

# Journal of Population Therapeutics & Clinical Pharmacology

RESEARCH ARTICLE DOI: 10.53555/jptcp.v31i6.6572

# ASSESSING MICROPLATE-BASED BIOCHEMICAL ANALYSIS FOR BACTERIAL IDENTIFICATION: A VALIDATION STUDY

Muodiaju Joan C<sup>1</sup>, Muhammad Rashid<sup>2</sup>, Inaam Ullah<sup>3</sup>, Dr. Tariq Rafique<sup>4\*</sup>, Dr. Mahmoud El Safadi<sup>5</sup>, Amna Mahmood<sup>6</sup>

<sup>1</sup>Laboratory Research Analyst, Department of Surgery, Center for Human Systems Immunology, Duke University, USA

<sup>2</sup>Student, University Institute of Medical Lab Technology – Faculty of Allied Health Sciences, The University of Lahore, Pakistan

<sup>3</sup>Student, Institute of Medical Lab Technology – Faculty of Allied Health Sciences, The University of Lahore, Pakistan

<sup>4\*</sup>Assistant Professor Dadabhoy Institute of Higher Education, Karachi, Pakistan

<sup>5</sup>Department of Chemistry, College of Science, United Arab Emirates University, P.O.BOX 15551, Al Ain, Abu Dhabi, United Emirates

<sup>6</sup>Assistant Professor Biochemistry, Department of Biochemistry, Faculty of Basic Sciences, NUR International University Lahore, Pakistan

\*Corresponding Author: Dr. Tariq Rafique,

\*Assistant Professor Biochemistry, Department of Biochemistry, Faculty of Basic Sciences, NUR International University Lahore, Pakistan

# **ABSTRACT:**

**Objective**: Bacterial infections present a significant global health threat, necessitating effective identification for appropriate treatment. Biochemical analysis traditionally classifies bacteria based on metabolic traits. This study aimed to validate biochemical studies on microplates by testing lactose, fructose, glucose, sucrose, and maltose metabolism.

**Methods**: Biochemical analysis was conducted using microplates to examine bacterial metabolism. Tests for lactose, fructose, glucose, sucrose, and maltose were performed, assessing color changes, turbidity, and pH alterations. The expected color alteration in carbohydrate metabolism was compared between microplate results and experimental tube observations.

**Results**: Microplate testing revealed unexpected outcomes, failing to demonstrate anticipated color changes during bacterial sugar metabolism. In contrast, experimental tubes exhibited the anticipated yellow color indicative of bacterial fermentation in carbohydrate substrates.

**Conclusion**: Despite the reliability of biochemical analysis in bacterial classification, microplate testing for carbohydrate metabolism showed inconsistencies. The discrepancy in color alteration between microplates and experimental tubes highlights the need for further validation and refinement of microplate-based biochemical assays in bacterial identification.

**KEYWORDS**: Microplate, Biochemical Testing, And Bacteria.

# **INTRODUCTION:**

In most impoverished nations, bacterial illnesses are among the most serious issues. The high rate of cases, together with morbidity and death, is indicative of both the sociocultural and economic

status as well as the hygienic and sanitary circumstances. The patient, the microbes that cause the infections, and the hospital setting are some of the causes of these diseases (Fajdek-Bieda, Pawlińska, Wróblewska, & Łuś, 2024; Jaber et al., 2024).

Since infectious diseases produced by multiple microbes possess distinct courses and effects, identifying and isolating these microbes in patients aids in the treatment process. Selecting antibiotics for therapy can be aided by testing the clinical isolates' susceptibility to antibiotics or determining the minimal inhibitory concentration. Antimicrobial resistance is a significant worldwide issue that is linked to longer hospital stays, higher treatment expenses, and higher rates of patient morbidity and death (Bacchetti, Schito, Milanese, Castellaro, & Alfei, 2024; Ramadan et al., 2024).

One major risk factor for the beginning and development of microbial resistance is the careless and inappropriate use of antibiotics in hospital and community settings. As a result, accurate and prompt identification can save time and resources. In this situation, the microbiologist makes sense because their job is to not only identify the infection's cause but also recommend the best course of action for treating it (Sibińska, Arendowski, Fijałkowski, Gabryś, & Pomastowski, 2024; Thomas et al., 2024).

Microbiology labs must have the equipment and resources necessary to identify microbiota and pollutants, identify microbes linked to diseases or for epidemiological purposes, and provide fast findings in emergencies for this to be done appropriately. Yeast and bacteria employ the anaerobic process of fermentation to produce energy. Glycolysis is the collection of early events for the breakdown of glucose, common to all fermentation forms (Saleh, Dheyab, Hadi, Hasan, & Jasim, 2024; Yu et al., 2024).

When this biochemical test is analyzed, it can be seen that the fermentation of the glucose results in a yellow color change, which is the outcome of the bacteria metabolizing sugar. Both bacteria and fungi are capable of anaerobic fermentation of fructose. Bacterial enzymes produce Carbon dioxide and ethanol from sugar (fructose or glucose). Its yellow color and turbidity are used to analyze its metabolism (Lee et al., 2024; Marciniak et al.).

Fermentation Type	Metabolized Substrate	Indicator of Fermentation	References
Glucose Fermentation	Glucose	Yellow color change	Lee et al., 2024; Marciniak et al.
Fructose Fermentation	Fructose	Yellow color change	Lee et al., 2024; Marciniak et al.
Lactic Acid Fermentation (Lactose)	Lactose	Yellow hue	Le et al., 2024; Ou et al., 2024
Sucrose Fermentation	Sucrose	Yellow color shift	Le et al., 2024; Ou et al., 2024
Maltose Fermentation	Maltose	Color change	Le et al., 2024; Ou et al., 2024

#### MICROBIAL FERMENTATION

Bacteria, namely lactobacilli, are responsible for lactic acid fermentation, which happens when glycolysis breaks down a lactose molecule and uses glucose and galactose as its primary mediator. A yellow hue is used to assess lactose fermentation. Fructose and glucose condense to generate sucrose, a molecule whose fermentation is indicated by a yellow color shift. Carbon, hydrogen, and oxygen atoms comprise the chemical molecule known as maltose, often glucose or sugar (Le, Duong, & Nguyen, 2024; Ou et al., 2024).

#### IMPACTS OF ANTIMICROBIAL RESISTANCE

Impacts	References
Longer hospital stays	Bacchetti et al., 2024
Higher treatment expenses	Bacchetti et al., 2024
Higher rates of patient morbidity and death	Bacchetti et al., 2024

Like other sugars, its metabolization can be determined by looking at the color change. Thus, this work aims to confirm the approach developed for Gram-positive bacteria microplate biochemical testing. The objective is to enhance the patient's bacterial treatment and identification while cutting expenses and raising throughput to enable more thorough patient analysis (Pagotto et al., 2024; Zhao et al., 2024).

### MATERIALS AND METHODS:

This study aims to confirm the approach of biochemical analysis for Gram-positive bacteria belonging to the Staphylococcus and Enterococcus genera on microplates through practical investigations of an experimental character. Abreu states that the initial inoculation was done in Cled Agar as well as Mannitol culture media to initiate the development of the microbes that will ultimately be utilized in the investigation (Golnari et al., 2024; Wu et al., 2024).

These media were prepared with a 10 ul loop, sowed, and baked for 24 hours at 35.2°C. 150 ml of deionized tap water, 50 ml of sodium chloride, and 3.2 g of phenol red broth were used to make the base solution. Then, using a magnetic stirrer, sugars (2.6 g lactose, 5 g glucose, 5 g fructose, 5 g maltose, and 5 g sucrose) were added and cooked until boiling. The culture medium was then transferred into test tubes and autoclaved for 15 minutes at 120°C (Min, Son, Jang, Yi, & Park, 2024; Semenzato et al., 2024).

The tests were performed on a tiny plate by putting 20 ul of solutions into each well and then adding 100 ul for each solution for the controls and the bacteria as soon as the tests were completed repeatedly in the laboratory tubes and the variations in color of the various mediums were noted. Of Staphylococcus and Enterococcus bacteria, using the Mc Farland scale (Cunha Del Vecchio et al., 2024; Gul, Rahman, Zafar, Abedien, & Malik, 2024).

To verify the fermentation process of the bacteria, triplicate experiments were conducted in each fundamental solution. Following the injection of the bacteria and control into the corresponding wells, the small plate was left to incubate at  $35\pm$  two °C for a whole day to see if fermentation occurred (Alharbi et al., 2024).

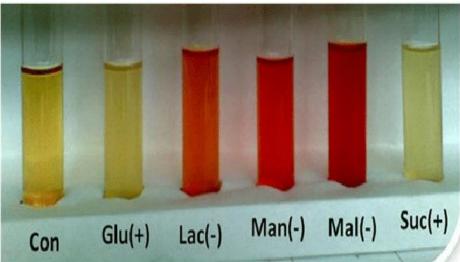
#### **RESULTS AND DISCUSSION:**

Following physiological and biochemical investigations and tests for catalase, coagulase, and novobiocin susceptibility, strains belonging to the Staphylococcus and Enterococcus genera were identified. The anaerobic bacterial identification system from Diagnostics s.r.o. is modeled after the Gram-positive bacterial identification system and the Corynebacterium identification system (Lenart-Boroń et al., 2024; Malú et al., 2024).

To determine whether a metabolic reaction was occurring, the reactions were first conducted in test tubes filled with the fundamental solutions of glucose, lactose, fructose, sucrose, and maltose, as seen in images 1 and 2 (Tiphaine et al., 2024).



**Image** 1 shows the initial reaction of Enterococcus faecalis bacteria in basic solutions. Net Negative Control is CN.



**Image 2** shows the initial response in the Staphylococcus aureus basic solution, with the control test tube on the right and the infected test tube on the left.

To validate fermentation, duplicate tests were run consecutively employing the two bacteria after examining the color shift. Three experiments were run in triplicate in total, yielding nine repeats in every base solution with identical outcomes as the initial responses for each microbe in the test tubes, as seen in photos 3 and 4 (Damoczi et al., 2024; Zahr Zahr, El Hajj, & Khalil, 2024).

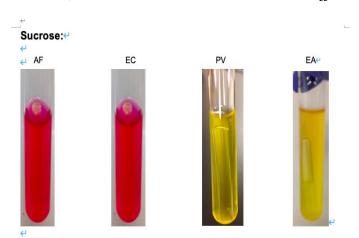


Image 3: Enterococcus faecalis fermentation in experiments conducted in triplicate.

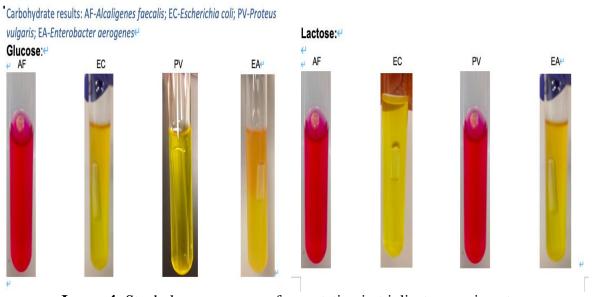


Image 4: Staphylococcus aureus fermentation in triplicate experiments.

The triplicate tests conducted in test tubes yielded satisfactory results, demonstrating that the bacteria had broken down the carbohydrates utilized and changed color.

Following the findings in the laboratory tubes, the procedures were repeated three times on the microplate, with a Gram-negative bacteria and the other sugars used as a comparative example, as seen in image 5 (Addisu, Fekadu, Hamza, & Adane, 2024; Mahony, 2024).

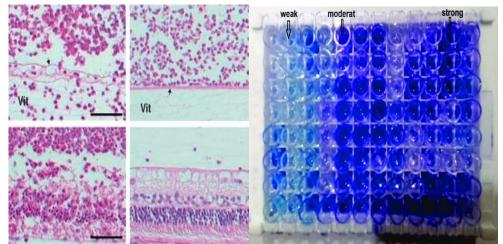


Image 5: Enterococcus faecalis, 1-3 Staphylococcus aureus, and 7-9 Gram-negative bacteria.

The carbohydrates were fermenting in the test tubes, but the microplate showed that the bacteria Staphylococcus aureus and Enterococcus faecalis were not metabolizing properly. In the microplate, Staphylococcus alone fermented maltose and fructose, and there was no noticeable change in color. The Enterococcus has fermented all the sugars, but the fructose has undergone some color change that makes it hard to see (Ben Yahia et al., 2024; Nye et al., 2024; Vitorino et al., 2024),

Comparatively speaking, the fermentation of the Gram-negative and Gram-positive bacteria was comparable. A common bacterial pathogen spread mostly by careless handling, Staphylococcus aureus is thought to be the microbe behind the majority of food poisoning outbreaks. Studies often concentrate on other sugars, including mannitol, and this bacterium is not known to ferment sucrose. Consequently, this fact might have played a role in the sugar's poor fermentation inside the microtiter plate (Alruwad, Salah El Dine, Gendy, Sabry, & El Hefnawy, 2024; Thuy et al., 2024; Weng et al., 2024).

Because the tubes are clear, which aids in a better study of the fermentation, more solution is also utilized. However, because the microplate has an impenetrable material and uses less solution, this may affect the results. The microplate results in the study that were utilized for evaluation assisted in recognizing the bacteria and caused the microplate to ferment the sugars as expected (Lei et al., 2024; Matei et al., 2024).

The effectiveness of the materials employed, the potential for bacterial contamination during injection, and the brief duration of the study could all have played a role in the dispersion of test findings used to verify this technique. Further research on this approach is highly relevant, as it is a biochemical analysis procedure that has not been accepted in the nation at large and has not been included in similar studies targeted at clinical diagnosis (Devanesan et al., 2024; Ferrando et al., 2024; Rammali et al., 2024).

#### **CONCLUSION:**

To increase the efficacy of antimicrobial medications and lower the death rate, it is essential to identify dangerous microorganisms and examine how they react to treatments.

Based on the results, it is evident that the biochemical evaluation technique for identifying Grampositive bacteria using microplates was not successfully validated because some sugars did not ferment, which made it difficult to determine the bacteria employing this methodology.

This method is distinguished by using less basic solutions than test tubes, which yield better results in metabolizing sugars and need a larger volume of solution. This may cause problems with the inability to metabolize particular sugars in the microplate. Execute the examinations. Because this is a novel methodology, more research is required to validate and enhance it.

#### **REFERENCE:**

- 1. Addisu, D., Fekadu, K., Hamza, S., & Adane, L. (2024). Antibacterial activity and phytochemical investigation of leaf and root extracts of Aloe gilbertii Reynolds.
- Alharbi, M., Azeez, Z. F., Alhussain, H. M., Shahlol, A. M., Albureikan, M. O., Elsehrawy, M., ... Ghareeb, A. (2024). Tapping The Biosynthetic Potential of Marine Bacillus Licheniformis LHG166, A Prolific Sulphated Exopolysaccharide Producer: Structural Insights, Bio-Prospecting Its Antioxidant, Antifungal, Antibacterial and Anti-Biofilm Potency As A Novel Anti-Infective Lead. Frontiers in Microbiology, 15, 1385493.
- 3. Alruwad, M. I., Salah El Dine, R., Gendy, A. M., Sabry, M. M., & El Hefnawy, H. M. (2024). Exploring the Biological and Phytochemical Potential of Jordan's Flora: A Review and Update of Eight Selected Genera from Mediterranean Region. Molecules, 29(5), 1160.
- 4. Bacchetti, F., Schito, A. M., Milanese, M., Castellaro, S., & Alfei, S. (2024). Anti Gram-Positive Bacteria Activity of Synthetic Quaternary Ammonium Lipid and Its Precursor Phosphonium Salt. International Journal of Molecular Sciences, 25(5), 2761.
- 5. Ben Yahia, H., Trabelsi, I., Arous, F., García-Vela, S., Torres, C., & Ben Slama, K. (2024). Detection of linezolid and vancomycin-resistant Enterococcus isolates collected from healthy chicken caecum. Journal of Applied Microbiology, 135(2), lxae027.
- Cunha Del Vechio, M. A., Bezerra, K., Estevam dos Santos, J., Midori Ono, J., Carvalho dos Santos, I., Dib Gonçalves, D., & Nunes Barbosa, L. (2024). Cymbopogon citratus Essential Oil in Controlling of Bacteria Associated with Oral Cavity of Dogs In Vitro and Dry Pet Feed. Foodborne Pathogens and Disease.
- Damoczi, J., Knoops, A., Martou, M.-S., Jamaux, F., Gabant, P., Mahillon, J., . . . Hols, P. (2024). Uncovering the class II-bacteriocin predatiome in salivarius streptococci. bioRxiv, 2024.2003. 2004.583286.
- 8. Devanesan, S., David, H. A., Ranjitsingh, A. J., Alzahim, T., Selvam, R., & AlSalhi, M. S. (2024). Efficient biogenesis of calcium oxide nanoparticles using the extract of Eleusine coracana seeds and their application against multidrug-resistant ocular bacterial pathogens. Environmental Research, 118632.
- 9. Fajdek-Bieda, A., Pawlińska, J., Wróblewska, A., & Łuś, A. (2024). Evaluation of the

Antimicrobial Activity of Geraniol and Selected Geraniol Transformation Products against Gram-Positive Bacteria. Molecules, 29(5), 950.

- Ferrando, N., Pino-Otín, M. R., Ballestero, D., Lorca, G., Terrado, E. M., & Langa, E. (2024). Enhancing Commercial Antibiotics with Trans-Cinnamaldehyde in Gram-Positive and Gram-Negative Bacteria: An In Vitro Approach. Plants, 13(2), 192.
- Golnari, M., Bahrami, N., Milanian, Z., Rabbani Khorasgani, M., Asadollahi, M. A., Shafiei, R., & Fatemi, S. S.-A. (2024). Isolation and characterization of novel Bacillus strains with superior probiotic potential: comparative analysis and safety evaluation. Scientific Reports, 14(1), 1457.
- 12. Gul, K., Rahman, K., Zafar, N., Abedien, Z. U., & Malik, A. (2024). ASSESSMENT OF MULTIDRUG RESISTANCE IN FALCON BACTERIAL ISOLATES: IMPLICATIONS FOR THERAPEUTIC APPROACHES. Journal of Population Therapeutics and Clinical Pharmacology, 31(3), 34-42.
- Jaber, D., Younes, N., Khalil, E., Albsoul-Younes, A., Zawiah, M., & Al-Bakri, A. G. (2024). Studying Microbial Ecology of Diabetic Foot Infections: Significance of PCR Analysis for Prudent Antimicrobial Stewardship. The International Journal of Lower Extremity Wounds, 15347346241230288.
- 14. Le, T. T., Duong, H. T., & Nguyen, C. H. (2024). Characterization of in vitro antimicrobial activity of gyrophoric acid isolated from Parmotrema indicum on methicillin-resistant Staphylococcus aureus. Journal of Applied Pharmaceutical Science, 14(3), 045-054.
- Lee, R. T., Evanowski, R. L., Greenbaum, H. E., Pawloski, D. A., Wiedmann, M., & Martin, N. H. (2024). Troubleshooting high laboratory pasteurization counts in organic raw milk requires characterization of dominant thermoduric bacteria, which includes non-sporeformers as well as sporeformers. Journal of Dairy Science.
- 16. Lei, Y., Yan, Y., Zhong, J., Zhao, Y., Xu, Y., Zhang, T., . . . Zhang, K. (2024). Enterococcus durans 98D alters gut microbial composition and function to improve DSS-induced colitis in mice. Heliyon.
- 17. Lenart-Boroń, A., Stankiewicz, K., Bulanda, K., Czernecka, N., Heliasz, M., Hunter, W., ... Khachatryan, G. (2024). In Vitro Antibacterial Activity of Ozonated Olive Oil against Bacteria of Various Antimicrobial Resistance Profiles Isolated from Wounds of Companion Animals. International Journal of Molecular Sciences, 25(6), 3557.
- 18. Mahony, J. (2024). Biological and bioinformatic tools for the discovery of unknown phagehost combinations. Current Opinion in Microbiology, 77, 102426.
- 19. Malú, Q., Caldeira, G. I., Catarino, L., Indjai, B., da Silva, I. M., Lima, B., & Silva, O. (2024). Ethnomedicinal, Chemical, and Biological Aspects of Lannea Species—A Review. Plants, 13(5), 690.
- 20. Marciniak, T., Kirchner, L., Wolf, S. A., Walther, B., Bischler, T., Nyasinga, J., . . . Whitelaw, A. Emergence of a Transferable Daptomycin Resistance Mechanism in Gram-Positive Bacteria: A Novel Membrane-Associated ABC Transporter Mediates High-Level Daptomycin Resistance in Staphylococci.
- Matei, S.-C., Dumitru, C. S., Fakhry, A. M., Ilijevski, N., Pešić, S., Petrović, J., . . . Olariu, S. (2024). Bacterial Species Involved in Venous Leg Ulcer Infections and Their Sensitivity to Antibiotherapy—An Alarm Signal Regarding the Seriousness of Chronic Venous Insufficiency C6 Stage and Its Need for Prompt Treatment. Microorganisms, 12(3), 472.
- 22. Min, J., Son, Y., Jang, I., Yi, C., & Park, W. (2024). Managing two simultaneous issues in concrete repair: Healing microcracks and controlling pathogens. Construction and Building Materials, 416, 135125.
- 23. Nye, T. M., Zou, Z., Obernuefemann, C. L., Pinkner, J. S., Lowry, E., Kleinschmidt, K., . . . Flores-Mireles, A. L. (2024). Microbial co-occurrences on catheters from long-term catheterized patients. Nature communications, 15(1), 61.
- 24. Ou, Y.-H., Chang, Y.-T., Chen, D.-P., Chuang, C.-W., Tsao, K.-C., Wu, C.-H., . . . Huang, C.-G. (2024). Benefit analysis of the auto-verification system of intelligent inspection for

microorganisms. Frontiers in Microbiology, 15, 1334897.

- 25. Pagotto, A., Campanile, F., Conti, P., Prataviera, F., Della Siega, P., Flammini, S., . . . Sartor, A. (2024). An Aminoglycoside-Sparing Regimen with Double Beta-Lactam to Successfully Treat Granulicatella adiacens Prosthetic Aortic Valve Endocarditis—Time to Change Paradigm? Infectious Disease Reports, 16(2), 249-259.
- 26. Ramadan, A., Abdel-Monem, M. O., El-Dougdoug, N. K., Mekky, A. E., Elaskary, S. A., Al-Askar, A. A., . . . Saied, E. (2024). Fully Characterized Effective Bacteriophages Specific against Antibiotic-Resistant Enterococcus faecalis, the Causative Agent of Dental Abscess. Medicina, 60(3), 501.
- Rammali, S., Rahim, A., El Aalaoui, M., Bencharki, B., Dari, K., Habach, A., . . . Khattabi, A. (2024). Antimicrobial potential of Streptomyces coeruleofuscus SCJ isolated from microbiologically unexplored garden soil in Northwest Morocco. Scientific Reports, 14(1), 3359.
- 28. Saleh, R. O., Dheyab, A. S., Hadi, B. H., Hasan, R. N., & Jasim, S. A. (2024). Effect of Ethanolic Extract of Syzygium Aromaticum Plant Against Enterococcus faecalis Isolated from Women with Urinary Tract Infections. Archives of Clinical Infectious Diseases, 19(1).
- Semenzato, G., Bernacchi, A., Amata, S., Bechini, A., Berti, F., Calonico, C., . . . Piccionello, A. P. (2024). Antibacterial Properties of Bacterial Endophytes Isolated from the Medicinal Plant Origanum heracleoticum L. Frontiers in Bioscience-Landmark, 29(3), 111.
- 30. Sibińska, E., Arendowski, A., Fijałkowski, P., Gabryś, D., & Pomastowski, P. (2024). Comparison of the Bruker Microflex LT and Zybio EXS2600 MALDI TOF MS systems for the identification of clinical microorganisms. Diagnostic Microbiology and Infectious Disease, 108(2), 116150.
- Thomas, J. K., Clark, J., Arora, V., Burgess, D. S., Burgess, D. R., Mynatt, R. P., ... Cotner, S. E. (2024). Performance of ePlex<sup>®</sup> Blood Culture Identification Panels in Clinical Isolates and Characterization of Antimicrobial Stewardship Opportunities. Diagnostic Microbiology and Infectious Disease, 116269.
- 32. Thuy, T. T. D., Lu, H.-F., Bregente, C. J. B., Huang, F.-C. A., Tu, P.-C., & Kao, C.-Y. (2024). Characterization of the broad-spectrum antibacterial activity of bacteriocin-like inhibitory substance-producing probiotics isolated from fermented foods. BMC microbiology, 24(1), 85.
- 33. Tiphaine, G., Céline, D.-C., Anne-Laure, R., Eve, T., Adeline, B.-D., Camille, C., . . . Hélène, M. (2024). A prospective multicenter evaluation of BioFire® Joint Infection Panel for the rapid microbiological documentation of acute arthritis. Clinical Microbiology and Infection.
- Vitorino, I. R., Pinto, E., Martín, J., Mackenzie, T. A., Ramos, M. C., Sánchez, P., . . . Reyes, F. (2024). Uncovering the biotechnological capacity of marine and brackish water Planctomycetota. Antonie van Leeuwenhoek, 117(1), 26.
- 35. Weng, S., Leng, G., Gao, J., Wang, Y., Yao, J., Li, X., . . . Tang, W. (2024). Garvicin-SHAMU-LG6, A Novel Bacteriocin from Lactococcus garvieae That Exert Broad Antimicrobial Activity Against Drug-Resistant Pathogens.
- 36. Wu, M., Kang, J., Tao, J., Yang, Y., Li, G., & Jia, W. (2024). Clinical Characteristics and Drug Resistance Mechanisms of Linezolid-Non-Susceptible Enterococcus in a Tertiary Hospital in Northwest China. Infection and Drug Resistance, 485-494.
- 37. Yu, J., Zhang, L., Gao, D., Wang, J., Li, Y., & Sun, N. (2024). Comparison of metagenomic next-generation sequencing and blood culture for diagnosis of bloodstream infections. Frontiers in Cellular and Infection Microbiology, 14.
- 38. Zahr, R., Zahr, S., El Hajj, R., & Khalil, M. (2024). Characterization of Actinobacteria strains in Lebanese soil with an emphasis on investigating their antibacterial activity. Brazilian Journal of Microbiology, 1-13.
- 39. Zhao, M., He, S., Wen, R., Li, C., Chen, X., Lin, X., . . . Tang, Y. (2024). Membrane vesicles derived from Enterococcus faecalis promote the co-transfer of important antibiotic resistance genes located on both plasmids and chromosomes. Journal of Antimicrobial Chemotherapy, 79(2), 320-326.