uses random components to specify the distribution of the response variable and a smooth invertible linearizing link function to transform its expectation to a function of the regressors (Hardin and Hilbe 2007). GLM is an extension of the familiar regression models that make use of normal distribution as conditional distribution for response variables. Non-normal distributions used to extend the usual regression framework include Poisson, gamma, inverse-Gaussian, and binomial (McCullagh and Nelder 1989).

In previously published works, the model inputs were mostly selected arbitrarily. However, to obtain optimum results, it might help to make the choice via appropriate empirical methods (Tomić et al. 2018). For instance, Ahmadi et al. (2018) investigated the influence of model inputs selected using different combinations of methods which include principal component analysis, gamma test, and correlation model on the performance of a BOD model. The authors reported the choice of inputs have significant effects on the model prediction.

In this study, a procedure for the nonlinear dynamical system identification of a biological wastewater treatment process using ANFIS and GLM regression is described and exemplified. Models for the prediction of the effluent BOD and COD were both developed from past inputs and output variables. For each model, a brute-force search also known as exhaustive search was employed in selecting the most adequate combination of the variables for the ANFIS models while a penalized regression method termed *least absolute shrinkage and selection operator* (LASSO) was used for the GLMs.

Materials and methods

The case study

The data used in this study was collected from the WWTP of Seven-Up Bottling Company in Lagos, Nigeria. This company is one of the largest beverage manufacturing industries in the country. The company produces different brands of soft drinks which are widely consumed in the country. The plant depends on a large volume of water in its daily operations which include, but not limited to, the preparation of feedstock and purification of packaging bottles.

The process wastewater contains oil, solid objects, suspended particles, dissolved chemicals, and organic substances. The wastewater is disposed of in the treatment operation cycle before discharged into the main water body in the environment. The treatment objective is to remove or reduce objectionable materials to a level that conforms to environmental regulations and standards which limits the COD and 5-day BOD of effluent discharges into surface waters and drainages to 30 and 6 mg/l, respectively.

The treatment process consists of three main stages, viz. physical, chemical, and biological treatment processes. The biological treatment step is an activated sludge system where dissolved biodegradable organic matter is degraded by micro-organisms to an acceptable and ecologically safe limit. This is a typical example of a process whose dynamics is difficult to fully comprehend and manage due to perturbations in flow, organic load, and concentration of micro-organisms per unit time (Pons et al. 1999; Gernaey et al. 2004; Moral et al. 2008; Cristea et al. 2009).

Five-day BOD and COD are the organic content indicators of the wastewater in this case study. These parameters are monitored through sampling and laboratory analyses before and after treatment. Some of the analyses involve the use of hazardous reagents such as mercury sulfate, chromium trioxide, sulfuric acid, and potassium dichromate. They are exacting and requiring sophisticated laboratory equipment and gadgets. Also, the determination of COD can take few hours while BOD 5 days. For these reasons, accurate predictive models developed from the historical data can facilitate real-time monitoring and effective management of the WWTP. Hence, it will be of important use in maintaining environmental balance and ensuring the safeness of

water bodies (Dürrenmatt and Gujer 2012; Heddam et al. 2016).

This study is primarily concerned with the activated sludge system. Three hundred fifty-nine datasets were collected over the period of a year, as seasonal changes were considered important in capturing possible variations in the studied variables (Mjalli et al. 2007; Nasr et al. 2012; Heddam et al. 2016). The collected data include the influent and effluent BOD and COD recorded on a daily basis, and cover the peak (dry season) and the lowest period (rainy season) of the company's products demand.

Data preprocessing

To produce models that satisfactorily map-out the process behavior, the data was preprocessed to eliminate some likely unusable information which are probably introduced into the WWTP database as a result of measurement errors (Oliveira-Esquerre et al. 2002; Hamed et al. 2004; Rustum and Adeloje 2007; Yel and Yalpir 2011; Tangirala 2015). Firstly, the dataset was subjected to a statistical tool to sort out and remove outliers which were classified as data-points lying outside $\pm 2\sigma$, standard deviation of the group mean (Mjalli et al. 2007). This procedure reduced the dataset to 324 data-points.

Also, for adequate process representation, the presence of noise in the data is undesirable as it causes non-smoothing in the data trend which is counterproductive. Thus, a data smoothing technique was applied to reduce noise and normalize gaps caused by the removal of the outliers (Rustum and Adeloje 2007). In the smoothing process, each data-point was replaced with the average of the neighboring data-points defined within a span using moving average filter which is mathematically expressed as:

$$y_s(i) = \frac{1}{2N+1} (y(i+N) + y(i+N-1) + \dots + y(i-N)) \quad (1)$$

In Eq. (1), $y_s(i)$ is the smoothed value of the i -th data-point, N is the number of neighboring data-points on either side of $y_s(i)$, and $2N+1$ is the span. Most researchers use three-, four-, or five-point moving averages (span) (Mjalli et al. 2007). In this study, five-point moving averages (span = 5) was used because it gave satisfactory result when compared to others. The statistical information of the preprocessed dataset is presented in Fig. 1.

Brief description of system identification

This is the process of building models of dynamic systems from observed input–output data. It involves the use of the past inputs and outputs, known as regressors, as model inputs (Passino and Yurkovich 1998; Tangirala 2015). A linear mapping of this form is defined as:

$$y(k) = \sum_{i=1}^q \theta_{ai}y(k-i) + \sum_{i=0}^p \theta_{bi}u(k-i-1) \tag{2}$$

In Eq. (2), $y(k)$ and $u(k)$ are the system output and input variables at time interval $k \geq 0$ respectively. The number of the past output and input variables used are represented by q and p , respectively. The parameters associated with the past output and input variables are θ_a and θ_b , respectively. When Eq. (2) is expressed in a matrix notation, it becomes:

$$y(k) = f(x|\theta) = \theta^T x \tag{3}$$

Where,

$$\left. \begin{aligned} x &= [y(k-1), \dots, y(k-q), u(k), \dots, u(k-p-1)]^T \\ \theta &= [\theta_{a1}, \dots, \theta_{aq}, \theta_{b0}, \dots, \theta_{bp}]^T \end{aligned} \right\} \tag{4}$$

In Eqs. (3) and (4), x is the regression vector, and θ is the vector of unknown parameters to be estimated. Each vector is of dimension $N \times 1$ where $N = q + p + 1$. System identification involves adjusting θ making use of the information contained in the observed data until the error between the predicted and observed values is reduced to an acceptable tolerance. Fuzzy identification follows the same procedure by utilizing the predefined regression vector x , but with functional capabilities which enable it to achieve more accurate identification of nonlinear systems than the linear map defined in Eq. (2) (Passino and Yurkovich 1998; Araromi et al. 2014).

ANFIS architecture

Basically, neuro-fuzzy system, which only supports Takagi–Sugeno–Kang (TSK) model, is a fuzzy inference system (FIS) constructed in the framework of a neural network. It employs the easy interpretability and expansiveness of fuzzy logic and the robust learning ability of artificial neural networks as it extracts fuzzy rules from observed data into a rule-base (Passino and Yurkovich 1998; Guillaume 2001; Karray and De Silva 2004; Mingzhi et al. 2009; Nadiri et al. 2018). A rule in

a first-order TSK (with consequent linear function) is as expressed in Eq. (5).

$$\begin{aligned} p_{th} : & \text{IF } X_1 \text{ is } A_1^p \text{ AND } X_2 \text{ is } A_2^p \dots \text{ AND } X_n \text{ is } A_n^p \text{ THEN } O_p \\ & = \alpha_0^p + \alpha_1^p X_1 + \dots + \alpha_n^p X_n \end{aligned} \tag{5}$$

In Eq. (5), X_i is the i -th input linguistic variable of the p -th rule. The corresponding linguistic value is A_i^p and n is the number of input linguistic variables. A_i^p is linked to a membership function (MF) $\mu_{A_i^p}(X_i)$. In each rule consequent, O_p denotes the output of the inference process, and $\alpha_0^p, \dots, \alpha_n^p$ are Sugeno parameters. For a zero-order TSK, each rule consequent is a constant (α_0^p) as other Sugeno parameters are set to zero. Moreover, each rule in ANFIS has unity weight and one output.

Fig. 2 shows the structure of a first-order TSK model formulated as a five-layer feed-forward network. The model in the figure has two inputs, four rules, and an output which is determined by weighted average defuzzification method. Each layer is a step in a fuzzy identification process and their functionality is described as follows (Guillaume 2001; Karray and De Silva 2004):

Layer 1: Fuzzification

Each node i , characterized by a MF, converts the crisp inputs into membership values.

$$O_i^1 = \mu_{A_i^p}(X) \tag{6}$$

Layer 2: Rule nodes

To evaluate the degree of fulfillment W_p of each rule, T-norm, and T-conorm operators are used to express the AND and OR connectives respectively. An example of a T-norm operation is:

$$\begin{aligned} W_p &= \mu_{A_i^p}(X_1) \otimes \mu_{A_j^p}(X_2), \quad i = 1, 2; j = 1, 2; p \\ &= 1, \dots, 4 \end{aligned} \tag{7}$$

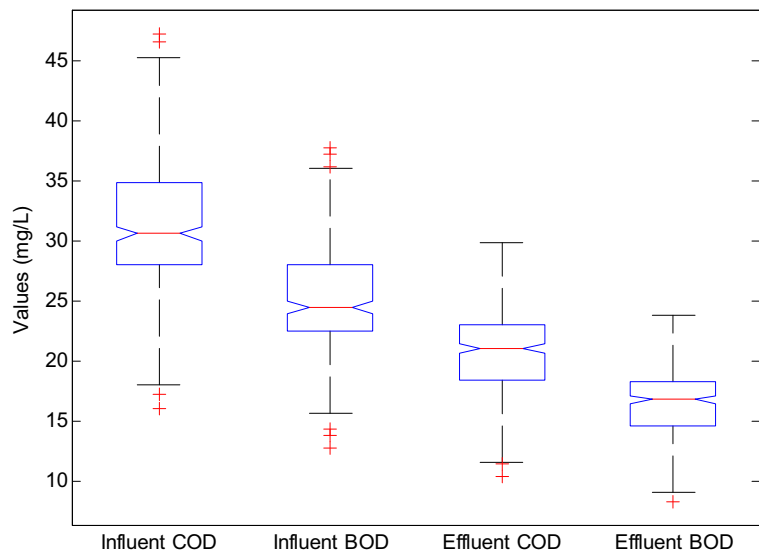
Layer 3: Normalization

The normalized firing strength of the p -th rule is calculated as the ratio of its firing strength to the sum of all rules firing strengths.

$$\bar{W}_p = \frac{W_p}{\sum_{p=1}^4 W_p} \tag{8}$$

Layer 4: Rule consequent

Fig. 1 Statistical features of the modeled data



These nodes evaluate the consequent of each fuzzy rule. The values of the consequents are multiplied by normalized firing strengths.

$$O_p^4 = \bar{W}_p O_p = \bar{W}_p (\alpha_0^p + \alpha_1^p X_1 + \alpha_2^p X_2) \tag{9}$$

Layer 5: Summation

The final output is calculated as the sum of all incoming signals.

$$O^5 = \sum_{p=1}^4 \bar{W}_p (\alpha_0^p + \alpha_1^p X_1 + \alpha_2^p X_2) \tag{10}$$

Regressors formulation and selection

In this work, two multi-input single-output models were identified from the collected process data. One is for the prediction of the effluent COD (model 1) while the other is for the effluent BOD (model 2). In each case, four past outputs and six past inputs, organized into three groups, were considered as model input candidates as expressed in Eq. (11) where $u_1(k)$ and $u_2(k)$ depicts the influent BOD and COD respectively. And the effluent BOD or COD is represented by $y(k)$.

$$\mathbf{x} = \begin{cases} y(k-i), & 1 \leq i \leq 4 \\ u_1(k-j), & 1 \leq j \leq 6 \\ u_2(k-j) \end{cases} \tag{11}$$

It is quite difficult to determine how to select exogenous variables for adequate model approximation and the search for the combination of the input candidates which influences the output the most is a major challenge (Passino and Yurkovich 1998; Ahmadi et al. 2018;

Tomić et al. 2018). A fuzzy brute-force search was used to select the ANFIS models inputs while LASSO was used for the GLM regressors. The search space is exponential in size and the task is to find $\mathbf{x} \in \mathbf{X}$ which satisfies a condition ψ . This exhaustive search method iterates through all the elements in the search space, testing every possible candidate solution. It builds an ANFIS model for each combination, trains it for one epoch, and reports its performance.

Since it is logical that the regressors should not exclusively be from any of the process inputs or outputs, one regressor each was selected from the three sets highlighted in Eq. (11) (Passino and Yurkovich 1998; Tangirala 2015). This step was separately performed for the two models and 144 combinations were generated in each case as shown in Eq. (12):

$${}^4_1 C * {}^6_1 C * {}^6_1 C = 144 \tag{12}$$

Fuzzy exhaustive search algorithm identifies the combination with the least training RMSE. But in order to obtain models with minimal error and no over-fitting, two other criteria were considered simultaneously. They are low validation RMSE and relatively low positive difference between the training and validation RMSE. Figs. 3 and 4 show the exhaustive search results for ANFIS models 1 and 2 respectively. Incidentally, the combination of regressors $y(k-1)$, $u_1(k-6)$, and $u_2(k-5)$ satisfied these conditions for both models and thus stands out as the required regression vector. A rule in a first-order Sugeno model considered in this study is:

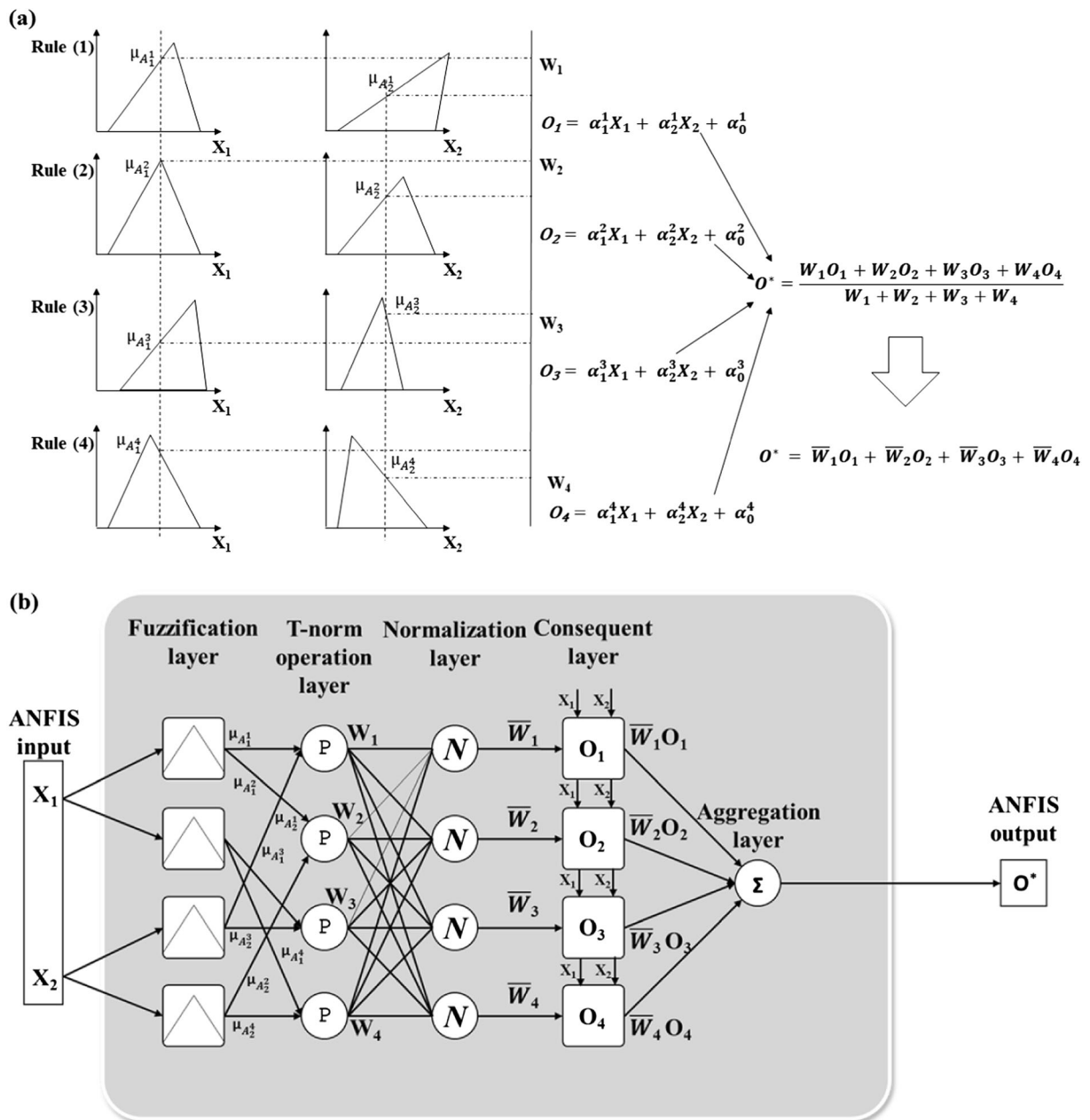


Fig. 2 Five-layer ANFIS architecture: **(a)** schematic representation of weighted average defuzzification method; **(b)** overview of ANFIS

$$\begin{aligned}
 p_{th} : & \text{IF } y(k-1) \text{ is } A_1^p \text{ AND } u_1(k-6) \text{ is } A_2^p \text{ AND } u_2(k-5) \text{ is } A_3^p \text{ THEN } O_p \\
 & = \alpha_0^p + \alpha_1^p y(k-1) + \alpha_2^p u_1(k-6) + \alpha_3^p u_2(k-5)
 \end{aligned}
 \tag{13}$$

For clear observation, only the combinations having comparatively low RMSE values are reported in the plots. In Fig. 3, it is observed that the aforesaid combination has the least validation error and a relatively low training error. The difference between the error values, which is illustrated as the distance between both points on the plot, is comparatively low. This scenario is also observed in the regressor selection process for ANFIS model 2 (Fig. 4).

Notably, the identified regressors reveal the time interval between a change in the process inputs and the first significant change in the response. In case of a process disturbance and tightening environmental constraints, this information can aid the WWTP operators in approximating the time needed to realize a set treatment quality before the treated wastewater is discharged into the environment. Eq. (13) shows that at any instant in time (day, k) the value of the effluent five-day BOD or COD, $y(k)$, can be obtained from $y(k - 1)$, $u_1(k - 6)$, and $u_2(k - 5)$ before new measurements are obtained. Since the measurement process takes time, this implies that by employing the described model inputs selection method, it is possible to approximate future effluent conditions based on measured past variables. The plant managers can also detect potential process disturbances, at least, a day earlier. In such cases, a number of activated sludge process variables, such as the hydraulic or sludge retention time and mixing rate, can be adjusted to control the situation and avoid a likely shutdown.

Fuzzy identification

Neuro-fuzzy systems are developed in two successive stages: structure and parameter learning. In the structure learning stage, a set of input–output numerical data is partitioned to define the rules structure. Each partition represents a rule and their boundaries overlap. Common partitioning techniques include grid-type and clustering. In grid partitioning, with n inputs and k input MFs, the number of rules is k^n . Though the number of rules increases exponentially, grid partitioning is effective in extracting all the rules for a small number of inputs (Guillaume 2001; Karray and De Silva 2004).

Grid partitioning was used in this study and two input MFs were affixed to each regressor. The number of MFs to be associated with each model input was determined by trial-and-error (Mjalli et al. 2007; Jang 1993). It was observed that any value higher than two increased the models' complexity and returned a very high training

and validation error. Thus, the number of rules generated is $k^n = 2^3 = 8$. Figure 5 shows the adopted FIS structure.

In the second stage, the values of the connection weights and MFs parameters are adjusted and optimized using learning algorithms. The hybrid learning algorithm proposed by Jang (1993) was adopted in this work: back-propagation method which is based on gradient-descent optimization technique was employed to tune the antecedent parameters while least squares method was used for the consequent part.

The preprocessed data was divided into two: one-half was used for model training and the other for validation. Because correct ANFIS model specifications are determined empirically, the effects of five different input MF types on the predicting performance of zero- and first-order Sugeno model structures was examined (Jang and Sun 1995; Guillaume 2001; Pai et al. 2011). The essence was to identify the model specifications required to realize a satisfactory performance. The considered input MFs are presented in Table 1.

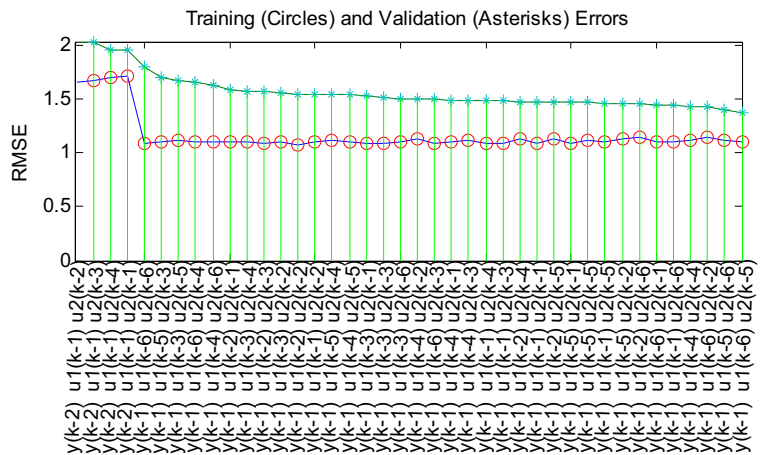
Generalized linear model (GLM) regression with LASSO regularization

GLM regressions extend linear model regressions by relaxing the assumption of linearity in the parameters with the introduction of a link function and the use of error distributions other than normal distribution. As shown in Eq. (14), the model is a function of mean μ with a linear combination $x\beta$ formed from regressors x and coefficient vector β . The mean μ_i of the response variable y_i is modeled as a monotonic nonlinear transformation of a linear function (η_i) of the regressors. For each set of observation, the linearizing invertible link function $g(\cdot)$ transforms the expected value of the response $\mu_i = E(y_i)$ to the linear predictor such that $\mu_i = g^{-1}(\eta_i) = E(y_i)$ (McCullagh and Nelder 1989; Hardin and Hilbe 2007).

$$g(\mu_i) = \eta_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} + \dots + \beta_k x_{ik} \tag{14}$$

The choice of the linearizing transformation which is partially separated from the distribution of the response is an advantage in GLM regression. Common GLM link functions include: identity, natural logarithm, inverse-square, inverse, square root, logit (sigmoid), and probit. The identity link function, as the name implies, returns

Fig. 3 Selection of regressors for ANFIS model 1



its argument unchanged such that $\mu_i = g^{-1}(\eta_i) = \eta_i = E(y_i)$.

Least absolute shrinkage and selection operator (LASSO) is a regularization method used with GLM regression to identify and select dominant regressors amongst redundant ones, and produce shrinkage estimates with potentially low prediction errors than the regular least squares. LASSO approach minimizes the prediction error subject to a non-differentiable constraint presented in terms of the L_1 -norm of the coefficients (Tibshirani 1996). A nonnegative regularization parameter λ , known as penalty, is introduced to constrain the size of the estimated coefficients and solve the objective function in Eq. (15) where N represents the number of data-points, β_0 the intercept and β the p -vector of the coefficients. D is the estimated expected deviance of the model applied to new data as evaluated by cross-validation. As λ increases, the number of nonzero elements in β decreases.

$$\min_{\beta_0, \beta} \left(\frac{1}{N} D(\beta_0, \beta) + \lambda \sum_{j=1}^p |\beta_j| \right) \quad (15)$$

Tibshirani (1996) proposed a k -fold cross-validation algorithm for the estimation of the prediction error in order to compute the best λ . In this method, LASSO is indexed in terms of λ and the prediction error is calculated over a range of values of λ from 0 to 1 inclusive. The value of λ which gives the least prediction error is selected. This method is often used for small parameters vector space. Tenfold cross-validation was used in this work. Figure 6 shows the cross-validated deviance of the LASSO fit. It is a plot of the absolute deviance against $\log \lambda$. The broken vertical line with the blue circle indicates the λ with the minimum deviance while the longest continuous vertical line with the green circle locates the minimum deviance plus one standard deviation.

Fig. 4 Selection of regressors for ANFIS model 2

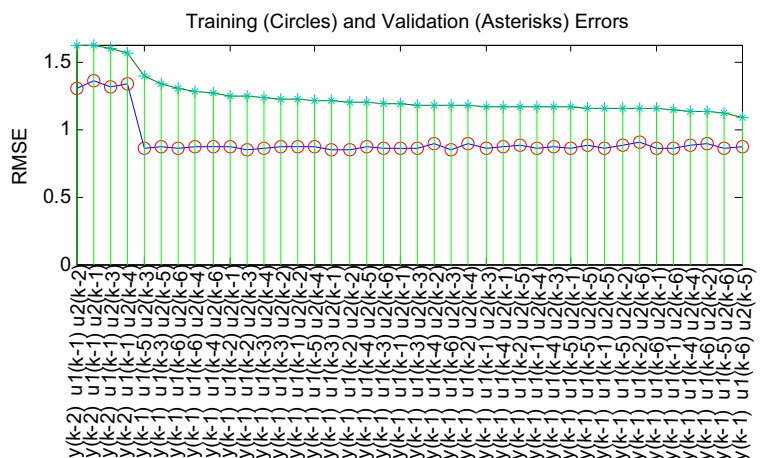


Fig. 5 Adopted ANFIS model structure

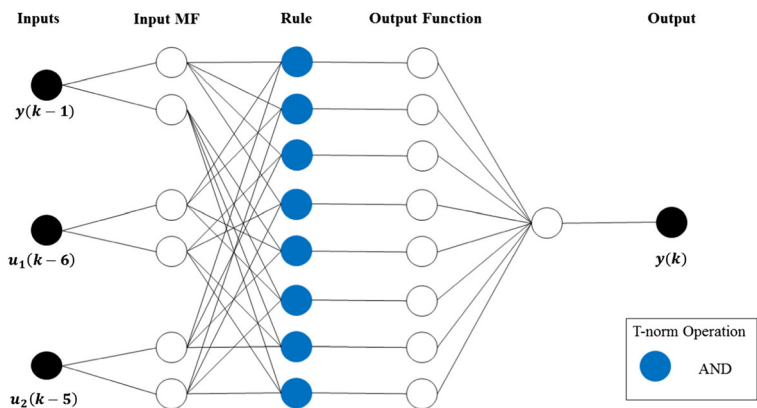


Figure 7 displays the nonzero model coefficients plotted as a function of $\log \lambda$. Each curve on the plot represents each of the 16 model input candidates (regressors) highlighted in Eq. (11). As observed, LASSO sets more coefficients to zero while few nonzero ones remain as λ increases. The regressors having nonzero model coefficients at the minimum deviance plus one standard deviation point are $y(k-1)$, $u_1(k-6)$, $u_2(k-5)$, and $u_2(k-6)$. The LASSO regularization was also separately performed for both models and the same regressors suggested by the fuzzy search, with the exception of $u_2(k-6)$, were obtained.

In the estimation of the GLMs, the responses were considered to have inverse-Gaussian distribution. The link functions used are identity, logarithmic, square root, inverse, and square. The inverse-Gaussian distribution,

otherwise known as Wald distribution, can be used to model nonnegative positively skewed data. Though this distribution emanated from the theory of Brownian motion, it has been utilized in modeling different phenomena (Jablonski et al. 2013). The density function is presented in Eq. (16) where μ is the expected value of the response and λ is the inverse of the dispersion parameter.

$$p(y) = \sqrt{\frac{\lambda}{2\pi y^3}} \exp\left[-\frac{\lambda}{2\mu^2 y}(y-\mu)^2\right] \text{ for } y > 0 \quad (16)$$

Models predicting performance metrics

RMSE and Pearson’s correlation coefficient (R -value) were used to statistically assess the predicting performance of the models: how well the predicted data approximates the modeled raw data. These metrics are mathematically expressed as:

$$RMSE = \sqrt{\frac{\sum_{i=1}^n (X_{\text{obsv},i} - X_{\text{pred},i})^2}{n}} \quad (17)$$

$$R\text{-value} = \frac{\sum_{i=1}^n (X_{\text{obsv},i} - \bar{X}_{\text{obsv}}) \cdot (X_{\text{pred},i} - \bar{X}_{\text{pred}})}{\sqrt{\sum_{i=1}^n (X_{\text{obsv},i} - \bar{X}_{\text{obsv}})^2 \cdot \sum_{i=1}^n (X_{\text{pred},i} - \bar{X}_{\text{pred}})^2}} \quad (18)$$

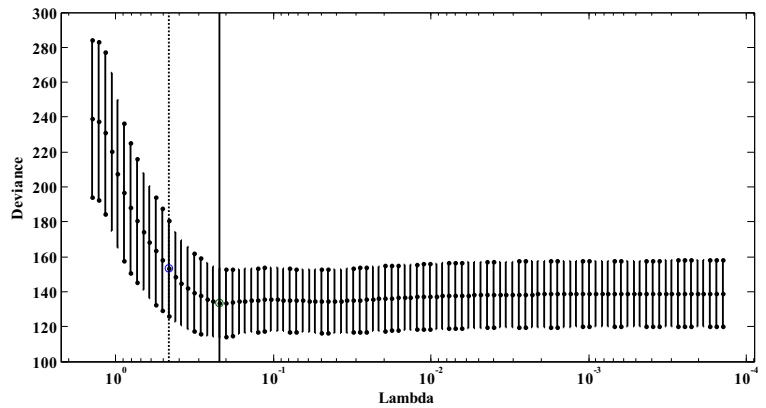
Where X_{obsv} is the measured or observed value, X_{pred} is the predicted value, and n is the number of observations. \bar{X}_{obsv} and \bar{X}_{pred} are the means of the observed and predicted values respectively. The reliability of the model depends on the closeness of the RMSE to zero and R -value to + 1 (Gontarski et al. 2000; Mjalli et al. 2007; Pai et al. 2011; Dürrenmatt and Gujer 2012; Heddard et al. 2016; Oke et al. 2017).

Table 1 Mathematical expressions of the considered MFs

Input MF	Notation	Membership value $[\mu_{A_i^p}(X)]$
Triangular	trimf	$\max\left(\min\left(\frac{X-a_i}{b_i-a_i}, \frac{X-a_i}{b_i-a_i}\right), 0\right)$
Trapezoidal	trapmf	$\max\left(\min\left(\frac{X-a_i}{b_i-a_i}, 1, \frac{X-a_i}{b_i-a_i}\right), 0\right)$
Generalized bell	gbellmf	$\frac{1}{1 + \left \frac{X-c_i}{a_i}\right ^{2b_i}}$
Gauss (one-sided)	gaussmf	$\exp\left(-\frac{(X-c_i)^2}{2\sigma_i}\right)$
Gauss (two-sided)	gauss2mf	$\exp\left(-\frac{(X-c_i)^2}{\sigma_i}\right)$

$a_i, b_i, c_i, d_i,$ and σ_i are the MFs parameters

Fig. 6 Tenfold cross-validated LASSO fit



Results and discussion

ANFIS modeling

As earlier stated, model 1 is the predictive model of the effluent COD while model 2 denotes that of the effluent BOD. Table 2 presents the predicting performance of ANFIS model 1 for different combinations of input and output MFs. Although the least difference between the training and validation errors was achieved when *trimf* and *linear* were used as the MFs and the corresponding RMSE values are also relatively low; it is observed that the *R*-value (0.947) is a bit lower than that obtained when *gbellmf* and *constant* were used as the input and output MFs respectively. Also, the *gbellmf* and *constant* MF combination produced the highest *R*-value (0.948). Figure 8 shows how the errors change as the training proceeds. It illustrates that convergence was achieved for both the training and validation errors before the 200th epoch. The resulting training (1.130) and validation RMSE (1.302) are also comparatively low and the positive difference between both does not suggest

model over-fitting. Therefore, the combination of *gbellmf* and *constant* MF types suitably identify the relationship that exists between the modeled variables for ANFIS model 1.

The predicting performances of ANFIS model 2 for different combinations of input and output MFs are presented in Table 3. From the table, the highest *R*-value (0.947) was realized in four different modeling scenarios. Two of these cases are significant because they gave lower validation errors. They are *trimf* and *linear*, and *gbellmf* and *constant* MFs combination, but the positive difference between the training (0.899) and validation RMSE (1.041) for the *gbellmf* and *constant* MFs combination is lower when compared to that obtained in the *trimf* and *linear* MFs combination. This implies less over-fitting in the resulting model, thus, making it the model configuration with the best predicting performance for ANFIS model 2. The training progress for this case is presented in Fig. 9. It shows the errors also converged before the 200th epoch. This informs that the data was not over-fitted and, hence, the produced model generalizations are reliable.

Fig. 7 Trace plot of coefficients fits by LASSO

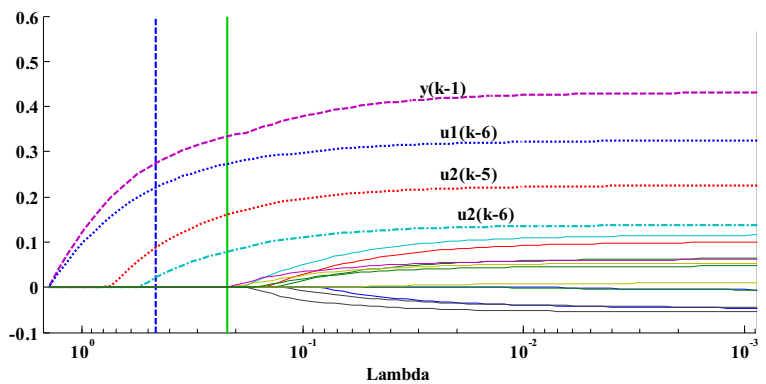


Table 2 Predicting performance using different MFs for ANFIS model 1

Input MF	RMSE				R-value	
	Output function				Output function	
	Constant		Linear		Constant	Linear
	Training	Validation	Training	Validation		
trimf	1.164	1.310	1.129	1.309	0.946	0.947
trapmf	1.152	1.336	1.097	1.359	0.945	0.946
gbellmf	1.130	1.302	1.094	1.361	0.948	0.946
gaussmf	1.178	1.329	1.092	1.355	0.944	0.946
gauss2mf	1.173	1.341	1.112	1.372	0.944	0.945

GLM regression

The influence of different transformations (link functions) on the predicting performance of the two GLMs was studied. The result of the performance evaluation is presented in Table 4. The maximum R-value (0.637) and

the least errors were realized when no transformation (identity link) was applied to the responses. This is consistent with that obtained in ANFIS modeling, where the same model specifications yielded the best predicting performance for both models. The GLM regression is expressed as:

$$\begin{aligned}
 y(k) = & \beta_0 + \beta_1 \cdot y(k-1) + \beta_2 \cdot u_1(k-6) + \beta_3 \cdot u_2(k-5) + \beta_4 \cdot u_2(k-6) + \beta_5 \cdot y(k-1)^2 + \beta_6 \cdot u_1(k-6)^2 \\
 & + \beta_7 \cdot u_2(k-5)^2 + \beta_8 \cdot u_2(k-6)^2 + \beta_9 \cdot y(k-1) \cdot u_1(k-6) + \beta_{10} \cdot y(k-1) \cdot u_2(k-5) + \beta_{11} \cdot y(k-1) \cdot u_2(k-6) \\
 & + \beta_{12} \cdot u_1(k-6) \cdot u_2(k-5) + \beta_{13} \cdot u_1(k-6) \cdot u_2(k-6) + \beta_{14} \cdot u_2(k-5) \cdot u_2(k-6)
 \end{aligned}
 \tag{19}$$

Fig. 8 Training progress for ANFIS model 1

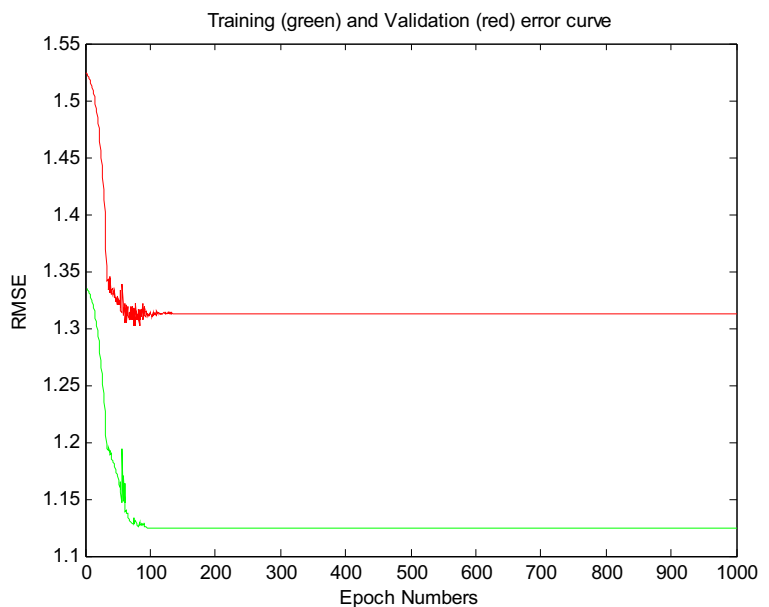


Table 3 Predicting performance using different MFs for ANFIS model 2

Input MF	RMSE				R-value	
	Output function				Output function	
	Constant		Linear		Constant	Linear
	Training	Validation	Training	Validation		
trimf	0.918	1.046	0.893	1.043	0.946	0.947
trapmf	0.934	1.074	0.864	1.086	0.944	0.946
gbellmf	0.899	1.041	0.860	1.081	0.947	0.947
gaussmf	0.930	1.060	0.861	1.078	0.945	0.947
gauss2mf	0.950	1.101	0.903	1.101	0.941	0.944

For both models, the estimated coefficients and their statistics are presented in Table 5. At 5% significance level, the intercept (β_0) and all the LASSO selected regressors, except $u_2(k-6)$, are significant, as the p -values $\ll 0.05$. This observation is also in agreement with the fuzzy brute-force search result where $y(k-1)$, $u_1(k-6)$, and $u_2(k-5)$ were selected as the most influential regressors. However, the R -values and RMSE values reported in Table 4 are undesirable. When these values are compared to those obtained in ANFIS modeling, it can be provisionally said that the ANFIS models perform better than the GLMs.

Significance of the results

The results obtained in this work are consistent. The same regressors were identified in the fuzzy exhaustive search and LASSO regularization processes and identical specifications produced the desired predicting performance in both models and modeling approaches. This is largely due to the modeled data structure, and affirms that the reliability of a black-box model hinges on the correctness of the data. It is noteworthy that appropriate model specifications are determined empirically which is a time-consuming process for higher number of inputs as it necessitates testing different combinations of the variables before arriving at one that

Fig. 9 Training progress for ANFIS model 2

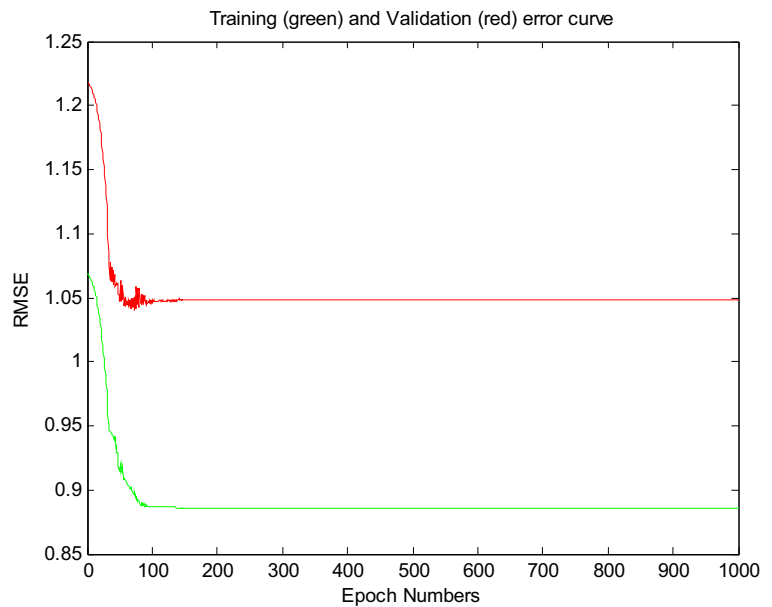


Table 4 GLM regression results

Transformation	Model 1		<i>R</i> -value	Model 2		<i>R</i> -value
	RMSE			RMSE		
	Training	Checking		Training	Checking	
Identity	2.895	3.042	0.637	2.307	2.415	0.637
Natural logarithm	2.915	3.083	0.631	2.319	2.441	0.632
Inverse	2.984	3.193	0.615	2.369	2.517	0.619
Square	2.929	3.090	0.629	2.338	2.466	0.626
Square root	2.899	3.055	0.635	2.309	2.422	0.635

gives the desired model performance. In this study, three input variables were used to adequately model a highly nonlinear behavior and the adopted model structure achieved this within a reasonably short simulation time, circumventing the curse of dimensionality peculiar to grid-partitioning techniques in ANFIS modeling (Guillaume 2001; Karray and De Silva 2004; Jang 1993).

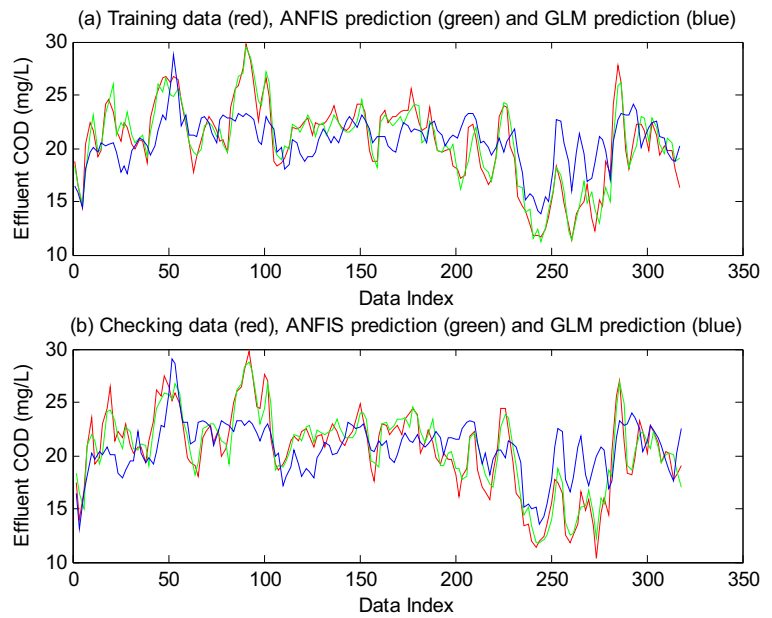
How good is a fit? One obvious metric is how close it is to the measured data-points. For visual inspection, fits of the measured data and model predictions of the effluent COD and BOD are presented in Figs. 10 and 11 respectively. For both model training and validation

datasets, the figures show the ANFIS predictions are closer to the corresponding measured values than the GLM predictions. Figure 12a, b, c, d illustrates the degree of linear relationship between the measured data and the corresponding predictions of GLM 1, GLM 2, ANFIS model 1, and ANFIS model 2 respectively. The linear fits reveal that the measured data and ANFIS predictions are in satisfactory agreement. These show the ANFIS models outperform the GLMs. The results also illustrate that wastewater treatment processes, such as this case study, are highly nonlinear and cannot be effectively modeled using mechanistic approaches or linear model structures (Oliveira-Esquerre et al. 2002;

Table 5 GLMs estimated coefficients and their statistics

Coefficient	Model 1				Model 2			
	Estimate	Standard error	<i>t</i> -statistics	<i>p</i> -value	Estimate	Standard error	<i>t</i> -statistics	<i>p</i> -value
β_0	-2.936	0.509	-5.766	8.089e-09	-1.326	0.567	-4.786	1.080e-07
β_1	0.249	0.607	4.099	4.150e-05	0.257	0.676	5.299	2.102e-06
β_2	0.355	0.633	4.028	5.616e-05	0.138	0.656	4.018	1.696e-08
β_3	0.529	0.542	4.231	2.320e-05	0.543	0.533	5.431	1.019e-05
β_4	0.488	0.503	0.969	0.332	0.128	0.603	0.899	0.602
β_5	0.013	0.621	0.990	0.322	0.213	0.672	0.879	0.702
β_6	-1.776	0.551	-5.036	4.740e-07	-2.656	0.551	-6.236	1.740e-06
β_7	-1.379	0.582	-4.083	4.436e-05	-1.324	0.575	-4.183	4.136e-06
β_8	0.778	0.577	1.347	0.178	0.718	0.582	3.247	0.778
β_9	0.615	0.621	0.990	0.322	0.934	0.571	0.800	0.923
β_{10}	-0.070	0.505	-0.140	0.888	-0.168	0.602	-0.210	0.801
β_{11}	0.021	0.576	3.506	4.537e-04	0.121	0.726	4.006	1.530e-04
β_{12}	0.795	0.559	1.423	0.155	0.495	0.537	1.523	0.113
β_{13}	1.268	0.554	2.287	0.022	0.668	0.532	2.587	0.122
β_{14}	1.780	0.543	3.278	0.104	2.780	0.556	4.008	0.234

Fig. 10 Comparison of the model predicted and measured effluent COD values: (a) training data; (b) validation/checking data



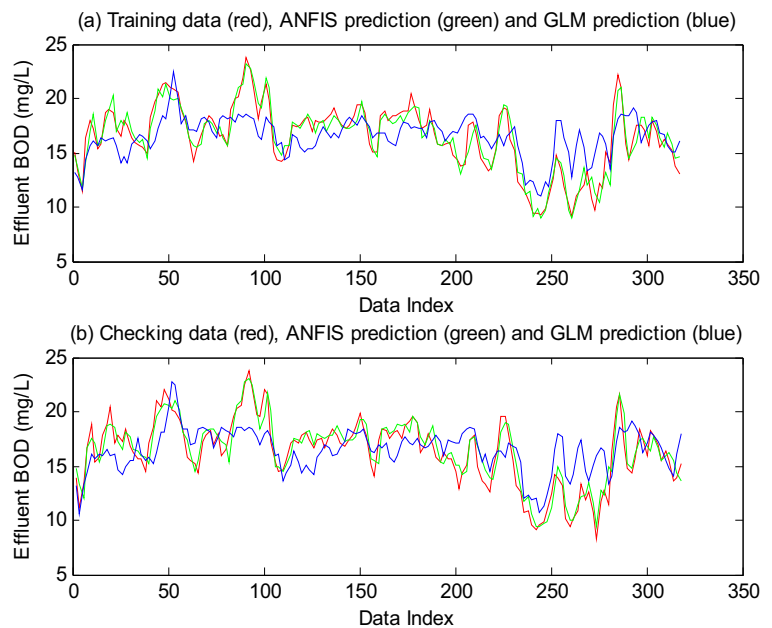
Gernaey et al. 2004; Moral et al. 2008; Singh et al. 2010; Heddam et al. 2016; Oke et al. 2017).

Conclusions

The application of mechanistic approaches in identifying the behavior of nonlinear processes is tasking. But ANFIS has yet proved its capability as an efficient

model approximating tool by adequately mapping-out a highly nonlinear biological wastewater treatment process based on previously measured inputs and output values of the studied variables. ANFIS predicted the process outputs with greater accuracy when compared with the GLM regression models. The data used in the study were pretreated to remove outliers and reduce noise as the predictive performance of black-box models depends on the exactness of data. Also, the influential

Fig. 11 Comparison of the model predicted and measured effluent BOD values: (a) training data; (b) validation/checking data



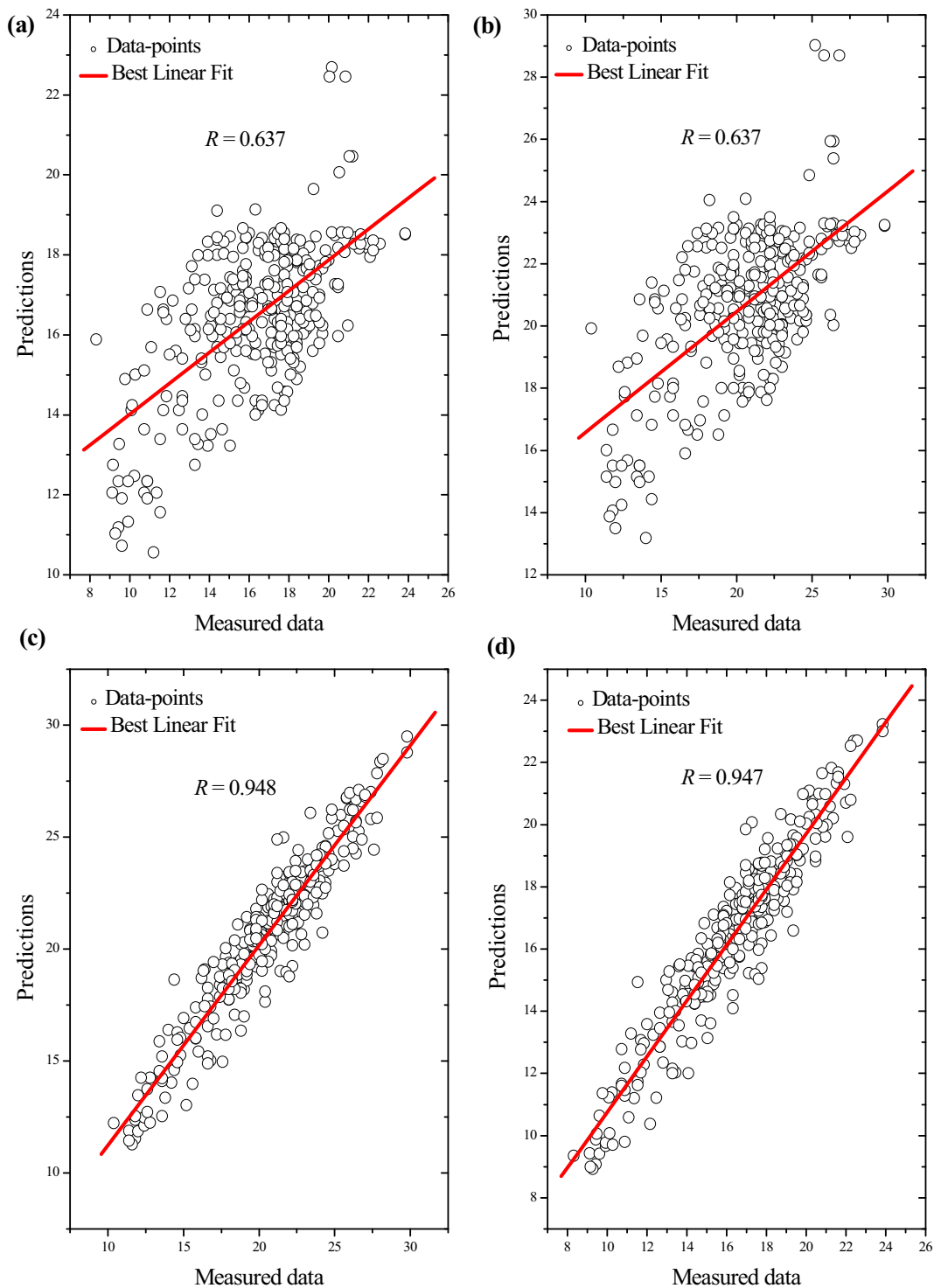


Fig. 12 Linear fit between the measured and predicted values: (a) GLM 1; (b) GLM 2; (c) ANFIS model 1; (d) ANFIS model 2

regressors identified indicate there are time lags in the treatment process. This is understandable as wastewater

undergo both chemical and biological changes before new process conditions are established. Therefore, the

ANFIS models can also serve as a reliable tool to estimate the time required to achieve a set treatment performance and also preempt process disturbances. The described modeling approach can be extended to other WWTPs as it would help the decision makers in effectively managing the plant operations in order to ensure the effluent quality consistently complies with the discharge regulations.

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