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# Application of a hybrid mechanistic/machine learning model for prediction of nitrous oxide $(N_2O)$ production in a nitrifying sequencing batch reactor



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### ABSTRACT

Nitrous oxide  $(N_2O)$  is a key parameter for evaluating the greenhouse gas emissions from wastewater treatment plants. In this study, a new method for predicting liquid  $N_2O$  production during nitrification was developed based on a mechanistic model and machine learning (ML) algorithm. The mechanistic model was first used for simulation of two 15-day experimental trials in a nitrifying sequencing batch reactor. Then, model predictions (NH<sub>4</sub>-N, NO<sub>2</sub>-N, NO<sub>3</sub>-N, MLSS, MLVSS) along with the recorded online measurements (DO, pH, temperature) were used as input data for the ML models. The data from the experiments at 20 °C and 12 °C, respectively, were used for training and testing of three ML algorithms, including artificial neural network (ANN), gradient boosting machine (GBM), and support vector machine (SVM). The best predictive model was the ANN algorithm and that model was further subjected to the 95% confidence interval analysis for calculation of the true data probability and estimating an error range of the data population. Moreover, Feature Selection (FS) techniques, such as Pearson correlation and Random Forest, were used to identify the most relevant parameters influencing liquid  $N_2O$  predictions. The results of FS analysis showed that NH<sub>4</sub>-N, followed by NO<sub>2</sub>-N had the highest correlation with the liquid  $N_2O$  production. With the proposed approach, a prompt method was obtained for enhancing prediction of the liquid  $N_2O$  concentrations for short-term studies with the limited availability of measured data.

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### 1. Introduction

Nitrous oxide ( $N_2O$ ) is one of the most significant greenhouse gases (GHGs) with an extremely high global warming potential (GWP), which is almost 300 times higher than that of carbon dioxide (IPCC, 2014). Wastewater treatment plants (WWTPs) are responsible for 3–5% of worldwide anthropogenic  $N_2O$  emissions (Mannina et al., 2019). In WWTPs,  $N_2O$  is primarily produced during biological nitrogen removal processes, including autotrophic nitritation (aerobic ammonium oxidation to nitrite) and heterotrophic denitritation (reduction of nitrite) (Su et al., 2019).

Moreover, carbon footprint (CF) is a measure of GHG emissions (Delre et al., 2019). The amount of  $N_2O$  produced in wastewater treatment operations has a significant impact on the overall CF of

\* Corresponding author, E-mail address: mohammad-javad.mehrani@pg.edu.pl (M.-J. Mehrani). WWTPs (Maktabifard et al., 2020). High shares of N<sub>2</sub>O emissions in the CF have been observed in biological nutrient removal (BNR) plants (Koutsou et al., 2018). Hence, an accurate estimation of N<sub>2</sub>O can help in better understanding of the process behavior and consequently, mitigation and control of this GHG in WWTPs. An N<sub>2</sub>O emission factor is a vital indicator of the long-term sustainability of WWTPs and environmental protection (Chen et al., 2020a; Vasilaki et al., 2019).

Mathematical models are a strong tool for process simulation, prediction, and optimization (Wisniewski et al., 2018). There are two possible approaches for  $N_2O$  modeling, including mechanistic models and machine learning (ML) techniques. In the area of wastewater treatment, the Activated Sludge Models (ASMs) (Henze et al., 2006) are the most common mechanistic models, which mathematically describe a hypothetical base for biological wastewater treatment processes. There has been a growing number of successful applications of mechanistic modeling for  $N_2O$  prediction

### 1. Data collection

Two operational periods (15 d each) of a nitrifying SBR at different temperatures (12 and 20 °C)

Sensors: N2O, pH, DO, and temperature Analytical data: Nitrogen components, and biomass (MLSS, MLVSS)

### 2. Mechanistic model

- Two-step nitrification and comammox model
- Expanding analytical data frequency from two days to 30 seconds (similar to sensor data)



# 5. N2O prediction with ML models

- ANN, SVM, and GBM algorithms
- Input data: sensors data and analytical data obtained from mechanistic model

Training = data-set at 20 °C Test = data-set at 12 ℃

# 3. Data pre-processing

- Min-max normalization (to normalize the dataset and transfer to 0-1)
- 4. Feature selection
- Pearson correlation (filter method)
- Random Forest (Wrapper method)



# 6. Model efficiency evaluation and comparison

Determination coefficient (R2), root mean square error (RMSE), mean absolute error (MAE), and Janus coefficient (J2)

# 7. 95% confidence interval for the selected ML model

Comparison of the population data scatter with the predicted model



in real WWTPs (Massara et al., 2018; Su et al., 2019; Wang et al., 2016; Zaborowska et al., 2019). However, these models have a few limitations. First of all, prediction of N2O requires an extensive knowledge of biological nitrogen transformations for model identification and calibration (Zaborowska et al., 2019). No generic mechanistic model is available, while various expansions and modifications have been added to the existing models (Vasilaki et al., 2019; Chen et al., 2020b). In addition, the traditional mechanistic models are over parameterized, highly sensitive to the changes in operational condition, and demand extensive efforts for calibration and validation. A calibration procedure for N<sub>2</sub>O is especially challenging as N2O is only an intermediate in the nitrogen transformation chain and its contribution to the nitrification process is low (Hwangbo et al., 2021).

On the other hand, ML is a tool for data analysis that can learn from input data and make decisions accordingly without any process equations and pathways (Al-Jamimi et al., 2018). ML algorithms recognize a specific pattern (during a training process) based on defined data (input data) for the prediction and/or classification purposes, which results in a more accurate output (Bagherzadeh et al., 2021; Osarogiagbon et al., 2021). In WWTPs, the ML prediction models have primarily been used for modeling influent/effluent wastewater characteristics. Those models are mainly artificial neural network (ANN) (Ryan et al., 2004; Shaahmadi et al., 2017) and support vector machine (SVM) (Alejo et al., 2018; Shaahmadi et al., 2017; Vasilaki et al., 2020b), while gradient boosting machine (GBM) has been used less frequently (Bagherzadeh et al., 2021). In addition, there are very limited studies on hybrid models (mechanistic models combined with the ML techniques) for forecasting influent/effluent wastewater components (Haimi et al., 2013; Hvala and Kocijan, 2020), but there has been no such a hybrid model applied for N<sub>2</sub>O prediction yet.

In terms of N<sub>2</sub>O, a predictive ML model of N<sub>2</sub>O emission, based on experimental data from an anoxic/aerobic bioreactor with ANN, was proposed by Sun et al. (2017). Moreover, two algorithms, including random forest (RF) and SVM, were used by Vasilaki et al. (2020b) to determine N<sub>2</sub>O emission factors. The SVM models performed better than RF in the training of the model to predict the expected range of N<sub>2</sub>O emission in WWTPs. Very recently, data-driven-based models, including long short-term memory (LSTM) and deep neural network, have been used to predict liquid N<sub>2</sub>O concentrations by big data from a full-scale WWTP (Hwangbo et al., 2021).

In this study, a predictive hybrid model for liquid N<sub>2</sub>O production was developed based on the data from a laboratory-scale nitrifying sequencing batch reactor (SBR). The new model overcame limitations of the pure mechanistic models or ML algorithms. This approach includes two major steps: i) mechanistic model simulation for expanding the short-term experimental data into an extensive data set with a very small interval (similar to the recordings of a liquid N2O sensor, and ii) liquid N2O predictions using three powerful ML algorithms (ANN, SVM, and GBM) to achieve a highly accurate model by producing input data from the previous step. To the



Statistical summary report of the input and output data used for the ML algorithms.

Data	Function	Mechanistic mode	el predictions				Online measure	ements		
		NH <sub>4</sub> -N (mg N/L)	NO <sub>3</sub> -N (mg N/L)	NO <sub>2</sub> -N (mg N/L)	MLSS (mg/L)	MLVSS (mg/L)	DO (mg O <sub>2</sub> /L)	Temp (°C)	pH -	N <sub>2</sub> O (mg N/L)
Training data	Min	2.00	4.86	4.92	545.4	350.9	0.20	18.74	7.11	0.00
	Max	49.06	37.29	12.97	2151.0	1351.0	2.22	21.51	8.35	0.57
	Mean	31.81	29.45	8.44	1176.5	744.54	0.61	20.08	7.55	0.15
Testing data	Min	0.00	19.22	0.00	545.4	350.9	0.150	11.74	7.09	0.00
_	Max	119.30	127.40	53.33	2151	1351.0	2.17	13.11	8.65	0.11
	Mean	38.89	93.11	22.57	1176	744.54	0.65	12.29	7.45	0.07

best of our knowledge, this hybrid approach has been used for the first time for N<sub>2</sub>O prediction in short-term studies. This research demonstrates a prompt method for enhancing prediction of liquid N<sub>2</sub>O concentrations with the limited availability of measured data.

### 2. Materials and methods

The modeling procedure includes two separate steps for the mechanistic model and ML algorithms. The first step was carried out for expanding the data-set, i.e., converting the communication interval of analytical data similar to the sensor data. The complete process of calibration and validation of the mechanistic model using GPS-X 8.0 software can be found in our previous study (Mehrani et al., 2021). In the second step, liquid N<sub>2</sub>O concentrations were predicted by three ML algorithms based on the data generated by the mechanistic model. The diagram in Fig. 1 presents the full modeling/ prediction approach.

### 2.1. Data collection for simulation

Two series of long-term washout experimental trials were carried out in the SBR. The experiments aimed at washing out NOB at decreasing solids retention times (SRTs) from 4d to 3d. The inoculum biomass was taken from the "Czajka" WWTP in Warsaw, Poland during winter and summer periods. The working volume of the SBR was 10 L and the reactor was operated for 15 days at the temperatures typical for winter and summer conditions, i.e., 12 °C and 20 °C. The temperature was kept constant during the experiment with a tolerance of  $\pm$  1.5 for 20 °C and  $\pm$  1.0 for 12 °C.

The experiments at 20 °C and 12 °C were selected for training and testing the ML algorithms, respectively (Table 1). The initial mixed liquid suspended solids (MLSS) and volatile fraction (MLVSS) concentrations were approximately 2000 mg/L and 1200 mg/L for the experiment at 12 °C, and 2500 mg/L and 1500 mg/L for the experiment at 20 °C.

The SBR was fed with ammonium-rich synthetic wastewater, including tracer elements, but without organic substrate (Mehrani et al., 2021). The volumetric nitrogen loading rates (NLRs) were kept stable at 0.02  $\pm$  0.01 and 0.05  $\pm$  0.01 g N/(L·d) at 12 °C and 20 °C, respectively. During the experiments, pH, temperature, DO concentration, and liquid N2O concentration were recorded every 30 s by online sensors (Table 1).

### 2.2. Simulations with a mechanistic model

For both experiments, simulations with the mechanistic model (two-step nitrification with comammox) were run with a communication time of 30 s (similar to the online sensor data). GPS-X 8.0 software (Hydromantis, 2021) was used as a simulation platform. The details of calibration and validation of the mechanistic model can be found in the previous study (Mehrani et al., 2021).

Simulation results, including nitrogen species (NH<sub>4</sub>-N, NO<sub>2</sub>-N, and NO<sub>3</sub>-N) and biomass components (MLSS, MLVSS), were selected as input data (> 50k data for each parameter) of the ML algorithms.

Table 1 shows a brief representation of the acquiesced input data set from the mechanistic model and online sensors data, separately for the training and testing data sets.

### 2.3. Data pre-processing for ML algorithms

Before ML prediction, the data obtained from the mechanistic model and online sensors were subjected to data engineering, i.e., cleaning the information with care taken to the missing or irregular records (Halim et al., 2021; Ranjan et al., 2021). Moreover, input data for training and testing the models were normalized and scaled between 0.0 and 1.0 values as:

$$Xn = \frac{Xi - Xmin}{Xmax - Xmin} \tag{1}$$

where Xn is normalized data, Xmax, and Xmin are the maximum and minimum values of the considered variable, and Xi is the value of the variable in each record. The normalization helps assign relevant weights for the ML models considering the value of each feature.

### 2.4. Feature selection for ML algorithms

Pearson correlation. Pearson correlation coefficient (PCC) is a feature selection (FS) filter method and is considered one of the most straightforward FS strategies. The PCC defines the linear relationship between two variables that range from + 1 to - 1, with 1 indicating total positive correlation, 0 indicating no correlation, while -1 showing the negative correlation (Ali et al., 2021; Alver and Altas, 2017). The PCC is computed as:

$$\sigma_{ij} = \frac{Cov(f_i, f_j)}{\sqrt{Var(f_i)Var(f_j)}}$$
(2)

where  $\sigma_{ii}$  is a correlation coefficient between a given feature  $f_i$  and all other features of the data set  $f_i$ , Cov is covariance, and Var is a variance.

Random Forest. The random forest (RF) is a ML filter method for ranking input variables according to their significance (Breiman, 2001). In this technique, several decision trees are created using random feature extraction and data set observations. The trees are de-correlated as a result of this random collection of records and features (bootstrapping). Each bootstrap is used to train a tree when there is a T number of trees in total (Breiman, 2001: Masmoudi et al., 2020). A small portion of data in each bootstrap is kept out of the box  $(oob_i)$  to evaluate the feature importance. Moreover, feeding the input feature f observations randomly to the tree will result in  $oob_i^f$ . Ultimately, the tree is able to predict the new values of the box by applying the mean squared error,  $MSE(oob_i^f)$ , and the feature importance is calculated as:

$$I(f) = \frac{1}{T} \sum_{i=1}^{T} \frac{MSE(oob_i^f) - MSE(oob_i)}{MSE(oob_i)}$$
(3)

Higher importance values show that the feature is more relevant to the target, and it can improve the prediction output.



### 2.5. Prediction of liquid N<sub>2</sub>O concentrations with ML models

After mechanistic modeling and data pre-processing, the ML algorithms were constructed for liquid  $N_2O$  prediction during the two experimental trials in the SBR. Python 3.8 open source programming language was used, while applying various libraries, such as Pandas, Matplotlib, Keras, and Scikitlearn. Each prediction algorithm is outlined in the following subsections and the details are given in the SI.

ANN algorithm. To locate the useful connection among dependent and independent variables, fully connected neural networks can be built up for prediction analysis (Yegnanarayana, 2009). For each neuron, a linear equation between its input and output is assumed. Due to the unpredictability of a non-linear model, more neurons are expected to anticipate the objective variable with a satisfactory precision (Eq. (4)):

$$y = \sigma(\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_n X_n)$$
(4)

where y is the output of the ANN,  $\sigma$  is the sigmoid function,  $X_i$  is the input number in each neuron,  $\beta_0$  is the sum of biases of each neuron, and  $\beta_i$  is the weight (trainable parameter) of the neuron.

An ANN is a multilayer perceptron (MLP) with three layers: input, hidden, and output (Kazemi et al., 2021). In a straightforward approach, the number of input layer neurons is the same as the size of a model dimensionality. To guarantee a smooth and precise link between the layers, the rectified linear unit (ReLU) technique was utilized for selecting a suitable number of the layers. The constructed ANN algorithm consists of seven input layers (MLSS was ignored due to a high correlation with MLVSS), three hidden layers with 10, 10, and 5 neurons respectively, and one neuron in the output layer. The optimization algorithm and the loss criterion were Adam and MSE, respectively.

SVM algorithm. The SVM is a versatile ML model that can do linear and nonlinear predictions, and even outlier detection (Vapnik et al., 1995; Géron, 2019). This method was originally designed to solve classification problems before being expanded to solve prediction problems. Since the cost function criteria for model building do not refer to attribute values that lie outside the margin, a model generated by the SVM is extremely depended on the subset of training data (Arshad et al., 2021). As shown in Fig S1, data points outside of the decision boundary will be ignored for developing the hyperplane (removing outliers). Similarly, the SVM model is only based on a subset of the training results (Vapnik et al., 1995). Any training data which are close to the model prediction (hyperplane) are ignored by the cost function method to prevent overfitting issues (Steinwart and Christmann, 2008).

SVM prediction (SVR) is a supervised learning model that uses the same SVM (classification) manner with minor editions. As it is difficult to predict a real number (infinite possibilities), a margin of error is considered for the prediction. The SVR transforms an input matrix to a higher dimensional feature space via a kernel. The following equation expresses the non-linear SVR function F(x) in a mathematical format (Eq. 5) (Awad and Khanna, 2015; Park et al., 2021):

$$F(x) = \sum_{i=1}^{M} (\alpha_i - \alpha_i^*) \quad K(X_i, X) + \delta$$
(5)

where M is the number of training records,  $\alpha_i$ ,  $\alpha_i^*$  are Lagrange multipliers, K is the transformation kernel that contains the dot product of mapped vectors of the support vectors  $(X_1 .... X_i)$ , and  $\delta$  is the sum of biases (Smola and Schölkopf, 2004).

*GBM algorithm.* The gradient boosting machine (GBM) is a kind of decision-tree ML model with a distinct ensemble formation of supportive technique (Ayyadevara, 2018). In this method, new trees are added to the ensemble sequentially based on the overall ensemble prediction error (Natekin and Knoll, 2013). The estimation

error for the dependent variable shrinks continuously by adding new trees until it reaches the highest possible accuracy (Bagherzadeh et al., 2021). The algorithm produces a new decision tree to minimize the prediction error, and finally, the output of all trees will be aggregated:

Data set = 
$$\{x^i, Y^i\}$$
  $\sum_{i=1}^{N}$  Minimizing  $\left\|\sum_{i=1}^{N} L(F(x^i), Y^i)\right\|$  (6)

where N is the number of records,  $x^i$  are independent variables,  $Y^i$  is the target variable in the training data set, L is the error function, and  $F(x^i)$  is the model output (Xenochristou et al., 2020).

The ultimate goal of GBM is to develop one strong model from several weak and smaller learning models (decision tree models). Considering Eq. (6), the GBM algorithm takes the training data and tries to minimize the error value. Decision trees divide the data set at each branch (node) to maximize the entropy. Each tree has several nodes and will split the data set until fulfilling the given hyperparameters (maximum tree depth).

Adjusting the hyper-parameters is a crucial step in designing a GBM model. Therefore, after many trial-and-error attempts, the following values were used in this study: a learning rate of 0.05, the number of 2000 trees for the forest, subsampling of 0.8, a tree depth of 6, a min sample leaf of 50, and minimum split samples as a 600.

### 2.6. Evaluation and comparison of the efficiency of ML models

The model performance can be evaluated with statistical methods. The dependent variable data are assumed as  $y_1$ ,  $y_2...y_n$  with the mean value  $\overline{y}$  and estimated values of this variable as  $f_1$ ,  $f_2...f_n$  (collectively known as  $y_i$  and  $f_i$ ). Then the sum square of residuals ( $SS_{res}$ ) and the total sum of squares ( $SS_{tot}$ ) are calculated, and the coefficient of determination (R-squared) indicates "goodness-of-fit" between the predicted feature and real values. Moreover, the mean absolute error (MAE), root mean square error (RMSE), and Janus coefficient describe the model errors and accuracy (Eqs. (7)–(12)) (Hauduc et al., 2015; Verma et al., 2013):

$$SS_{tot} = \sum_{i} (y_i - \bar{y})^2 \tag{7}$$

$$SS_{res} = \sum_{i} (y_i - f_i)^2 \tag{8}$$

$$R^2 = 1 - \frac{SS_{res}}{SS_{tot}} \tag{9}$$

$$MAE = \frac{1}{n} \sum_{i} (y_i - f_i)$$
(10)

$$RMSE = \sqrt{\frac{SS_{res}}{n}} \tag{11}$$

$$J^{2} = \sqrt{\frac{RMSE_{validation}^{2}}{RMSE_{calibration}^{2}}} \tag{12}$$

where  $y_i$  is the observed data in the data set,  $f_i$  is the model prediction,  $\overline{y}$  is the mean value of the observed data, and n is the number of observation samples.

### 2.7. Calculation of the confidence interval for the selected ML model

The confidence interval (*CI*) is a valuable measure to indicate the estimated range of error. In this study, there is a large number of observations with a normal error distribution. Therefore, the 95% CI was calculated with a z-score (Eq. (13)), considering that the standard deviation of the population is known due to having a high number of records (Hogg, 2012):



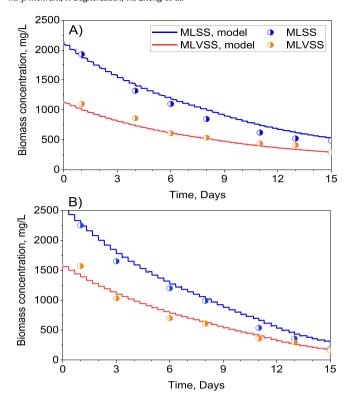


Fig. 2. Measured vs. predicted MLSS and MLVSS concentrations by the mechanistic model: A) Experiment at 12  $^{\circ}$ C, B) Experiment at 20  $^{\circ}$ C.

$$CI = \bar{X} \pm z \frac{\sigma}{\sqrt{n}} \tag{13}$$

where  $\bar{X}$  is the sample mean, z is the value from the standard normal distribution for the selected confidence level,  $\sigma$  is the standard deviation, and n is the total number of observations. If all the test points are within the 95% CI, it indicates a high level of precision (Abbas et al., 2018).

## 3. Results

### 3.1. Predictions of the mechanistic model (ML input data acquisition)

The results of predicted biomass concentrations (MLSS and MLVSS) and nitrogen species by the calibrated mechanistic model are shown in Fig. 2 and Fig. 3, respectively. The MLSS and MLVSS concentrations revealed a decreasing trend in both experiments due to the continuous biomass washout conditions (Fig. 2). Concerning the nitrogen species, in the experiment at 12 °C (Fig. 3a), NO<sub>3</sub>-N started to dilute after the first week resulting from biomass and NOB washout, while NO<sub>2</sub>-N stabilized at around 12-15 mg N/L to the end of the experiment. In the experiment at 20 °C (Fig. 3b), the NLRs were approximately doubled  $(0.05 \pm 0.01 \text{ g N/(L·d)})$  in response to the higher activity of bacteria at higher temperature and NO<sub>3</sub>-N dilution started after 10 days resulting from biomass and NOB washout, while NO<sub>2</sub>-N stabilized at around 40-45 mg N/L. The NO<sub>3</sub>-N and NO<sub>2</sub>-N production were more than double in Fig. 3b in comparison to Fig. 3a based on higher activity of bacteria in higher temperature.

### 3.2. Feature selection for ML algorithms

Fig. 4 shows a heatmap of the PCC between the variables (N species, biomass components, and online measurements) and  $N_2O$  concentration (target variable). The highest positive and negative correlation, 0.82 and -0.56, was obtained for NH<sub>4</sub>-N and NO<sub>2</sub>-N,

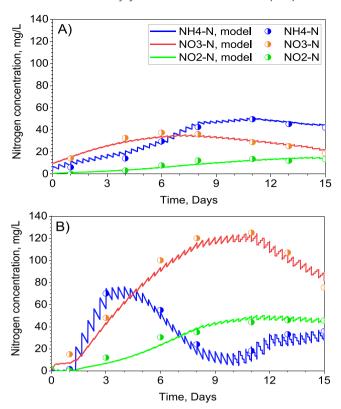


Fig. 3. Measured vs. predicted concentrations of nitrogen species by the mechanistic model: A) Experiment at 12  $^{\circ}$ C, B) Experiment at 20  $^{\circ}$ C.

respectively. The biomass components (MLSS and MLVSS) with the correlation factor of 0.48 were the next highest correlated variables. However, both parameters (MLSS and MLVSS) had a perfect correlation of 1.0 between each other, and thus only one of them (MLSS) was considered in the final subset of features to avoid multicollinearity issues.

The results of the RF method were in line with the Pearson correlation concerning the highest importance level of the nitrogen species with the target variable (Fig. 5). The maximum importance levels were obtained for  $NH_4$ -N (0.71) and  $NO_2$ -N (0.37). Moreover, among the online measurements, DO concentration had the highest correlation with  $N_2O$  concentration for both examined methods.

# 3.3. ML modeling results

### 3.3.1. Model predictions against training data

All three examined prediction models were trained and tested based on the data from the experiment at 20 °C and 12 °C, respectively. The comparative results for all the training models are shown in Fig. 6a. The overall performance of the models was within an acceptable range (Table 2) The noisy data points were not predicted accurately as they were treated as outliers.

Each algorithm required a specific approach and trial-and-error attempt to obtain the optimum parameter set, which ensured a high prediction accuracy. The ANN model was developed after looping over 300 epochs on the training data set. The SVM model was built with the regularization parameter (C=1), and SVR epsilon tube (epsilon=0.1). For the GBM, the following setting was selected: 4000 estimators, the learning rate of 0.01, min. sample leaf of 40, min. sample split of 30, and max. depth of 40.

### 3.3.2. Model predictions against test data

The trained prediction models were tested by another data set (experiment at 12 °C) (Fig. 6b). The SVM and GBM were overfitted,



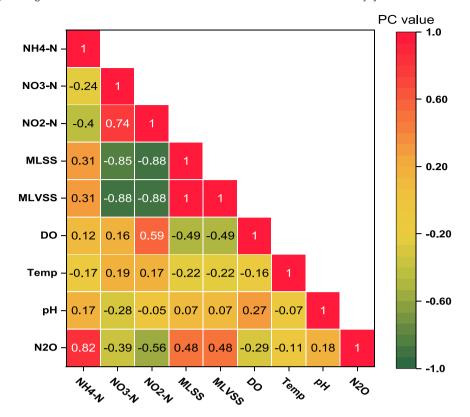
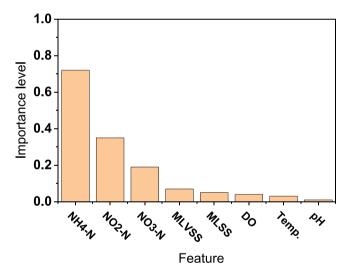


Fig. 4. Heatmap of Pearson correlation coefficient for input data with the target variable (N2O).

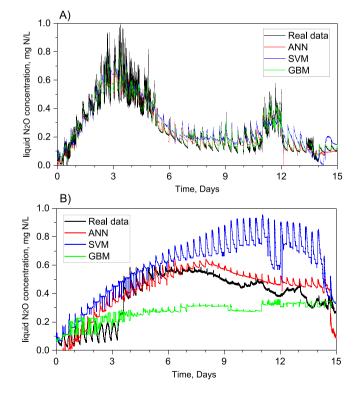


 $\textbf{Fig. 5.} \ \ \text{RF feature selection importance level of the features related to } \ N_2O \ production.$ 

even though they were able to capture the train data patterns. On the other hand, the ANN demonstrated its versatility under the different operational conditions and process patterns. As shown in Fig. 6b, the SVM overestimated the  $N_2O$  peak with a 6-day delay, and the GBM failed to predict any distinguishable climax.

# ${\it 3.3.3.}\ Model\ efficiency\ evaluation\ and\ comparison$

Table 2 presents the efficiency and error of each prediction algorithm. The ANN had the highest coefficient of determination, i.e.,  $R_{\rm Train}^2$ 0.93,  $R_{\rm Test}^2$ 0.67, and the lowest error indexes. These measures confirm that the ANN is the best model for predicting liquid  $N_2$ 0 concentrations during the experimental trials under different operational conditions. The SVM failed to predict the  $N_2$ 0 concentrations of unseen test data and the GBM partially detected the pattern



**Fig. 6.** Training and test results of the prediction models, (A) Train data (experiment at  $20\,^{\circ}$ C), and (B) Test data (experiment at  $12\,^{\circ}$ C).

of the real data at the beginning and end of the experiment. The high values of  $R^2_{\text{Train}}$  and low values of  $R^2_{\text{Test}}$  show that both SVM and GBM are overfitted and failed to predict the target variable. Among all the examined models, the ANN had also the best  $J^2$ , i.e., closest value to



**Table 2**Model efficiency criteria for the examined prediction models.

Model	Traini	ng data			Test c	lata			
	R <sup>2</sup>	MSE	RMSE	MAE	$\mathbb{R}^2$	MSE	RMSE	MAE	$J^2$
ANN	0.92	0.002	0.048	0.006	0.67	0.012	0.09	0.002	1.87
SVM	0.88	0.003	0.062	0.12	0.06	0.057	0.239	0.23	3.71
GBM	0.97	0.007	0.077	1.23	0.12	0.027	0.266	0.11	3.45

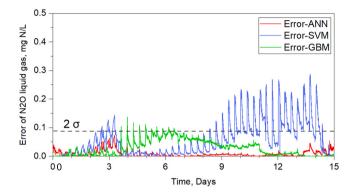


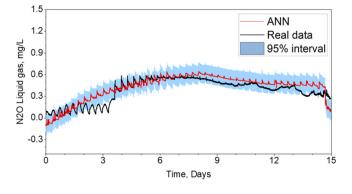
Fig. 7. Prediction errors of liquid  $N_2O$  concentrations for the examined prediction models from the test data ( $\sigma$  is the standard deviation of the test data set).

1.0. This confirms the ability of that model in finding data patterns under a highly dynamic behavior of the online measurements.

In Fig. 7, it can be seen that at the beginning of the experimental trial (days 2–4), the model outputs had considerable errors due to a sudden increase in  $N_2O$  production. After 4 days, when the stability in  $N_2O$  production was achieved, the errors (except for the SVM) became more stable. The ANN model output error was below the doubled standard deviation (2  $\sigma$ ) during the entire experiment, which indicated a valid and proper prediction (Fig. 7). Moreover, the variance of the prediction error in the ANN model was significantly lower than the SVM and GBM for the test data.

## 3.3.4. Confidence interval for the selected model (ANN)

The result of CI was evaluated based on the z-score as there was a large number of population and the Gaussian shape of the error distribution. Overall, the data values are mostly within the error range of  $\pm$  0.13  $N_2O$  mg/L as an upper and lower band of 95% CI (Fig. 8). Moreover, the variance of ANN model prediction is located in the confidence region (95% CI), showing its accuracy. This confirms that the ANN model predicted the measured data with 95% of the observed uncertainty.



 $\textbf{Fig. 8.} \ \ \text{Measured data, ANN model predictions and } 95\% \ \ \text{confidence interval of the predictions.}$ 

3.3.5. Effect of mechanistic modeling (data acquisition) on the prediction performance

For evaluation of the effect of the mechanistic model on the ML prediction performance, the models were fed with the direct experimental data (without mechanistic modeling). It can be seen in the SI (Fig. S4) that only the SVM could train the model with the acceptable accuracy ( $R^2$  =0.76), but failed to predict the test data set ( $R^2$  =0.0). The ANN and GBM could not train the models nor predict the test data-set. Table S1 shows the efficiency and error for each prediction ML algorithm without considering the mechanistic model. Mechanistic modeling prior to the ML procedure had a significant impact on the accuracy of training and testing the  $N_2$ O prediction. In the ANN model, after applying mechanistic modeling before the ML prediction, the accuracy increased dramatically from  $R_{Test}^2$  = -0.06 (Table S1) to  $R_{Test}^2$  = 0.67 (Table 2). This improvement can be justified by the fact that ML models need a sufficiently large data-set first to train and then predict accurately the test data-set.

### 4. Discussion

Results of the feature selection analysis on  $N_2O$  production show that the behavior of  $NH_4$ -N and  $NO_2$ -N plays an important role in predicting  $N_2O$  accumulation during nitrification. This finding was supported by the results of other studies (Duan et al., 2020; Li et al., 2015; Song et al., 2020). Song et al. (2020) observed that  $NH_4$ -N and the sum of  $NO_2$ -N and  $NO_3$ -N had the highest effect on  $N_2O$  emissions in a feature selection study of a full-scale WWTP. Strong positive correlations between  $NH_4$ -N oxidation and  $N_2O$  production during nitrification were also reported in a pilot-scale SBR (Li et al., 2015) and full-scale SBR (Duan et al., 2020). Moreover, in the study of (Duan et al., 2020), a strong positive correlation was also found between liquid  $N_2O$  and  $NO_2$ -N concentration (Pearson correlation of 0.93).

Furthermore, Duan et al. (2020) observed that  $N_2O$  emission exhibited a clear pattern that followed the DO profile in an intermittent aeration mode (Pearson correlation of 0.74). In the present study, a weak negative correlation (-0.29) was also found between the DO concentration and  $N_2O$  production during continuous aeration at the low DO setpoint of 0.6 mg  $O_2/L$ . It should be emphasized, however, that the well-established favorable conditions for liquid  $N_2O$  production comprise low DO and high  $NO_2-N$  concentrations (Mannina et al., 2017; Massara et al., 2018; Peng et al., 2014; Vasilaki et al., 2020a).

In the present study, the observed correlation of  $N_2O$  production with pH and temperature was lower than other evaluated parameters.  $N_2O$  production presented a weak positive correlation of 0.18 with pH which was kept in the range of 7.0–7.5 (Fig. 4). Law et al. (2011) reported that  $N_2O$  production fluctuated with pH in the range of 6.0 and 8.5 while keeping the pH between 6.4 and 7.0 reduced  $N_2O$  production in a partial nitrification system with aerobic conditions (Law et al., 2011). It can be seen in Fig. 5 that the online measurement data (temperature, DO, and pH) had a lower importance level than nitrogen species and biomass concentrations for the prediction of  $N_2O$  production. Vasilaki et al. (2020a) found that under similar DO and pH concentrations, the average liquid  $N_2O$  conditions can vary substantially.

The trend of  $N_2O$  production was different in experiments for train and test of ML Algorithms. However, the ANN model successfully predict the  $N_2O$  production in a test experiment (Fig. 6a,b). This higher production of  $N_2O$  (up to 1.0 mg N/L) in the first week of train experiment (Fig. 6a) can due to higher NLR of  $0.05 \pm 0.01 \text{ g N/(L.d)}$  in compare to the test experiment ( $0.02 \pm 0.01 \text{ g N/(L.d)}$ ) (Fig. 6b).

Table 3 presents various  $N_2O$  prediction studies with different modeling approaches (mechanistic and ML) in lab-scale, pilot-scale, or full-scale WWTPs. The mechanistic models were mostly used for the prediction of lab-scale or pilot-scale systems, while the ML

Table 3 Summary of different studies on  $N_2 O$  prediction in lab-scale or full-scale WWTP.

Sammary of american	definition of differences on 1/2 production in the Scarc of the Scarc	care training			
Prediction method	Model / Algorithm	Model Accuracy	System	Remarks	References
Mechanistic	A mathematical model based on AOB and HB processes	na	Lab- and pilot-scale	The model result showed that the contribution of heterotrophs to N <sub>2</sub> O production decreased with the increasing DO concentrations.	(Wang et al., 2016)
	ASM2d-N <sub>2</sub> O	na	Full-scale	N <sub>2</sub> O hotspots (important rise and falls) were predicted theoretically by an extended ASM considering all the biological pathways for N <sub>2</sub> O production.	(Massara et al., 2018)
	ASM2d	$R^2 = 0.65$	Lab-scale and pilot- scale	The model was used as a tool for assessing the multivariable problem of $N_2O$ mitigation in a combined N-P activated sludge system.	(Zaborowska et al., 2019)
	Mathematical model (two-step nitrification, four step HD)	$R^2 = 0.83 - 0.99$	Batch experiment	In the best-fit model, which combined two-step nitrification and denitrification pathways, the contribution of heterotrophs to $N_2O$ production was stronger than authrophs	(Domingo-Félez et al., 2017)
	ASM-ICE	$R^2 = 0.94 - 0.98$	Batch experiment	recorded predicted that N <sub>2</sub> O accumulation resulted from a faster drop of the N <sub>2</sub> O reduction rate than the NO <sub>2</sub> -N reduction rate.	(Ding et al., 2017)
	Extended ASM model	$R_{\rm Test}^2=0.88$	Full-scale	The integrated N <sub>2</sub> O model, considering the nitrification/denitrification pathways, could predict N <sub>2</sub> O production.	(Ni et al., 2015)
	Extended ASM model	$R^2 = 0.10 - 0.53$ RMSE= 0.4-1.14	Full-scale	Imbalanced distribution of flow rate between the two treatment lines did not result in a substantial increase in N <sub>2</sub> O emissions for the particular WWTP and operational parameters, according to the N <sub>2</sub> O simulation results.	(Solís et al., 2022)
	Extended ASM model	R <sup>2</sup> (NSE)= 0.3-0.33 RMSE= 0.38-0.41	Full-scale	The model estimated the N <sub>2</sub> O in terms of liquid and gaseous N <sub>2</sub> O emissions. This extended model indicated the essential function of heterotrophs. The N <sub>2</sub> O emission factor (EF) for the examined plant was between 0.9% and 0.94% of the influent TN-load.	(Maktabifard et al., 2022)
Machine Learning	Back Propagation-ANN	$R_{Test}^2 = 0.81-94$ MSE= 0.1-0.8	Full-scale / pilot-scale	The model was a convenient and feasible method for the prediction of N <sub>2</sub> O emissions in a nitrifying-denitrifying system.	(Sun et al., 2017)
	DNN and LSTM	$R_{\rm DNN}^2 = 0.86$ $R_{\rm CTM}^2 = 0.94$	Full-scale	In a comparative study, the LSTM-based forecasting model performed better than the DNN-based model in terms of model goodness-of-fit	(Hwangbo et al., 2021)
	SVM	па	Full-scale	The model predicted that $N_2O$ emissions accounted for 7.6% of the total $NH_4$ -N load (on average) in a sidestream SBR. $N_2O$ was correlated with DO, influent $NH_4$ -N load, and pH.	(Vasilaki et al., 2020a)
Hybrid method	Extended ASM1 and three different ML algorithms (ANN, SVM, and GBM)	ANN model:  R <sup>7</sup> rain=0.92  R <sup>7</sup> rest=0.67  RMSE <sub>Train</sub> = 0.048  RMSE <sub>Train</sub> =	Lab-scale	nd NO <sub>2</sub> -N had the highest correlation with N <sub>2</sub> O production. ANN revealed the formance among other algorithms.	This study

na: not available, AOB: Ammonia oxidation bacteria, HB: heterotrophic bacteria, HD: heterotrophic denitrification, DNN: deep neural network, SVM: support vector machines, ANN: artificial neural network, CSTR: continuously stirred tank reactor, ASM-ICE: Activated Sludge Model for Indirect Coupling of Electrons, NSE: Nash-Sutcliff coefficient.

methods considered data from full-scale systems. In general, the accuracy of  $N_2O$  prediction was higher for the pure ML models than the pure mechanistic models, and the developed hybrid model of this study revealed one of the highest prediction accuracies in comparison with the studies shown in Table 3.

An accurate prediction of  $N_2O$  can play a significant role in the mitigation of  $N_2O$  from WWTPs (Solís et al., 2022; Maktabifard et al., 2022). Hence, the proposed approach, i.e., expanding a data set by a mechanistic model and prediction with ML algorithms, can be useful for the limited amounts of data collected during  $N_2O$  measurement campaigns towards a mitigation of this hazardous gas from the WWTPs. With sufficiently big data set, the ML algorithms can ensure predictions with a satisfactory level of performance (without expanding the data by a mechanistic model).

The present approach still has some limitations. Only the limited experimental data from a lab-scale system were considered, while the hybrid model still requires validation based on experimental data from full-scale WWTPs. Furthermore, nitrogen transformations were only evaluated with respect to nitrification, whereas denitrification may also be an important source of  $N_2O$  production in full-scale WWTPs. Expanding the input data of ML algorithms by external software, such as GPS-X, requires experiments to validate the mechanistic model. Furthermore, applicability of the hybrid models still requires further validation with more variety of data-sets.

For future studies, a comparison between mechanistic modeling and ML predictions for liquid  $N_2O$  production and gas  $N_2O$  emission in a bigger data set is suggested. Future ML algorithms can be developed for prediction of specific  $N_2O$  production pathways, although this function is now applicable only by mechanistic models. In addition, estimation of an  $N_2O$  emission factor,  $EF_{N2O}$ , can be another interesting and useful application for full-scale WWTPs. This factor plays a critical role in determining the WWTP carbon footprint (Maktabifard et al., 2020).

### 5. Conclusions

A hybrid model, combining mechanistic and ML (ANN) models, accurately predicted the liquid N<sub>2</sub>O concentrations during two 15day experimental trials in a nitrifying SBR. This approach is novel in comparison with the previous attempts for finding a predictive model of N<sub>2</sub>O production during nitrification. The hybrid model successfully predicted unknown test data with an acceptable coefficient of determination ( $R_{TEST}^2 = 0.67$ ), showing its versatility in terms of variable operating conditions and the ability to generalize process patterns more accurately than the other two examined models (SVM, GBM). On the other hand, the SVM overfitted in the estimation of the test data and GBM failed to predict an acceptable model. Moreover, accounting for the level of uncertainty for the ANN model, the predicted values with more than 95% accuracy are reliable enough for delivering valuable information regarding the phenomenon for further research and practical applications. A hybrid modeling concept that combines mechanistic models of WWTPs (e.g., ASMs) with ML can be further expanded to predict N2O production/emission in full-scale WWTPs for N<sub>2</sub>O mitigation strategies.

### **Declaration of Competing Interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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## Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at doi:10.1016/j.psep.2022.04.058.

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