

Design and implementation of a software system for surveillance of antibiotics concentrations in wastewater

Yousuf Al-Hakim¹, Kosmas Dragos¹^a, Kay Smarsly¹^b, Silvio Beier², Claudia Klümper³

¹*Institute of Digital and Autonomous Construction, Hamburg University of Technology, Germany*

²*Chair of Urban Bioengineering for Resource Recovery, Bauhaus University Weimar, Germany*

³*Laboratory of Environmental Analysis, Department HAM 2, Hamm-Lippstadt University of Applied Sciences, Germany*
{yousuf.al-hakim, kosmas.dragos, kay.smarsly}@tuhh.de, silvio.beier@uni-weimar.de, claudia.kluemper@hshl.de


Keywords: Antibiotics, Antibiotic Resistance, Monitoring, Software Design, Wastewater Surveillance.


Abstract: Antibiotics are important drugs for treating infectious diseases. The extensive use of antibiotics for human, veterinary, and agricultural purposes has led to the permanent release of antibiotics into the environment, particularly into municipal wastewater. In turn, the widespread release of antibiotics into the environment has led to the emergence of antibiotic-resistant bacteria and antibiotic-resistant genes (collectively referred to as “antibiotic resistance”), which reduce the effectivity of antibiotic treatment. To counteract antibiotic resistance, surveillance of the release of antibiotics into the environment is necessary. Municipal wastewater surveillance may provide insights into the release of antibiotics into the environment. Current municipal wastewater surveillance systems, dedicated to antibiotics concentrations, rely on the ad-hoc use of third-party software, which may compromise the efficiency and user-friendliness of municipal wastewater surveillance systems. Designing software systems dedicated to the surveillance of antibiotics concentrations in municipal wastewater, based on well-established software design concepts, has received scarce research attention. In this study, a software system is proposed, which serves as a technological basis for the surveillance of the concentration of antibiotics in municipal wastewater in an efficient and user-friendly manner. The software system implements well-established software design concepts and is capable of conducting on-demand data analysis, as well as providing various user interfaces. The software system is validated using both data derived from simulations and real-world wastewater data recorded from a wastewater treatment plant. The results showcase the efficiency and user-friendliness of the proposed software system for the surveillance of antibiotics concentrations in municipal wastewater.

1 INTRODUCTION

Antibiotics are used for treating infectious diseases in human and veterinary medicine as well as for agricultural purposes (Davies, 2010). Since the introduction of the antimicrobial agent sulfonamide in the 1930s, usage of antibiotics has increased (Adler, 2018), leading to permanent release of antibiotics into the environment (Rizzo, 2013). Through feces of animals that have received antibiotics treatment, as well as through wastewater treatment plants, antibiotics are released into fields, soils, and local waters. The release of antibiotics into the environment poses risks to human and environmental health (Paulus, 2019). One of the most significant health risks is the emergence of antibiotic-

resistant bacteria (ARB) and antibiotic-resistant genes (ARG), collectively referred to as “antibiotic resistance” (Nguyen, 2021). Antibiotic resistance (AR) limits the effectiveness of antibiotics for treating infectious diseases (CDC, 2021). National and international institutions have realized the risk of the emergence of AR and have introduced measures for reducing the impact of AR on the health of humans, animals, and the environment (Aminov, 2010; Manzetti, 2014). However, since neither the EU nor other international and national institutions have set standards for the maximum allowable concentration of antibiotics in municipal wastewater (WHO, 2020), surveillance of antibiotics concentrations in the environment is rarely conducted. Gaining insights into the release of

^a <https://orcid.org/0000-0002-4204-6547>

^b <https://orcid.org/0000-0001-7228-3503>

antibiotics into the environment could stand to benefit from existing approaches on the surveillance of municipal wastewater at wastewater treatment plants, the most important literature on which is reviewed in the following paragraph, focusing on the underlying software systems.

Seminal works discussing the software of surveillance systems in general include, for example, surveillance approaches for geospatial data (Mutuku, 2022) and approaches for livestock surveillance (Mena, 2019). Generally, the literature shows the case-specific manner in which software for surveillance systems is designed, which is underlined by software development without clear software design concepts. Regarding the surveillance of municipal wastewater, a surveillance system proposed by Selisteanu et al. (2020) has been designed using three third-party software environments. Martinez et al. (2020) have used third-party middleware for designing surveillance systems. Finally, studies that have focused on the surveillance of antibiotics concentrations in municipal wastewater have been conducted by Mtetwa et al. (2021) with wastewater samples from wastewater treatment plants in South Africa, by Majlander et al. (2021) with wastewater samples from two hospitals in Finland, and by Huijbers et al. (2020), with wastewater samples from municipal wastewater treatment plants of several European countries. In summary, the studies show that municipal wastewater surveillance systems typically lack underlying software that is designed specifically for the surveillance of municipal wastewater. To improve efficiency and user-friendliness, dedicated software systems based on well-established design concepts are necessary.

This study proposes a software system for efficient and user-friendly surveillance of antibiotics concentrations in municipal wastewater. The software system supports executing multiple tasks simultaneously, such as data storage and management, data analysis, and data visualization. In data storage and management, the software system stores wastewater data in a database, specifically designed for the surveillance of antibiotics concentrations. In data analysis, algorithms are implemented to analyze wastewater data and identify parameters that are critical to the surveillance of antibiotics concentrations in municipal wastewater. In data visualization, the parameters are visualized in multiple views. The software system is validated with data derived from simulations as well as from a real-world municipal wastewater treatment plant (WWTP). The software system increases the efficiency and user-friendliness of the surveillance of antibiotics concentrations in municipal wastewater by organizing and bundling tasks into one software system. With the software system, surveillance of

municipal wastewater may be used as a reliable tool to limit the release of antibiotics concentration. The remainder of this paper is structured as follows: The design of the software system is discussed in the next section, followed by the implementation. Then, the software system is validated and, finally, conclusions are drawn, and an outlook on future research is given.

2 DESIGN OF THE SOFTWARE SYSTEM

In this section, the design of the software system is discussed. Upon defining the requirements, the software design is presented, showing how the requirements are met. For designing the software system for the surveillance of antibiotics concentrations in municipal wastewater, functional and non-functional software requirements are defined.

Functional requirements are directed to project-related properties and ensure high quality and feasibility of the software system (Glinz, 2007). The main functional requirements for the software system are:

- The software system must allow accessibility through web browsers and smartphone applications;
- The visualization of the wastewater data must be adaptable to the requirements of every user;
- The software system must consider data analysis functionalities with the option to switch between algorithms;
- The concepts of the software system must be valid generally, meaning that the software system can be used in different environments.

The main non-functional software requirements for the software system are correctness, robustness, extensibility, and reusability. For further information on non-functional software requirements, interested readers are referred to the IEEE Standard Glossary of Software Engineering Terminology (IEEE, 1990).

The software system for the surveillance of antibiotics concentrations in municipal wastewater is presented in Figure 1. The rectangles represent main components of the software system. The squares resemble ports (P1-P3), to which interfaces are connected. The ports allow extensions to the software system, which enable users to access the software system through web browsers and smartphone applications. The full circles resemble interfaces provided by a component, and the half circles resemble interfaces required for a component. For the software design, the model-view-controller (MVC)

software design concept is pursued because MVC separates the core of the software system into three components (Smarsly et al., 2023): The model component, the view component, and the controller component. Separating the software system reduces the complexity, since different functionalities are separated accordingly. The aspect of separation is an advantage of MVC over other software design concepts, such as the microkernel software design concept or the event-driven software design concept.

In the <<Core>> component, which resembles the software architecture of the software system, the port P1 enables the view component to visualize wastewater data with dashboards, charts, or spreadsheets, according to the requirements of respective users. The port P2 enables switching between algorithms, with which wastewater data is processed and analyzed. The port P3 offers interfaces to connect databases to the model component of the software system, allowing analysis and visualization of wastewater data from multiple sources. To ensure general validity of the database of the software system, switching between different databases as a whole is enabled by P3, as well.

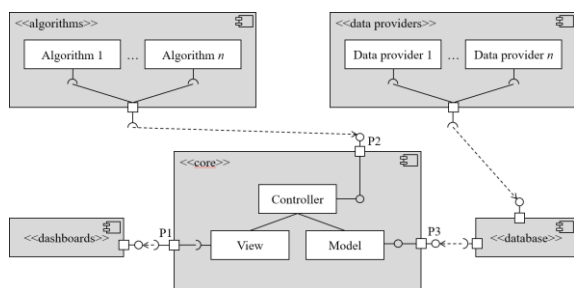


Figure 1: Software system for the surveillance of antibiotics concentrations in municipal wastewater.

3 IMPLEMENTATION OF THE SOFTWARE SYSTEM

This section describes the implementation of the software system. The components of the software system are listed, and the functionalities are explained in detail.

3.1 Model component

The database system of the software system resides in a cloud, on which wastewater data can be stored and shared with project partners. The cloud of the software system is based on the client-server software *Nextcloud*, with which clouds can individually be scaled and used on various systems

such as small microcontrollers or large data centers. Additionally, *Nextcloud* offers a web-based application and a smartphone application. *Nextcloud* enables the assignment of different rights to different users. By sharing individual access codes, or by including users to access lists, the users may access the wastewater data to the degree designated. The cloud is managed by a data manager, which is responsible for the following tasks:

- Assigning rights to users;
- Ensuring the wastewater data is always up to date, old versions are deleted or backed up, and only the newest version is passed to the model component;
- Unifying data file types, for example *.csv* or *.xls*, and assuring that the providers of wastewater data conform to the formats of the predefined data file types.
- Performing any non-automated task necessary for the operation of the database system, such as maintaining the database system structure, identifying and correcting errors, or performing updates of hardware and software.

3.2 Controller component

Once the wastewater data has been collected, the wastewater data is analyzed. The controller component performs data analysis tasks, including correcting the wastewater data and preparing the wastewater data for forecasting. The controller component is built using the *Python* programming language. The *Python* programming language is used, because the native libraries of the *Python* programming language include a variety of different algorithms (Bogdanchikov, 2013), which reduce the need to install external libraries when analyzing the wastewater data. Wastewater data is analyzed in the controller component in four steps, described in the following sub sections. A flowchart, showcasing the workflow of the data analysis, in the controller component is shown in Figure 2.

3.2.1 Data correction

Raw wastewater data is often not ready for data analysis immediately after the data is created. Several factors can lead to gaps in the wastewater data, such as:

- Disturbances during municipal wastewater sampling, or even loss of municipal wastewater samples;
- The human factor, such as illness of the worker on the day in which collection of municipal wastewater samples is scheduled;

- Errors in devices for extracting the data samples.

First, the controller component locates gaps in the wastewater data. For analyzing wastewater data, particularly when advanced data analysis methods are applied to the wastewater data, gaps are detrimental. The controller component applies linear interpolation to fill the gaps.

3.2.2 Calculation of moving statistics

Once the wastewater data is corrected, moving statistics are calculated, since moving statistics are used for modelling, forecasting, and gaining insights into the wastewater data. First, the controller component calculates the moving average (*MA*) over a range of antibiotics concentrations. Upon computing the moving average, the controller component calculates the moving standard deviation (*MSD*)

$$MSD = \sqrt{\frac{1}{n} \sum_{i=1}^n (y_i - MA)^2}, \quad (1)$$

where y_i is the wastewater data point. In addition to the *MA* and the *MSD*, the controller calculates the trend and the seasonality of the wastewater data.

The seasonality is determined by calculating the seasonal variation (*SV*) of the wastewater data

$$SV = y_i - MA. \quad (2)$$

Plotted over time, the seasonal variation shows the recurring, short-term cycle of the wastewater data.

3.2.3 Data normalization and stationarity

Calculating the statistics *MA* and *MSD*, as well as the trend and *SV*, provides insight into the wastewater data. With the *MA*, the wastewater data can be examined for stationarity. Stationarity is defined as follows (NIST, 2023):

A stationary process has the property that the mean, variance and autocorrelation structure do not change over time.

Stationarity is particularly important for modeling and for forecasting the wastewater data. If the wastewater data is non-stationary, the wastewater data is normalized to achieve a constant *MA*. The controller component checks the following normalizations for stationarity:

The logarithmic normalization:

$$f(x) = \log(x); \quad (3)$$

The square-root normalization:

$$f(x) = \sqrt{x}; \quad (4)$$

The cube-root normalization:

$$f(x) = \sqrt[3]{x}. \quad (5)$$

The Augmented Dickey-Fuller Test (ADF) (Paparoditis, 2013) is performed to select the appropriate normalization with respect to stationarity. The normalized dataset is used for modeling and forecasting of the wastewater data. To increase efficiency of the software system, the controller component automatically finds the appropriate normalization with respect to stationarity.

3.2.4 Forecasting

To estimate the future behavior of the wastewater data, the controller component performs forecasting. A model of the wastewater data is built in the controller component using linear regression, which is an efficient way to model time series data (Myers, 1990). The *Python* library *XGBoost* (XGB) is used for forecasting. XGB is a decision-tree machine learning library, used for regression, classification, and ranking problems (He, 2016). To train the regression model, the wastewater data is split into three parts. The first two parts are used by the controller component for training the regression model. The last part is used to test the regression model. The metric, with which the controller component trains the regression model is the root mean squared error (*RMSE*):

$$RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^n (y_i - \hat{y}_i)^2} \quad (6)$$

where y_i are the actual measurements and \hat{y}_i are the corresponding predictions of the wastewater data. The regression model runs multiple training iterations to reduce the *RMSE* as much as possible. For efficiency, training of the model stops automatically as soon as the *RMSE* reaches the lowest value, with a minimum of 10 training iterations, as an empirical value.

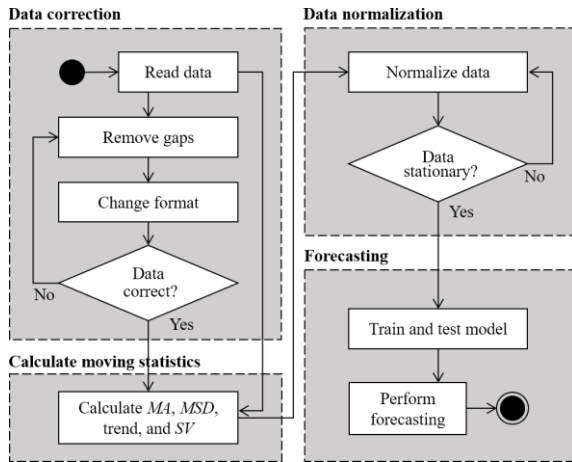


Figure 2: Flowchart of the data analysis conducted by the controller component.

3.3 View component

Once the data analysis has been completed, the controller component passes the wastewater data to the view component of the software system, where the wastewater data is visualized. To visualize the wastewater data, the software system uses the *Python* library *Matplotlib*. *Matplotlib* is a library for creating static, animated, and interactive visualizations with *Python*, without the need to install third-party software. The wastewater data is visualized using interactive figures that are updated automatically and that can be formatted individually. In addition to the visualization techniques provided by *Matplotlib*, the *Python* programming language provides interfaces that allow several third-party software packages to be used for visualizing the wastewater data, such as the *Python* extension *Streamlit*, which allows visualizing wastewater data with dashboards containing interactive figures and graphics.

3.4 Validation tests

This section discusses the tests, which have been devised to validate the software system. The validation tests consist of a test using data derived from simulations, as will be explained in the next paragraph, and a test using real-world municipal wastewater data.

For validating the software system with data derived from simulations, a wastewater data set is simulated using the Monte Carlo Simulation (MCS). The MCS is used to simulate different outcomes of a process that cannot be predicted due to the intervention of random values. The MCS is conducted in four steps, on which details may be found in Harrison (2009). The real-world wastewater

data has been measured at the Neugut WWTP in Dübendorf, Switzerland. The Neugut WWTP treats wastewater from 150,000 people, with an industrial contribution of approximately 50%. The flow of the Neugut WWTP averages 19,000 m³/day at dry weather conditions (Enz, 1992).

The database system is located on the *Nextcloud* server of Hamburg University of Technology. To validate the functionality of the database system, some users are assigned the right to download and reshare the folder, in which the wastewater data is stored, while other users are assigned the right to edit the wastewater data and create new files of the wastewater data. Users of the database have received a personalized invitation link, with which access to the database is granted. After reading the wastewater data, the model component passes the wastewater data to the controller component, where the wastewater data is corrected. No gaps are present in the data derived from simulations, which is why the controller component skips the data correction step in the first validation test.

Next, the controller component performs data analysis on the wastewater data. The *MA* and the *MSD* are calculated, as well as the trend *m* and the seasonal variation *SV*. To calculate the trend and seasonal variation, the controller component uses the *Python* library *Statsmodels*, which contains statistical functions for data analysis. The original wastewater data and the trend are depicted in Figure 3. For the visualization of the trend, visualization techniques provided by the internal *Python* library *Matplotlib* are used.

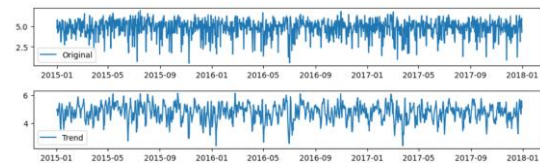


Figure 3: Line graphs for the original wastewater data, derived from simulations, and the trend.

Since the MCS is used, random values are present in the wastewater data. The random values of the data derived from simulations are apparent in the trend, as the trend line is highly periodic. To achieve stationarity, the controller component identifies the appropriate normalization by applying the ADF. The result of the ADF shows that the cube root normalization is appropriate with respect to stationarity.

Next, the controller component uses the data derived from simulations for training and testing the regression model. The wastewater data is split into three parts, the first two being used for training. The

controller component stops the training phase after 18 training iterations, since the *RMSE* reaches the lowest value. Finally, the controller component checks the forecasting capabilities of the regression model, using the third part of the data derived from simulations for testing the model.

The software system then performs the data analysis of the real-world wastewater data similarly to the data derived from simulations. The controller component normalizes the real-world wastewater data with the logarithmic normalization. Next, the controller component splits the normalized real-world wastewater data into three parts. When training the regression model with the real-world wastewater data, the controller component stops after 39 training iterations. The visualization and the analysis results of the real-world wastewater data are conducted using a dashboard, provided by the view component which uses the *Streamlit* extension of *Python*. The dashboard is shown in Figure 4.

The dashboard of the software system bundles multiple visualization techniques in one interactive screen. The wastewater data is visualized, alongside the forecast conducted by the regression model. The data analysis results, such as *MA* and *MSD* of antibiotics concentrations, are depicted in the form of line charts. The visualization techniques in the dashboard can be changed interactively, according to user preferences.

As a result, the validation tests with data derived from simulations show the efficient and user-friendly manner with which the database system is designed. The cloud allows assigning different rights to different users. Users are able to perform data management (depending on the rights assigned) from web-browsers or the *Nextcloud* smartphone application. A particularly efficient feature of the controller component is identifying the normalization for modelling. The visualization of the real-world wastewater data increases the user-friendliness of the software system. In conclusion, the software system significantly improves the efficiency and user-friendliness of the surveillance of antibiotics in municipal wastewater. The database allows user-friendly data management through the folder structure and rights assignment. The controller component analyses data efficiently, without the need of intervention between calculation steps. The view component enables user-friendly visualization of antibiotics concentrations in wastewater data and the analysis results through a variety of visualization techniques, including dashboards.

4 SUMMARY AND CONCLUSION

Antibiotic resistance poses serious health risks to humans, animals, and the environment. High concentrations of AR, particularly in municipal wastewater, threaten the potential effectivity of antibiotics in treating infectious diseases. Surveillance of municipal wastewater is, therefore, a vital step to reduce the release of antibiotics in the environment. This study has proposed a software system for efficient and user-friendly surveillance of antibiotics concentrations in municipal wastewater. The software system is based on the model-view-controller software design concept. The MVC concept separates the core of the software system into three components. In the model component, wastewater data is stored and managed, the controller component is responsible for data analysis, and the view component is designed for visualization. The software system has been validated using data derived from simulations as well as wastewater data obtained from a real-world wastewater treatment plant.

The validation tests have demonstrated the high efficiency and user-friendliness of the surveillance of municipal wastewater using the proposed software system. In particular, the MVC software design concept has exhibited its positive attributes within every component of the software system: (i) the model component has allowed user-friendly data management with an efficient folder structure, (ii) the controller component has conducted the data analysis in an efficient manner, including normalizations to achieve stationary wastewater data, and (iii) the view component has allowed a variety of data visualizations, while enabling multiple project partners at different locations to collaborate. The study aspires to provide a tool that helps adhere to standards and regulations regarding antibiotics concentrations in municipal wastewater. Future work may include coupling the software system with digital models of wastewater treatment plants, building upon previous studies reported in Söbke et al. (2018; 2021). The software system focusses on connecting to dashboards and similar visualization techniques. Future work may focus on integrating the monitoring data into 3D models, such as building information models. Also, future work may focus on optimizing the algorithms used for analyzing the wastewater data, enabling the software system to predict antibiotics concentrations in municipal wastewater with high accuracy.



Figure 4: Dashboard created by the view component of the software system.

ACKNOWLEDGEMENTS

This collaborative research is partially funded by Thüringer Aufbaubank (grant number 2021 FE 9143/44) through Bauhaus University Weimar, Germany (grant number 2360200276-2/21), by the Deutsche Bundesstiftung Umwelt (DBU) under grant number 38351/01-25, by the German Research Foundation (DFG) under grant SM 281/20-1, and by the German Federal Ministry for Digital and Transport (BMDV) within the mFUND program under grant 19FS2059C. The authors would like to sincerely thank the Swiss Federal Institute of Aquatic Science and Technology (EAWAG) for openly sharing data and accelerating scientific progress through transparency. Any opinions, findings, conclusions, or recommendations expressed in this paper are those of the authors and do not necessarily reflect the views of EAWAG or the sponsors.

REFERENCES

- Adler, N., Balzer, F., Blondzik, K., Brauer, F., Chorus, I., et al. (2018). *Antibiotics and Antibiotics Resistances in the environment – Background, challenges and options for action*. German Environment Agency, Dessau-Roßlau, Germany.
- Aminov, R.I. (2010). A brief history of the antibiotic era: Lessons learned and challenges for the future. *Frontiers in Microbiology*. Vol. 1, ID: 134.
- Bogdanchikov, A., Zhaparov, M., Suliyev, R. (2013). Python to learn programming. In: *Proc. of the International Conference on Science & Engineering in Mathematics, Chemistry and Physics*. Jakarta, Indonesia, 01/25/2013.
- Centers for Disease Control and Prevention, CDC. (2021). *Actions to fight antimicrobial resistance*. Available at: <https://www.cdc.gov/drugresistance/actions-to-fight.html> (Accessed on: 10/03/2023).
- Davies, J., Davies, D. (2010). Origins and evolution of antibiotic resistance. *Microbiology and Molecular Biology Reviews*. Vol. 74, No. 3, p. 417–433.
- Enz, D. (1992). Dübendorf, Kläranlage Neugut, Digital Equipment. Available at: https://www.e-pics.ethz.ch/index/ETHBIB.Bildarchiv/ETHBIB.Bildarchiv_26755.html (Accessed on: 10/03/2023).
- Glinz, M. (2007). On non-functional requirements. In: *Proc. of the 15th IEEE International Requirements Engineering Conference*. Delhi, India, 10/15/2007.
- Harrison, R.L. (2009). Introduction to Monte Carlo Simulation. In: *Proc. of the Nuclear Physics Methods and Accelerators in Biology and Medicine: Fifth Int. Summer School on Nuclear Physics Methods and Accelerators in Biology and Medicine*. Bratislava, Slovakia, 07/06/2009.
- He, T. (2016). *XGBoost: Scalable and Flexible Gradient Boosting*. Available at: <https://xgboost.ai/> (Accessed on: 10/01/2023)
- Huijbers, P., Larsson, J., Flach, C.-F. (2020). Surveillance of antibiotic resistant *Escherichia coli* in human populations through urban wastewater in ten European countries. *Environmental Pollution*. Vol. 261, ID: 114200.
- Majlander, J., Anttila, V.-J., Nurmi, W., Seppälä, A., Tiedje, J., Muziasari, W. (2021). Routine wastewater-based monitoring of antibiotic resistance in two Finnish hospitals: Focus on carbapenem resistance genes and genes associated with bacteria causing hospital-acquired infections. *Journal of Hospital Infection*. Vol. 117, p. 157–164.
- Manzetti, S., Ghisi, R. (2014). The environmental release and fate of antibiotics. *Marine Pollution Bulletin*. Vol. 79, No. 1–2, p. 7–15.
- Martínez, R., Vela, N., El Aatik, A., Murray, E., Roche, P., Navarro, J.M. (2020). On the use of an IoT integrated system for water quality monitoring and management in wastewater treatment plants. *Water*. Vol. 12, No. 4, ID: 1096.
- Mena, M., Corall, A., Iribarne, L., Criado, J. (2019). A progressive web application based on microservices combining geospatial data and the IoT. *IEEE Access*. Vol. 7, p. 104577-104590.
- Mtetwa, H.N., Amoah, I.D., Kumari, S., Bux, F., Reddy, P. (2021). Wastewater-based surveillance of antibiotic resistance genes associated with tuberculosis treatment regimen in KwaZulu Natal, South Africa. *Antibiotics*. Vol. 10, No. 11, ID: 1362.
- Mutuku, C., Gazdag, Z., Melegh, S. (2022). Occurrence of antibiotics and bacterial resistance genes in wastewater: Resistance mechanisms and antimicrobial resistance control approaches. *World Journal of Microbiology and Biotechnology*. Vol. 38, No. 9, ID: 152.

- Myers, R.H. (1990). *Classical and modern regression with applications*. Duxbury/Thompson Learning, Pacific Grove, CA, USA.
- National Institute of Standards and Technology (NIST) (2023). *Stationarity*. In: “NIST/SEMATECH e-Handbook of Statistical Methods”. Available at: <https://www.itl.nist.gov/div898/handbook/pmc/section4/pmc442.htm> (Accessed on 07/19/2023).
- Nguyen, A.Q., Vu, H. P., Nguyen, L.N., Wang, Q., Djordjevic, S.P., Donner, E., Yin, H., Nghiem, L.D. (2021). Monitoring antibiotic resistance genes in wastewater treatment: Current strategies and future challenges. *Science of The Total Environment*. Vol. 783, ID: 146964.
- Paparoditis, E., Politis, D.N. (2013). The asymptotic size and power of the augmented Dickey-Fuller test for a unit root. *Econometric Reviews*. Vol. 37, No. 9, p. 955-973.
- Paulus, G.K., Hornstra, L.M., Alygizakis, N., Slobodnik, J., Thomaidis, N., Medema, G. (2019). The impact of on-site hospital wastewater treatment on the downstream communal wastewater system in terms of antibiotics and antibiotic resistance genes. In: *International Journal of Hygiene and Environmental Health*. Vol. 222, No. 4, p. 635–644.
- Rizzo, L., Manaia, C., Merlin, C., Schwartz, T., Dagot, C., Ploy, M.C., Michael, I., Fatta-Kassinos, D. (2013). Urban wastewater treatment plants as hotspots for antibiotic resistant bacteria and genes spread into the environment: A review. *Science of The Total Environment*. Vol. 447, p. 345–360.
- Selișteanu, D., Petre, E., Prejbeanu, R., Popescu, I.M., Mehedințeanu, S. (2020). Software solutions for simulation, monitoring and data acquisition in wastewater treatment plants. In: *Proc. of the 21th International Carpathian Control Conference (ICCC)*. High Tatras, Slovakia, 10/27/2020.
- Smarsly, K., Al-Hakim, Y., Peralta, P., Beier, S., Klümper, C. (2023). A systematic review and recommendation of software architectures for SARS-CoV-2 monitoring. In: *Proc. of the 16th International Conference on Health Informatics*. Lisbon, Portugal, 02/18/2023.
- Söbke, H., Peralta, P., Smarsly, K., Armbruster, M. (2021). An IFC schema extension for BIM-based description of wastewater treatment plants. *Automation in Construction*. Vol. 129, ID: 103777.
- Söbke, H., Theiler, M., Tauscher, E., Smarsly, K. (2018). BIM-based description of wastewater treatment plants. In: *Proc. of the 16th International Conference on Computing in Civil and Building Engineering (ICCCBE)*. Tampere, Finland, 06/05/2018.
- The Institute of Electrical and Electronics Engineering (1990). *IEEE Standard Glossary of Software Engineering Terminology*. IEEE, New York, NY, USA.
- World Health Organization (2020). *Antibiotic resistance*. Available at: <https://www.who.int/news-room/fact-sheets/detail/antibiotic-resistance>. (Accessed on: 03/10/2023).