

FIȘĂ DE VERIFICARE A ÎNDEPLINIRII CRITERIILOR PENTRU ABILITARE
COMISIA BIOLOGIE ȘI BIOCHIMIE

ORDIN 6129/2016

Candidat: CS I dr. Cristian COMAN

Necesar	Existent
A1. Calificarea profesională: titlul de Doctor în specialitatea disciplinei postului sau înrudită cu aceasta	Doctor în Biologie, conferit în baza Ordinului Ministrului Educației, Cercetării, Tineretului și Sportului nr. 6468/07.12.2011.
A2. Articole științifice ca autor principal: pentru Profesor (CS I; Abilitare): minimum 4 articole în reviste cotate ISI cu AIS cumulat mai mare sau egal cu 4, din care 2 articole cu AIS de cel puțin 0,3 în ultimii 5 ani.	În ultimii 5 ani am publicat 8 articole în reviste cotate ISI, ca autor principal, cu AIS cumulat = $9 \geq 4$. Dintre acestea, 7 articole au AIS $\geq 0,3$.
A3. Coordonare proiecte de cercetare obținute prin competiție națională sau internațională: - pentru Profesor (CS I; Abilitare): minimum două granturi naționale de cercetare în calitate de director (sau responsabil de proiect în cazul parteneriatelor) sau unul național (în calitate de director) și unul internațional (în calitate de responsabil național).	Am coordonat 6 proiecte: 2 granturi internaționale, 2 granturi naționale, 1 proiect de tip COST și 1 bursă POSDRU.
B.1. Evaluarea activității de cercetare – Recunoaștere internațională minim 150 puncte	456,11 puncte
B.1. Evaluarea activității de cercetare – Performanță totală minim 250 puncte	520,32 puncte

Data,

Semnătura,

**Informații privind proiectele coordonate și calculul punctajelor de recunoaștere
internațională și performanță totală**

A2. Articole științifice ca autor principal: pentru Profesor (CS I; Abilitare): minimum 4 articole în reviste cotate ISI cu AIS cumulat mai mare sau egal cu 4, din care 2 articole cu AIS de cel puțin 0,3 în ultimii 5 ani.

În ultimii 5 ani am publicat 8 articole în reviste cotate ISI, ca autor principal (detaliate mai jos), cu AIS cumulat = $9 \geq 4$. Dintre acestea, 7 articole au AIS $\geq 0,3$.

1. Remizovschi, A., Carpa, R., Forray, F.L., Chiriac, C., Roba, C.-A., Beldean-Galea, S., Andrei, A.-S., Szekeres, E., Baricz, A., Lupan, I., Knut, R., **Coman, C.** 2020. Mud volcanoes and the presence of PAHs. *Scientific Reports*, 10:1253.
2. Baricz, A., Teban, A., Chiriac, C.M., Szekeres, E., Farkas, A., Nica, M., Dascălu, A., Oprișan, C., Lavin, P., **Coman, C.** 2018. Investigating the potential use of an Antarctic variant of *Janthinobacterium lividum* for tackling antimicrobial resistance in a One Health approach. *Scientific Reports*, 8(1):15272. (AIS=1,9)
3. Szekeres, E., Chiriac, C.M., Baricz, A., Szőke-Nagy, T., Lung, I., Soran, M.-L., Rudi, K., Dragos, N., **Coman, C.** 2018. Investigating antibiotics, antibiotic resistance genes, and microbial contaminants in groundwater in relation to the proximity of urban areas. *Environmental Pollution*, 236:734-744. (AIS=1,3)
4. Chiriac, C.M., Baricz, A., Szekeres, E., Rudi, K., Dragoș, N., **Coman, C.** 2018. Microbial composition and diversity patterns in deep hyperthermal aquifers from the Western Plain of Romania. *Microbial Ecology*, 75(1): 38-51. (AIS=1).
5. Chiriac, C.M., Szekeres, E., Rudi, K., Baricz, A., Hegedus, A., Dragoș, N., **Coman, C.** 2017. Differences in temperature and water chemistry shape distinct diversity patterns in thermophilic microbial communities. *Applied and Environmental Microbiology*, 83(21): e01363-17. (AIS=1,2)
6. Soran, M.-L., Lung, I., Opriș, O., Floare-Avram, V., **Coman, C.** 2017. Determination of Antibiotics in Surface Water by Solid-Phase Extraction and High-Performance Liquid Chromatography with Diode Array and Mass Spectrometry Detection. *Analytical Letters*, 50(7): 1209-1218. (AIS=0,2)
7. Opriș, O., Soran, M.L., Lung, I., Trușcă, M.R.C., Szoke-Nagy, T., **Coman, C.** 2017. The optimization of the antibiotics extraction from wastewaters and manure using Box–Behnken experimental design. *International Journal of Environmental Science and Technology*, 14(3): 473-480. (AIS=0,4).
8. Szekeres, E., Baricz, A., Chiriac, C.M., Farkas, A., Opris, O., Soran, M.-L., Andrei, A.-S., Rudi, K., Balcázar, J.L., Dragos, N., **Coman, C.** 2017. Abundance of antibiotics, antibiotic resistance genes and bacterial community composition in wastewater effluents from different Romanian hospitals. *Environmental Pollution*, 225:304-315. (AIS=1,11)

A3. Coordonare proiecte de cercetare obținute prin competiție națională sau internațională:

- pentru Profesor (CS I; Abilitare): minimum două granturi naționale de cercetare în calitate de director (sau responsabil de proiect în cazul parteneriatelor) sau unul național (în calitate de director) și unul internațional (în calitate de responsabil național); nu se iau în considerare granturi finanțate de propria instituție, granturi pentru participare la congrese, granturi de cercetare din finanțarea de baza de ex. programul Nucleu.

Granturi internaționale ca director de proiect/responsabil național:

1. Biodiversity as an ecological barrier for the spread of clinically relevant antibiotic resistance in the environment – Antiversa.

H2020 ERA-Net, contract nr. 117/2020. Implementare: 2020-2023. Valoare: 980.000 RON/200.000 Euro.

2. Ghid metodologic de monitorizare a antibioticelor și a rezistenței antimicrobiene în mediu ca instrument suport pentru îmbunătățirea managementului calității apelor de suprafață și a pânzei freactice – EnviroAMR.

EEA Grants, contract nr. 3499/20.05.2015. Implementare: 2015-2016. Valoare: 4.444.649 RON/1.007.285 Euro

Granturi naționale ca director de proiect:

1. Bacterii antarctice contra patogenilor umani: în căutare de noi compuși antimicrobieni – AntarcticPharma.

Contract nr. 140PED/2017. Implementare: 2017-2018. Valoare: 600.000 RON.

2. Biodiversitatea bacteriilor și archaeonilor dintr-un stromatolit modern din România ca posibili indicatori ai procesului de mineralizare.

Contract nr. PD 104/2012. Implementare: 2012-2013. Valoare: 300.000 RON

Alte proiecte coordonate:

1. Cyanobacteria blooms and toxins in water resources: Occurrence, impacts and management.

ESSEM COST Action ES1105.

Responsabil din partea ICB Cluj-Napoca

2. Biodiversitatea și profilul metabolic al comunităților microbiene implicate în formarea de stromatolite moderne investigate prin studii de metagenomică și metatranscriptomică.

Bursă POSDRU/159/1.5/S/133391.

B. Criterii și standarde minimale

B.1. Evaluarea activității de cercetare.

A fost efectuată utilizând baza de date Scopus (www.scopus.com), accesată la data de 09.03.2020:

Punctaj recunoaștere internațională: **458,76 puncte > 150 puncte.**

Punctaj performanță totală: **522,22 puncte > 250 puncte.**

Articole în reviste cotate ISI, ca autor principal:

1. Remizovschi, A., Carpa, R., Forray, F.L., Chiriac, C., Roba, C.-A., Beldean-Galea, S., Andrei, A.-Ș., Szekeres, E., Baricz, A., Lupan, I., Knut, R., Coman, C. 2020. Mud volcanoes and the presence of PAHs. Scientific Reports, 10(1):1253. (AIS=1,9)

Citări: 0

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,9)] = 17,3$ puncte

2. Baricz, A., Teban, A., Chiriac, C.M., Szekeres, E., Farkas, A., Nica, M., Dascălu, A., Opreșan, C., Lavin, P., Coman, C. 2018. Investigating the potential use of an Antarctic variant of *Janthinobacterium lividum* for tackling antimicrobial resistance in a One Health approach. Scientific Reports, 8(1):15272. (AIS=1,9)

Citări:

Oh W.T., Giri S.S., Yun S., Kim H.J., Kim S.G., et al. 2019. *Janthinobacterium lividum* as an emerging pathogenic bacterium affecting rainbow trout (*Oncorhynchus mykiss*) fisheries in Korea. Pathogens, 8(3):146.

Rajawat M.V.S., Singh R., Singh D., Saxena A.K. 2019. Psychrotrophs of the genus *Janthinobacterium* with potential to weather potassium aluminosilicate mineral. 3 Biotech, 9(4):142.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,9) + 2] = 19,3$ puncte

3. Szekeres, E., Chiriac, C.M., Baricz, A., Szőke-Nagy, T., Lung, I., Soran, M.-L., Rudi, K., Dragos, N., Coman, C. 2018. Investigating antibiotics, antibiotic resistance genes, and microbial contaminants in groundwater in relation to the proximity of urban areas. Environmental Pollution, 236:734-744. (AIS=1,3)

Citări:

Ding H., Qiao M., Zhong J., Zhu Y., Guo C., et al. 2020. Characterization of antibiotic resistance genes and bacterial community in selected municipal and industrial sewage treatment plants beside Poyang Lake. Water Research, 174:115603.

Wu D.-L., Zhang M., He L.-X., Zou H.-Y., Liu Y.-S., et al. 2020. Contamination profile of antibiotic resistance genes in ground water in comparison with surface water. Science of the Total Environment, 715:136975.

- Wang S., Ma X., Liu Y., Yi X., Du G., Li J. 2020. Fate of antibiotics, antibiotic-resistant bacteria, and cell-free antibiotic-resistant genes in full-scale membrane bioreactor wastewater treatment plants. *Bioresource Technology*, 302:122825.
- Xiang Y., Yang X., Xu Z., Hu W., Zhou Y., et al. 2020. Fabrication of sustainable manganese ferrite modified biochar from vinasse for enhanced adsorption of fluoroquinolone antibiotics: Effects and mechanisms. *Science of the Total Environment*, 709:136079.
- Qin L.-T., Pang X.-R., Zeng H.-H., Liang Y.-P., Mo L.-Y., et al. 2020. Ecological and human health risk of sulfonamides in surface water and groundwater of Huixian karst wetland in Guilin, China. *Science of the Total Environment*, 708:134552.
- Xiang S., Wang X., Ma W., Liu X., Zhang B., et al. 2020. Response of microbial communities of karst river water to antibiotics and microbial source tracking for antibiotics. *Science of the Total Environment*, 706: 135730.
- Wang W.-N., Zhang C.-Y., Zhang M.-F., Pei P., Zhou W., et al. 2020. Precisely photothermal controlled releasing of antibacterial agent from Bi₂S₃ hollow microspheres triggered by NIR light for water sterilization. *Chemical Engineering Journal*, 381:122630.
- Wang J., Zhuang R. 2020. Degradation of antibiotics by advanced oxidation processes: An overview. *Science of the Total Environment*, 701:135023.
- Felis E., Kalka J., Sochacki A., Kowalska K., Bajkacz S., et al. 2020. Antimicrobial pharmaceuticals in the aquatic environment - occurrence and environmental implications. *European Journal of Pharmacology*, 866:172813.
- Gao Y., Xia J., Liu D., Kang R., Yu G., Deng S. 2019. Synthesis of mixed-linker Zr-MOFs for emerging contaminant adsorption and photodegradation under visible light. *Chemical Engineering Journal*, 378:122118.
- Na G., Wang C., Gao H., Li R., Jin S., et al. 2019. The occurrence of sulfonamide and quinolone resistance genes at the Fildes Peninsula in Antarctica. *Marine Pollution Bulletin*, 149:110503.
- Jiang X., Liu S., Yang M., Rasooly A. 2019. Amperometric genosensor for culture independent bacterial count. *Sensors and Actuators, B: Chemical*, 299:126944.
- Nnadozie C.F., Odume O.N. 2019. Freshwater environments as reservoirs of antibiotic resistant bacteria and their role in the dissemination of antibiotic resistance genes. *Environmental Pollution*, 254:113067.
- Harb M., Wang P., Zarei-Baygi A., Plumlee M.H., Smith A.L. 2019. Background Antibiotic Resistance and Microbial Communities Dominate Effects of Advanced Purified Water Recharge to an Urban Aquifer. *Environmental Science and Technology Letters*, 6(10):578-584.
- Zheng X., Jiang B., Lang H., Zhang R., Li Y., et al. 2019. Effects of Antibiotics on Microbial Communities Responsible for Perchlorate Degradation. *Water, Air, and Soil Pollution*, 230(10):244.
- Kurwadkar S. 2019. Occurrence and distribution of organic and inorganic pollutants in groundwater. *Water Environment Research*, 91(10):1001-1008.

- Fitzpatrick M.C., Bauch C.T., Townsend J.P., Galvani A.P. 2019. Modelling microbial infection to address global health challenges. *Nature Microbiology*, 4(10):1612-1619.
- Veiga-Gómez M., Nebot C., Miranda J.M., Vázquez B., Verdes S., et al. 2019. Consumption of pharmaceuticals by dairy cows via watering trough: Uncontrolled intake. *Agriculture, Ecosystems and Environment*, 280:95-101.
- Ye M., Sun M., Huang D., Zhang Z., Zhang H., et al. 2019. A review of bacteriophage therapy for pathogenic bacteria inactivation in the soil environment. *Environment International*, 129:488-496.
- Peng C., Ba J., Hu F., Pan X., Jiao Y., et al. Typical antibiotic pollution characteristics and ecological risk assessment of Huixian Karst wetland in Guangxi, China. *Huanjing Kexue Xuebao/Acta Scientiae Circumstantiae*, 39(7):2207-2217.
- Bytesnikova Z., Richtera L., Smerkova K., Adam V. 2019. Graphene oxide as a tool for antibiotic-resistant gene removal: a review. *Environmental Science and Pollution Research*, 26(20):20148-20163.
- Hu Y., Zhang T., Jiang L., Luo Y., Yao S., et al. 2019. Occurrence and reduction of antibiotic resistance genes in conventional and advanced drinking water treatment processes. *Science of the Total Environment*, 669:777-784.
- Wang Y.-F., Fan J., Meng Z.-H., Xue M., Qiu L.-L. 2019. Fabrication of an antibiotic-sensitive 2D-molecularly imprinted photonic crystal. *Analytical Methods*, 11(22):2875-2879.
- Yu K., Sun C., Zhang B., Hassan M., He Y. 2019. Size-dependent adsorption of antibiotics onto nanoparticles in a field-scale wastewater treatment plant. *Environmental Pollution*, 248:1079-1087.
- Du H., Shi W., Zhang H., Liu Y., Li J. 2019. Distribution of antibiotic resistance genes in sediments of Hetao irrigation district. *Huanjing Kexue Xuebao/Acta Scientiae Circumstantiae*, 39(4):1257-1265.
- Sun K., Sun Y., Gao B., Xu H., Wu J. 2019. Effect of cation type in mixed Ca-Na systems on transport of sulfonamide antibiotics in saturated limestone porous media. *Environmental Science and Pollution Research*, 26(11):11170-11178.
- Schorr B., Ghanem H., Rosiwal S., Geißdörfer W., Burkovski A. 2019. Elimination of bacterial contaminations by treatment of water with boron-doped diamond electrodes. *World Journal of Microbiology and Biotechnology*, 35(3):48.
- Li D., Gu A.Z. 2019. Antimicrobial resistance: A new threat from disinfection byproducts and disinfection of drinking water? *Current Opinion in Environmental Science and Health*, 7(83):91.
- Zou H., He J.-T., He B.-N., Lao T.-Y., Liu F., Guan X.-Y. 2019. Sensitivity assessment of denitrifying bacteria against typical antibiotics in groundwater. *Environmental Science: Processes and Impacts*, 21(9):1570-1579.
- Bartley P.S., Domitrovic T.N., Moretto V.T., Santos C.S., Ponce-Terashima R., et al. 2019. Antibiotic resistance in enterobacteriaceae from surface waters in Urban Brazil highlights

the risks of poor sanitation. *American Journal of Tropical Medicine and Hygiene*, 100(6):1369-1377.

- Amarasiri M., Sano D., Suzuki S. 2019. Understanding human health risks caused by antibiotic resistant bacteria (ARB) and antibiotic resistance genes (ARG) in water environments: Current knowledge and questions to be answered. *Critical Reviews in Environmental Science and Technology*, DOI: 10.1080/10643389.2019.1692611.
- Jurado A., Walther M., Díaz-Cruz M.S. 2019. Occurrence, fate and environmental risk assessment of the organic microcontaminants included in the Watch Lists set by EU Decisions 2015/495 and 2018/840 in the groundwater of Spain. *Science of the Total Environment*, 663:285-296.
- Qiu H., Ling C., Yuan R., Liu F., Li A. 2019. Bridging effects behind the coadsorption of copper and sulfamethoxazole by a polyamine-modified resin. *Chemical Engineering Journal*, 362:422-429.
- Hao, H., Shi, D.-Y., Yang, D., Yang, Z.-W., Qiu, Z.-G., et al. 2019. Profiling of intracellular and extracellular antibiotic resistance genes in tap water. *Journal of Hazardous Materials*, 365:340-345.
- Sun, M., Ye, M., Zhang, Z., Zhang, S., Zhao, Y., et al. 2019. Biochar combined with polyvalent phage therapy to mitigate antibiotic resistance pathogenic bacteria vertical transfer risk in an undisturbed soil column system. *Journal of Hazardous Materials*, 365:1-8.
- Kumar M., Jaiswal S., Sodhi K.K., Shree P., Singh D.K., et al. 2019. Antibiotics bioremediation: Perspectives on its ecotoxicity and resistance. *Environment International*, 124: 448-461.
- Wen, X., Lu, J., Wu, J., Lin, Y., Luo, Y. 2019. Influence of coastal groundwater salinization on the distribution and risks of heavy metals. *Science of the Total Environment*, 652:267-277.
- Huang, F., Zou, S., Deng, D., Lang, H., Liu, F. 2019. Antibiotics in a typical karst river system in China: Spatiotemporal variation and environmental risks. *Science of the Total Environment*, 650:1348-1355.
- Sharma, B.M., Bečanová, J., Scheringer, M., Sharma, A., Bharat, G.K., et al. 2019. Health and ecological risk assessment of emerging contaminants (pharmaceuticals, personal care products, and artificial sweeteners) in surface and groundwater (drinking water) in the Ganges River Basin, India. *Science of the Total Environment*, 646:1459-1467.
- Zhou, M., Zhu, T., Fei, X. 2018. Enhanced adsorption performance of oxytetracycline by desugared reed residues. *International Journal of Environmental Research and Public Health*, 15(10): 2229.
- Lodeiro, C., Capelo, J.L., Oliveira, E., Lodeiro, J.F. 2019. New toxic emerging contaminants: beyond the toxicological effects. *Environmental Science and Pollution Research*. 26:1-4.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,3) + 41] = 54,1$ puncte

4. Chiriac, C.M., Baricz, A., Szekeres, E., Rudi, K., Dragoș, N., Coman, C. 2018. Microbial composition and diversity patterns in deep hyperthermal aquifers from the Western Plain of Romania. Microbial Ecology, 75(1): 38-51. (AIS=1).

Citări: 0

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1)] = 11$ puncte

5. Chiriac, C.M., Szekeres, E., Rudi, K., Baricz, A., Hegedus, A., Dragoș, N., Coman, C. 2017. Differences in temperature and water chemistry shape distinct diversity patterns in thermophilic microbial communities. Applied and Environmental Microbiology, 83(21): e01363-17. (AIS=1,2)

Citări:

Remizovschi A., Carpa R., Forray F.L., Chiriac C., Roba C.-A., et al. 2020. Mud volcanoes and the presence of PAHs. *Scientific Reports*, 10(1):1253.

Palmer-Young E.C., Ngor L., Burciaga Nevarez R., Rothman J.A., Raffel T.R., McFrederick Q.S. 2019. Temperature dependence of parasitic infection and gut bacterial communities in bumble bees. *Environmental Microbiology*, 21(12):4706-4723.

Zhang L., Cai Y., Jiang M., Dai J., Guo X., et al. 2019. The levels of microbial diversity in different water layers of saline Chagan Lake, China. *Journal of Oceanology and Limnology*, DOI: 10.1007/s00343-019-9027-7.

Tang J., Liang Y., Jiang D., Li L., Luo Y., et al. 2018. Temperature-controlled thermophilic bacterial communities in hot springs of western Sichuan, China. *BMC Microbiology*, 18(1):134.

Oliverio A.M., Power J.F., Washburne A., Cary S.C., Stott M.B., Fierer N. 2018. The ecology and diversity of microbial eukaryotes in geothermal springs. *ISME Journal*, 12(8):1918-1928.

Valeriani F., Crognale S., Protano C., Gianfranceschi G., Orsini M., et al. 2018. Metagenomic analysis of bacterial community in a travertine depositing hot spring. *New Microbiologica*, 41(2):126-135.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,2) + 6] = 18,4$ puncte

6. Soran, M.-L., Lung, I., Opreș, O., Floare-Avram, V., Coman, C. 2017. Determination of Antibiotics in Surface Water by Solid-Phase Extraction and High-Performance Liquid Chromatography with Diode Array and Mass Spectrometry Detection. Analytical Letters, 50(7): 1209-1218. (AIS=0,2)

Citări:

Gao L., Qin D., Huang X., Wu S., Chen Z., Tang S., Wang P. 2019. Determination of pesticides and Pharmaceuticals from Fish Cultivation Water by parallel solid-phase extraction (SPE) and liquid chromatography–quadrupole time-of-flight mass spectrometry (LC-QTOF-MS). *Analytical Letters*, 52(6):983-997.

Duan X.-Y., Zhang Y., Yan J.-Q., Zhou Y., Li G.-H., Feng X.-S. 2019. Progress in Pretreatment and Analysis of Cephalosporins: An Update Since 2005. *Critical Reviews in Analytical Chemistry*, DOI: 10.1080/10408347.2019.1676194.

Nagy-Kovács Z., László B., Fleit E., Czihat-Mártonné K., Till G., et al. 2018. Behavior of organic micropollutants during river bank filtration in Budapest, Hungary. *Water (Switzerland)*, 10(12):1861.

Szekeres E., Chiriac C.M., Baricz A., Szőke-Nagy T., Lung I., et al. 2018. Investigating antibiotics, antibiotic resistance genes, and microbial contaminants in groundwater in relation to the proximity of urban areas. *Environmental Pollution*, 236:734-744.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 0,2) + 4] = 9,4$ puncte

7. Opreș, O., Soran, M.L., Lung, I., Trușcă, M.R.C., Szoke-Nagy, T., Coman, C. 2017. The optimization of the antibiotics extraction from wastewaters and manure using Box–Behnken experimental design. *International Journal of Environmental Science and Technology*, 14(3): 473-480. (AIS=0,4).

Citări:

Styszko, K.aEmail Author, Szczurowski, J.a, Czuma, N.a, Makowska, D.a, Kistler, M.b, Uruski, Ł. 2018. Adsorptive removal of pharmaceuticals and personal care products from aqueous solutions by chemically treated fly ash. *International Journal of Environmental Science and Technology*, 15(3): 493-506.

Soran, M.-L., Opreș, O., Lung, I., Kacso, I., Poray, A.S., Stan, M. 2017. The efficiency of the multi-walled carbon nanotubes used for antibiotics removal from wastewaters generated by animal farms. *Environmental Science and Pollution Research*, 24(19): 16396-16406.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 0,4) + 2] = 8,8$ puncte

8. Szekeres, E., Baricz, A., Chiriac, C.M., Farkas, A., Opreș, O., Soran, M.-L., Andrei, A.-S., Rudi, K., Balcázar, J.L., Dragos, N., Coman, C. 2017. Abundance of antibiotics, antibiotic resistance genes and bacterial community composition in wastewater effluents from different Romanian hospitals. *Environmental Pollution*, 225:304-315. (AIS=1,11)

Citări:

Shi W., Liu Y., Li J., Zhang H., Shi R., et al. 2020. Distribution pattern of antibiotic resistance genes and bacterial community in agricultural soil samples of Wuliangshuai watershed, China. *Agriculture, Ecosystems and Environment*, 295:106884.

Sun J., Yang P., Huang S., Li N., Zhang Y., et al. 2020. Enhanced removal of veterinary antibiotic from wastewater by photoelectroactive biofilm of purple anoxygenic phototroph through photosynthetic electron uptake. *Science of the Total Environment*, 713:136605.

Wei T., Yao H., Sun P., Cai W., Li X., et al. 2020. Mitigation of antibiotic resistance in a pilot-scale system treating wastewater from high-speed railway trains. *Chemosphere*, 245:125484.

- Li H., Cheng W., Li B., Xu Y., Zheng X. 2020. The fate of antibiotic resistance genes during co-composting of swine manure with cauliflower and corn straw. *Bioresource Technology*, 300:122669.
- Makowska N., Philips A., Dabert M., Nowis K., Trzebny A., et al. 2020. Metagenomic analysis of β -lactamase and carbapenemase genes in the wastewater resistome. *Water Research*, 170:115277.
- Chen M.X., Zhang Y.Q., Chang J.L., Ma X.G., Lei T., et al. 2020. Occurrence of Antibiotic Resistance Genes in a Small Township Wastewater Treatment Plant and the Receiving River. *IOP Conference Series: Earth and Environmental Science*, 435(1):12012.
- Bardhan T., Chakraborty M., Bhattacharjee B. 2020. Prevalence of colistin-resistant, carbapenem-hydrolyzing proteobacteria in hospital water bodies and out-falls of West Bengal, India. *International Journal of Environmental Research and Public Health*, 17(3):1007.
- Chopyk J., Nasko D.J., Allard S., Bui A., Treangen T., et al. 2020. Comparative metagenomic analysis of microbial taxonomic and functional variations in untreated surface and reclaimed waters used in irrigation applications. *Water Research*, 169: 115250.
- Wielens Becker R., Ibáñez M., Cuervo Lumbaque E., Wilde M.L., Flores da Rosa T., et al. 2020. Investigation of pharmaceuticals and their metabolites in Brazilian hospital wastewater by LC-QTOF MS screening combined with a preliminary exposure and in silico risk assessment. *Science of the Total Environment*, 699:134218.
- Wang M., Liu H., Dai X. 2020. Dosage effects of lincomycin mycelial residues on lincomycin resistance genes and soil microbial communities. *Environmental Pollution*, 256:113392.
- Collins F., Rozhkovskaya A., Outram J.G., Millar G.J. 2020. A critical review of waste resources, synthesis, and applications for Zeolite LTA. *Microporous and Mesoporous Materials*, 291:109667.
- Xu K., Wang J., Gong H., Li Y., Zhou L., Yan M. 2019. Occurrence of antibiotics and their associations with antibiotic resistance genes and bacterial communities in Guangdong coastal areas. *Ecotoxicology and Environmental Safety*, 186:109796.
- Du B., Yang Q., Wang R., Wang R., Wang Q., Xin Y. 2019. Evolution of antibiotic resistance and the relationship between the antibiotic resistance genes and microbial compositions under long-term exposure to tetracycline and sulfamethoxazole. *International Journal of Environmental Research and Public Health*, 16(23):4681.
- Cabrera-Reina A., Martínez-Piernas A.B., Bertakis Y., Xekoukoulotakis N.P., Agüera A., Sánchez Pérez J.A. 2019. TiO₂ photocatalysis under natural solar radiation for the degradation of the carbapenem antibiotics imipenem and meropenem in aqueous solutions at pilot plant scale. *Water Research*, 166: 115037.
- Mills M.C., Lee J. 2019. The threat of carbapenem-resistant bacteria in the environment: Evidence of widespread contamination of reservoirs at a global scale. *Environmental Pollution*, 255:113143.

- Li Y., Wang X., Li J., Wang Y., Song J., et al. 2019. Effects of struvite-humic acid loaded biochar/bentonite composite amendment on Zn(II) and antibiotic resistance genes in manure-soil. *Chemical Engineering Journal*, 375:122013.
- Galindo-Miranda J.M., Guízar-González C., Becerril-Bravo E.J., Moeller-Chávez G., León-Becerril E., Vallejo-Rodríguez R. 2019. Occurrence of emerging contaminants in environmental surface waters and their analytical methodology - A review. *Water Science and Technology: Water Supply*, 19(7):1871-1884.
- Lye Y.L., Bong C.W., Lee C.W., Zhang R.J., Zhang G., et al. 2019. Anthropogenic impacts on sulfonamide residues and sulfonamide resistant bacteria and genes in Larut and Sangga Besar River, Perak. *Science of the Total Environment*, 688:1335-1347.
- Zaha D.C., Bungau S., Aleya S., Tit D.M., Vesa C.M., et al. 2019. What antibiotics for what pathogens? The sensitivity spectrum of isolated strains in an intensive care unit. *Science of the Total Environment*, 687:118-127.
- Limayem A., Wasson S., Mehta M., Pokhrel A.R., Patil S., et al. 2019. High-Throughput Detection of Bacterial Community and Its Drug-Resistance Profiling From Local Reclaimed Wastewater Plants. *Frontiers in Cellular and Infection Microbiology*, 9:303.
- Quintela-Baluja M., Abouelnaga M., Romalde J., Su J.-Q., Yu Y., et al. 2019. Spatial ecology of a wastewater network defines the antibiotic resistance genes in downstream receiving waters. *Water Research*, 162:347-357.
- Cacace D., Fatta-Kassinos D., Manaia C.M., Cytryn E., Kreuzinger N., et al. 2019. Antibiotic resistance genes in treated wastewater and in the receiving water bodies: A pan-European survey of urban settings. *Water Research*, 162:320-330.
- Wang Q., Wang D., Wang J., Cui Y., Xu H. 2019. Recent progress and novel perspectives of electrochemical sensor for cephalosporins detection. *International Journal of Electrochemical Science*, 14(9):8639-8649.
- Shi W., Zhang H., Li J., Liu Y., Shi R., et al. 2019. Occurrence and spatial variation of antibiotic resistance genes (ARGs) in the Hetao Irrigation District, China. *Environmental Pollution*, 251:792-801.
- Zhang H., He H., Chen S., Huang T., Lu K., et al. 2019. Abundance of antibiotic resistance genes and their association with bacterial communities in activated sludge of wastewater treatment plants: Geographical distribution and network analysis. *Journal of Environmental Sciences (China)*, 82:24-38.
- Miazek K., Brozek-Pluska B. 2019. Effect of phrs and pcps on microalgal growth, metabolism and microalgae-based bioremediation processes: A review. *International Journal of Molecular Sciences*, 20(10):2492.
- Farkas A., Tarco E., Butiuc-Keul A. 2019. Antibiotic resistance profiling of pathogenic enterobacteriaceae from cluj- napoca, Romania. *GERMS*, 9(1):17-27.
- Cioca G., Munteanu F.-D. 2019. Estimation of the amount of disposed antibiotics. *Sustainability (Switzerland)*, 11(6):1800.

- Jiang X., Cui X., Xu H., Liu W., Tao F., et al. 2019. Whole genome sequencing of extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli* isolated from a wastewater treatment plant in China. *Frontiers in Microbiology*, 10:1797.
- Eckstrom K., Barlow J.W. 2019. Resistome metagenomics from plate to farm: The resistome and microbial composition during food waste feeding and composting on a Vermont poultry farm. *PLoS ONE*, 14(11):e0219807.
- Amarasiri M., Sano D., Suzuki S. 2019. Understanding human health risks caused by antibiotic resistant bacteria (ARB) and antibiotic resistance genes (ARG) in water environments: Current knowledge and questions to be answered. *Critical Reviews in Environmental Science and Technology*, DOI: 10.1080/10643389.2019.1692611.
- Waseem H., Williams M.R., Jameel S., Hashsham S.A. 2018. Antimicrobial resistance in the environment. *Water Environment Research*, 90(10):865-884.
- Zhao Y., Ye L., Zhang X.-X. 2018. Emerging pollutants—part i: Occurrence, fate and transport. *Water Environment Research*, 90(10):1301-1322.
- Zad T.J., Astuti M.P., Padhye L.P. 2018. Fate of environmental pollutants. *Water Environment Research*, 90(10):1104-1170.
- Ao X., Sun W., Li S., Yang C., Li C., Lu Z. 2019. Degradation of tetracycline by medium pressure UV-activated peroxymonosulfate process: Influencing factors, degradation pathways, and toxicity evaluation. *Chemical Engineering Journal*, 361: 1053-1062.
- Chu L., Zhuan R., Chen D., Wang J., Shen Y. 2019. Degradation of macrolide antibiotic erythromycin and reduction of antimicrobial activity using persulfate activated by gamma radiation in different water matrices. *Chemical Engineering Journal*, 361: 156-166.
- Ory J., Bricheux G., Robin F., Togola A., Forestier C., Traore O. 2019. Biofilms in hospital effluents as a potential crossroads for carbapenemase-encoding strains. *Science of the Total Environment*, 657: 7-15.
- Zhao R., Feng J., Liu J., Fu W., Li X., Li B. 2019. Deciphering of microbial community and antibiotic resistance genes in activated sludge reactors under high selective pressure of different antibiotics. *Water Research*, 151: 388-402.
- Shen Y., Chu L., Zhuan R., Xiang X., Sun H., Wang J. 2019. Degradation of antibiotics and antibiotic resistance genes in fermentation residues by ionizing radiation: A new insight into a sustainable management of antibiotic fermentative residuals. *Journal of Environmental Management*, 232: 171-178.
- 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. *International Journal of Hygiene and Environmental Health*, S1438-4639(18)30828-9.
- Yi, X., Lin, C., Ong, E.J.L., Wang, M., Zhou, Z. 2019. Occurrence and distribution of trace levels of antibiotics in surface waters and soils driven by non-point source pollution and anthropogenic pressure. *Chemosphere*, 216:213-223.

- Wang, J., Zhuan, R., Chu, L. 2019. The occurrence, distribution and degradation of antibiotics by ionizing radiation: An overview. *Science of the Total Environment*, 646:1385-1397.
- Baricz, A., Teban, A., Chiriac, C.M., Szekeres, E., Farkas, A., et al. 2018. Investigating the potential use of an Antarctic variant of *Janthinobacterium lividum* for tackling antimicrobial resistance in a One Health approach. *Scientific Reports*, 8(1):15272.
- Yu, Z., Yin, D., Hou, M., Zhang, J. 2018. Effects of food availability on the trade-off between growth and antioxidant responses in *Caenorhabditis elegans* exposed to sulfonamide antibiotics. *Chemosphere*, 211:278-285.
- Zhang, Q.-Q., Tian, G.-M., Jin, R.-C. 2018. The occurrence, maintenance, and proliferation of antibiotic resistance genes (ARGs) in the environment: influencing factors, mechanisms, and elimination strategies. *Applied Microbiology and Biotechnology*, 102(19):8261-8274.
- Li, A., Chen, L., Zhang, Y., Tao, Y., Xie, H., Li, S., Sun, W., Pan, J., He, Z., Mai, C., Fan, Y., Xian, H., Zhang, Z., Wen, D. 2018. Occurrence and distribution of antibiotic resistance genes in the sediments of drinking water sources, urban rivers, and coastal areas in Zhuhai, China. *Environmental Science and Pollution Research*, 25(26):26209-26217.
- Ethica, S.N., Saptaningtyas, R., Muchlissin, S.I., Sabdono, A. 2018. The development method of bioremediation of hospital biomedical waste using hydrolytic bacteria. *Health and Technology*, 8(4):239-254.
- Bürgmann, H., Frigon, D., Gaze, W.H., Manaia, C.M., Pruden, A., et al. 2018. Water and sanitation: An essential battlefront in the war on antimicrobial resistance. *FEMS Microbiology Ecology*, 94(9): fiy101.
- Li, G., Yang, H., An, T., Lu, Y. 2018. Antibiotics elimination and risk reduction at two drinking water treatment plants by using different conventional treatment techniques. *Ecotoxicology and Environmental Safety*, 158:154-161.
- Petrillo, J.E., Ogunseitan, O.A. 2018. Emerging issues in the environmental context of antibiotic-resistance. *Environment International*, 116:39-42.
- Buelow, E., Bayjanov, J.R., Majoor, E., Willems, R.J.L., Bonten, M.J.M., et al. 2018. Limited influence of hospital wastewater on the microbiome and resistome of wastewater in a community sewerage system. *FEMS Microbiology Ecology*, 94(7): fiy087.
- Wang, G., Li, S., Ma, X., Qiao, J., Li, G., Zhang, H., Wang, J., Song, Y. 2018. A novel Z-scheme sonocatalyst system, $\text{Er}^{3+}:\text{Y}_3\text{Al}_5\text{O}_{12}@\text{Ni}(\text{Fe}_{0.05}\text{Ga}_{0.95})_2\text{O}_4\text{-Au-BiVO}_4$, and application in sonocatalytic degradation of sulfanilamide. *Ultrasonics Sonochemistry*, 45: 150-166.
- Szekeres, E., Chiriac, C.M., Baricz, A., Szőke-Nagy, T., Lung, I., et al. 2018. Investigating antibiotics, antibiotic resistance genes, and microbial contaminants in groundwater in relation to the proximity of urban areas. *Environmental Pollution*, 236: pp. 734-744.
- Zhang, Y., Niu, Z., Zhang, Y., Zhang, K. 2018. Occurrence of intracellular and extracellular antibiotic resistance genes in coastal areas of Bohai Bay (China) and the factors affecting them. *Environmental Pollution*, 236:126-136.

- Azanu, D., Styriahave, B., Darko, G., Weisser, J.J., Abaidoo, R.C. 2018. Occurrence and risk assessment of antibiotics in water and lettuce in Ghana. *Science of the Total Environment*, 622-623:293-305.
- Zhang, B., Wang, M.M., Wang, B., Xin, Y., Gao, J., Liu, H. 2018. The effects of bio-available copper on macrolide antibiotic resistance genes and mobile elements during tylosin fermentation dregs co-composting. *Bioresource Technology*, 251:230-237.
- Reina, A.C., Martínez-Piernas, A.B., Bertakis, Y., Brebou, C., Xekoukoulotakis, N.P., et al. 2018. Photochemical degradation of the carbapenem antibiotics imipenem and meropenem in aqueous solutions under solar radiation. *Water Research*, 128: 61-70.
- Choudri, B.S., Charabi, Y., Ahmed, M. 2018. Health effects associated with wastewater treatment, reuse and disposal. *Water Environment Research*, 90(10):1759-1776.
- Gao, H., Zhang, L., Lu, Z., He, C., Li, Q., Na, G. 2018. Complex migration of antibiotic resistance in natural aquatic environments. *Environmental Pollution*, 232:1-9.
- Adegoke A.A., Amoah I.D., Stenström T.A., Verbyla M.E., Mihelcic J.R. Epidemiological evidence and health risks associated with agricultural reuse of partially treated and untreated wastewater: A review. *Frontiers in Public Health*, 6:337.
- Chiriac, C.M., Szekeres, E., Rudi, K., Baricz, A., Hegedus, A., et al. 2017. Differences in temperature and water chemistry shape distinct diversity patterns in thermophilic microbial communities. *Applied and Environmental Microbiology*, 83(21): e01363-17.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,11) + 60] = 71,77$ puncte

9. Coman, C., Chiriac, C.M., Robeson, M., Ionescu, C., Dragoș, N., Barbu-Tudoran, L., Andrei, Ș.A., Banciu, H.L., Sicora, C., Podar, M., 2015. Structure, mineralogy, and microbial diversity of geothermal spring microbialites associated with a deep oil drilling in Romania. *Frontiers in Microbiology*, 6:253. (AIS=1,39)

Citări:

- Gómez-Acata E.S., Centeno C.M., Falcón L.I. 2019. Methods for extracting 'omes from microbialites. *Journal of Microbiological Methods*, 160(1):10.
- Zhang, B., Xu, X., Zhu, L. 2019. Activated sludge bacterial communities of typical wastewater treatment plants: distinct genera identification and metabolic potential differential analysis. *AMB Express*, 8:184.
- Liu Y.-C., Huang R.-M., Bao J., Wu K.-Y., Wu H.-Y., et al. 2018. The unexpected diversity of microbial communities associated with black corals revealed by high-throughput Illumina sequencing. *FEMS Microbiology Letters*, 365(15):fny167.
- Liang, X., Whitham, J.M., Holwerda, E.K., Shao, X., Tian, L., et al. 2018. Development and characterization of stable anaerobic thermophilic methanogenic microbiomes fermenting switchgrass at decreasing residence times. *Biotechnology for Biofuels*, 11(1):243.

- Thiel, V., Tank, M., Bryant, D.A. 2018. Diversity of Chlorophototrophic Bacteria Revealed in the Omics Era. *Annual Review of Plant Biology*, 69:21-49.
- Valdespino-Castillo, P.M., Hu, P., Merino-Ibarra, M., López-Gómez, L.M., Cerqueda-García, D., et al. 2018. Exploring biogeochemistry and microbial diversity of extant microbialites in Mexico and Cuba. *Frontiers in Microbiology*, 9:510.
- Louyakis, A.S., Gourelé, H., Casaburi, G., Bonjawo, R.M.E., Duscher, A.A., Foster, J.S. 2018. A year in the life of a thrombolite: comparative metatranscriptomics reveals dynamic metabolic changes over diel and seasonal cycles. *Environmental Microbiology*, 20(2):842-861.
- Proemse, B.C., Eberhard, R.S., Sharples, C., Bowman, J.P., Richards, K., et al. 2017. Stromatolites on the rise in peat-bound karstic wetlands. *Scientific Reports*, 7(1):15507.
- Chiriac, C.M., Szekeres, E., Rudi, K., Baricz, A., Hegedus, A., et al. 2017. Differences in temperature and water chemistry shape distinct diversity patterns in thermophilic microbial communities. *Applied and Environmental Microbiology*, 83(21): e01363-17.
- Mahajan, G.B., Balachandran, L. 2017. Sources of antibiotics: Hot springs. *Biochemical Pharmacology*, 134:35-41.
- Rempfert, K.R., Miller, H.M., Bompard, N., Nothaft, D., Matter, J.M., et al. 2017. Geological and geochemical controls on subsurface microbial life in the Samail Ophiolite, Oman. *Frontiers in Microbiology*, 8:56.
- Greer, H.F., Zhou, W., Guo, L. 2017. Reversed crystal growth of calcite in naturally occurring travertine crust. *Crystals*, 7(2):36.
- Rojas, C., Gutierrez, R.M., Bruns, M.A. 2016. Bacterial and eukaryal diversity in soils forming from acid mine drainage precipitates under reclaimed vegetation and biological crusts. *Applied Soil Ecology*, 105:57-66.
- Li, P., Lin, W., Liu, X., Wang, X., Luo, L. 2016. Environmental factors affecting microbiota dynamics during traditional solid-state fermentation of chinese daqu starter. *Frontiers in Microbiology*, 7:1237.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 1,39) + 14] = 27,73$ puncte

10. Coman, C., Drugă, B., Hegedus, A., Sicora, C., Dragoș, N., 2013. Archaeal and bacterial diversity in two hot spring microbial mats from a geothermal region in Romania. *Extremophiles*, 17:523-534. (AI=0,7)

Citări:

- Mechri S., Bouacem K., Zaráf Jaouadi N., Rekik H., Ben Elhoul M., et al. 2019. Identification of a novel protease from the thermophilic *Anoxybacillus kamchatkensis* M1V and its application as laundry detergent additive. *Extremophiles*, 23(6):687-706.
- Joseph E.P., Frey H.M., Manon M.R., Onyeali M.-M.C., DeFranco K., et al. 2019. Update on the fluid geochemistry monitoring time series for geothermal systems in Dominica, Lesser

- Antilles island arc: 2009–2017. *Journal of Volcanology and Geothermal Research*, 376(86):103.
- Zhang L., Cai Y., Jiang M., Dai J., Guo X., et al. 2019. The levels of microbial diversity in different water layers of saline Chagan Lake, China. *Journal of Oceanology and Limnology*, DOI: 10.1007/s00343-019-9027-7.
- Rotaru A.-E., Posth N.R., Löscher C.R., Miracle M.R., Vicente E., et al. 2019. Interspecies interactions mediated by conductive minerals in the sediments of the iron rich meromictic Lake La Cruz, Spain. *Limnetica*, 38(1):21-40.
- Patwardhan, S., Foustoukos, D.I., Giovannelli, D., Yücel, M., Vetriani, C. 2019. Ecological succession of sulfur-oxidizing epsilon- And gammaproteobacteria during colonization of a shallow-water gas vent. *Frontiers in Microbiology*, 9:2970.
- Prieto-Barajas, C.M., Alcaraz, L.D., Valencia-Cantero, E., Santoyo, G. 2018. Life in Hot Spring Microbial Mats Located in the Trans-Mexican Volcanic Belt: A 16S/18S rRNA Gene and Metagenomic Analysis. *Geomicrobiology Journal*, 35(8):704-712.
- Ming, D.-S., Chen, Q.-Q., Chen, X.-T. 2018. Analysis of resistance genes of clinical *Pannonibacter phragmitetus* strain 31801 by complete genome sequencing. *Archives of Microbiology*, 200(7):1101-1109.
- Paduano, S., Valeriani, F., Romano-Spica, V., Bargellini, A., Borella, P., Marchesi, I. 2018. Microbial biodiversity of thermal water and mud in an Italian spa by metagenomics: A pilot study. *Water Science and Technology: Water Supply*, 18(4):1456-1465.
- Selvarajan, R., Sibanda, T., Tekere, M. 2018. Thermophilic bacterial communities inhabiting the microbial mats of “indifferent” and chalybeate (iron-rich) thermal springs: Diversity and biotechnological analysis. *MicrobiologyOpen*, 7(2):e00560.
- Lavrentyeva, E.V., Radnagurueva, A.A., Barkhutova, D.D., Belkova, N.L., Zaitseva, S.V., et al. 2018. Bacterial Diversity and Functional Activity of Microbial Communities in Hot Springs of the Baikal Rift Zone. *Microbiology (Russian Federation)*, 87(2), 272-281.
- Subudhi, E., Sahoo, R.K., Gaur, M., Singh, A., Das, A. 2018. Shift in Cyanobacteria Community Diversity in Hot Springs of India. *Geomicrobiology Journal*, 35(2):141-147.
- Prieto-Barajas, C.M., Valencia-Cantero, E., Santoyo, G. 2018. Microbial mat ecosystems: Structure types, functional diversity, and biotechnological application. *Electronic Journal of Biotechnology*, 31:48-56.
- Rozanov, A.S., Bryanskaya, A.V., Ivanisenko, T.V., Malup, T.K., Peltek, S.E. 2017. Biodiversity of the microbial mat of the Garga hot spring. *BMC Evolutionary Biology*, 17:254.
- Wang M., Zhang X., Jiang T., Hu S., Yi Z., et al. 2017. Liver abscess caused by *Pannonibacter phragmitetus*: Case report and literature review. *Frontiers in Medicine*, 4:48.
- Kalita, D., Joshi, S.R. 2017. Study on bioremediation of Lead by exopolysaccharide producing metallophilic bacterium isolated from extreme habitat. *Biotechnology Reports*, 16:48-57.

- Zhou, Y., Jiang, T., Hu, S., Wang, M., Ming, D., Chen, S. 2017. Genomic insights of *Pannonibacter phragmitetus* strain 31801 isolated from a patient with a liver abscess. *MicrobiologyOpen*, 6(6):e00515.
- Schuler, C.G., Havig, J.R., Hamilton, T.L. 2017. Hot spring microbial community composition, morphology, and carbon fixation: Implications for interpreting the ancient rock record. *Frontiers in Earth Science*, 5:97.
- Chiriac, C.M., Szekeres, E., Rudi, K., Baricz, A., Hegedus, A., et al. 2017. Differences in temperature and water chemistry shape distinct diversity patterns in thermophilic microbial communities. *Applied and Environmental Microbiology*, 83(21): e01363-17.
- DeCastro, M.-E., Escuder, J.-J., Becerra, M., Rodríguez-Belmonte, E., González, M.-I. 2017. Archaeal biocommunication in Hot Springs revealed by metagenomics. *Biocommunication of Archaea*, pp. 85-101.
- Albers, S.-V., Quax, T.E.F. 2017. Archaeal surface structures and their role in communication with the extracellular environment. *Biocommunication of Archaea*, pp. 67-84.
- Ghilamical, A.M., Budambula, N.L.M., Anami, S.E., Mehari, T., Boga, H.I. 2017. Evaluation of prokaryotic diversity of five hot springs in Eritrea. *BMC Microbiology*, 17(1):203.
- Hayashida, G., Schneider, C., Espíndola, L., Arias, D., Riquelme, C., et al. 2017. Characterization of a Chlorophyta microalga isolated from a microbial mat in Salar de Atacama (northern Chile) as a potential source of compounds for biotechnological applications. *Phycological Research*, 65(3):202-211.
- Gupta, P., Manjula, A., Rajendhran, J., Gunasekaran, P., Vakhlu, J. 2017. Comparison of Metagenomic DNA Extraction Methods for Soil Sediments of High Elevation Puga Hot Spring in Ladakh, India to Explore Bacterial Diversity. *Geomicrobiology Journal*, 34(4):289-299.
- Saxena, R., Dhakan, D.B., Mittal, P., Waiker, P., Chowdhury, A., et al. 2017. Metagenomic analysis of hot springs in central India reveals hydrocarbon degrading thermophiles and pathways essential for survival in extreme environments. *Frontiers in Microbiology*, 7:2123.
- Sciuto, K., Moro, I. 2016. Detection of the new cosmopolitan genus *Thermoleptolyngbya* (Cyanobacteria, Leptolyngbyaceae) using the 16S rRNA gene and 16S–23S ITS region. *Molecular Phylogenetics and Evolution*, 105:15-35.
- Panda, A.K., Bisht, S.S., De Mandal, S., Kumar, N.S. Bacterial and archaeal community composition in hot springs from Indo-Burma region, North-east India. 2016. *AMB Express*, 6 (1):111.
- Aubé, J., Senin, P., Pringault, O., Bonin, P., Deflandre, B., et al. 2016. The impact of long-term hydrocarbon exposure on the structure, activity, and biogeochemical functioning of microbial mats. *Marine Pollution Bulletin*, 111(1-2):115-125.
- Kraková, L., Šoltys, K., Budiš, J., Grivalský, T., Ďuriš, F., et al. 2016. Investigation of bacterial and archaeal communities: novel protocols using modern sequencing by Illumina MiSeq and traditional DGGE-cloning. *Extremophiles*, 20(5):795-808.

- Drewniak, L., Krawczyk, P.S., Mielnicki, S., Adamska, D., Sobczak, A., et al. 2016. Physiological and metagenomic analyses of microbial mats involved in self-purification of mine waters contaminated with heavy metals. *Frontiers in Microbiology*, 7:1252.
- Valeriani, F., Biagini, T., Giampaoli, S., Crognale, S., Santoni, D., Romano Spica, V. 2016. Draft genome sequence of *Tepidimonas taiwanensis* strain VT154-175. *Genome Announcements*, 4(5):e00942.
- Yilmaz Cankiliç, M. 2016. Determination of cyanobacterial composition of Eynal (Sımav) hot spring in Kütahya, Turkey. *Applied Ecology and Environmental Research*, 14(4):607-622.
- Alcamán, M.E., Fernandez, C., Delgado, A., Bergman, B., Díez, B. 2015. The cyanobacterium *Mastigocladus fulfills* the nitrogen demand of a terrestrial hot spring microbial mat. *ISME Journal*, 9(10):2290-2303.
- Urbietta, M.S., González-Toril, E., Bazán, Á.A., Giaveno, M.A., Donati, E. 2015. Comparison of the microbial communities of hot springs waters and the microbial biofilms in the acidic geothermal area of Copahue (Neuquén, Argentina). *Extremophiles*, 19(2):437-450.
- Coman, C., Chiriac, C.M., Robeson, M.S., Ionescu, C., Dragos, N., et al. 2015. Structure, mineralogy, and microbial diversity of geothermal spring microbialites associated with a deep oil drilling in Romania. *Frontiers in Microbiology*, 6:253.
- Pohlschroder, M., Esquivel, R.N. 2015. Archaeal type IV pili and their involvement in biofilm formation. *Frontiers in Microbiology*, 6:190.
- Amarouche-Yala, S., Benouadah, A., El Ouahab Bentabet, A., López-García, P. 2015. Morphological and phylogenetic diversity of thermophilic cyanobacteria in Algerian hot springs. *Extremophiles*, 18(6):1035-1047.
- Pujalte, M.J., Lucena, T., Ruvira, M.A., Arahall, D.R., Macián, M.C. 2014. The family Rhodobacteraceae. *The Prokaryotes: Alphaproteobacteria and Betaproteobacteria*, pp. 439-512.
- Oren, A. 2014. The family Methanosarcinaceae. *The Prokaryotes: Other Major Lineages of Bacteria and The Archaea*, pp. 259-281.
- Farías, M.E., Contreras, M., Rasuk, M.C., Kurth, D., Flores, M.R., et al. 2014. Characterization of bacterial diversity associated with microbial mats, gypsum evaporites and carbonate microbialites in thalassic wetlands: Tebenquiche and La Brava, Salar de Atacama, Chile. *Extremophiles*, 18(2):311-329.
- Németh, A., Szirányi, B., Krett, G., Janurik, E., Kosáros, T., et al. 2014. Prokaryotic phylogenetic diversity of Hungarian deep subsurface geothermal well waters. *Acta Microbiologica et Immunologica Hungarica*, 61(3):363-377.
- Meyer-Dombard, D.R., Amend, J.P. 2014. Geochemistry and microbial ecology in alkaline hot springs of Ambitle Island, Papua New Guinea. *Extremophiles*, 18(4):763-778.
- Deep, K., Poddar, A., Das, S.K. 2013. *Anoxybacillus suryakundensis* sp. nov., a moderately thermophilic, alkalitolerant bacterium isolated from hot spring at Jharkhand, India. *PLoS ONE*, 8(12):e85493.

Cha, I.-T., Min, U.-G., Kim, S.-J., Yim, K.J., Roh, S.W., Rhee, S.-K. 2013. *Methanomethylovorans uponensis* sp. nov., a methylotrophic methanogen isolated from wetland sediment. *Antonie van Leeuwenhoek, International Journal of General and Molecular Microbiology*, 104(6):1005-1012.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 0,7) + 42] = 50,9$ puncte

11. Coman, C., Bica, A., Drugă, B., Barbu-Tudoran, L., Dragoș N., 2011. Methodological constraints in the molecular biodiversity study of a thermomineral spring cyanobacterial mat: a case study. *Antonie van Leeuwenhoek*, 99:271-281. (AIS=0,6)

Citări:

Tu J., Chen L., Gao S., Zhang J., Bi C., et al. 2019. Obtaining genome sequences of mutualistic bacteria in single *Microcystis* colonies. *International Journal of Molecular Sciences*, 20(20):5047.

Drugă, B., Ukrainczyk, N., Weise, K., Koenders, E., Lackner, S. 2018. Interaction between wastewater microorganisms and geopolymer or cementitious materials: Biofilm characterization and deterioration characteristics of mortars. *International Biodeterioration and Biodegradation*, 134:58-67.

Costa, N.B., Kolman, M.A., Giani, A., Moisaner, P. 2016. Cyanobacteria diversity in alkaline saline lakes in the Brazilian Pantanal wetland: A polyphasic approach. *Journal of Plankton Research*, 38(6):1389-1403.

El Khalloufi, F., Oufdou, K., Bertrand, M., Lahrouni, M., Oudra, B., et al. 2016. Microbiote shift in the *Medicago sativa* rhizosphere in response to cyanotoxins extract exposure. *Science of the Total Environment*, 539:135-142.

Coman, C., Chiriac, C.M., Robeson, M.S., Ionescu, C., Dragoș, N., et al. 2015. Structure, mineralogy, and microbial diversity of geothermal spring microbialites associated with a deep oil drilling in Romania. *Frontiers in Microbiology*, 6:253.

Cai, H., Jiang, H., Krumholz, L.R., Yang, Z. 2014. Bacterial community composition of size-fractionated aggregates within the phycosphere of cyanobacterial blooms in a eutrophic freshwater lake. *PLoS ONE*, 9(8):e102879.

Coman, C., Drugă, B., Hegedus, A., Sicora, C., Dragoș, N. 2013. Archaeal and bacterial diversity in two hot spring microbial mats from a geothermal region in Romania. *Extremophiles*, 17(3):523-534.

Shi, L., Cai, Y., Kong, F., Yu, Y. 2012. Specific association between bacteria and buoyant *Microcystis* colonies compared with other bulk bacterial communities in the eutrophic Lake Taihu, China. *Environmental Microbiology Reports*, 4(6):669-678.

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $1 \times [4 + (7 \times 0,6) + 6] = 14,2$ puncte

Articole în reviste cotate ISI, ca și contributor:

1. Baricz, A., Chiriac, C.M., Andrei, A.-Ș., Bulzu, P.-A., Levei, E.A., Cadar, O., Battes, K.P., Cîmpean, M. Șenilă, M., Cristea, A., Muntean, V., Alexe, M., Coman, C., Szekeres, E.K., Sicora, C.I., Ionescu, A., Blain, D., O'Neill, W.K., Edwards, J., Hallsworth, J.E., Banciu, H.L. 2019. Spatio-temporal insights into microbiology of the freshwater-to-hypersaline, oxic-hypoxic-euxinic waters of Ursu Lake. DOI: 10.1111/1462-2920.14909. (AIS=2,1)

Citări: 0

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $0,7x[4+(7x2,1)]=13,09$ puncte

2. Szöke-Nagy, T., Porav, A.S., Coman, C., Cozar, B.I., Dina, N.E., Tripon, C. 2019. Characterization of the Action of Antibiotics and Essential Oils against Bacteria by Surface-Enhanced Raman Spectroscopy and Scanning Electron Microscopy. *Analytical Letters*, 52(1):190-200. (AIS=0.2)

Citări: 0

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $0,7x[4+(7x0,2)]=3,78$ puncte

3. Dragoș, N., Chiriac, C., Porav, S., Szöke-Nagy, T., Coman, C., Török, L., Hegedűs, A. 2019. *Desmodesmus tropicus* (Chlorophyta) in the Danube Delta—reassessing the phylogeny of the series *Maximi*. *European Journal of Phycology*, 54(3):300-314.

Citări: 0

Punctaj articol conform formulei (1) din ordinul 6.129/2016: $0,7x[4+(7x0,7)]=6,23$ puncte

4. Andrei, A.-Ș., Baricz, A., Robeson, M.S., II, Păușan, M.R., Tămaș, T., Chiriac, C., Szekeres, E., Barbu-Tudoran, L., Levei, E.A., Coman, C., Podar, M., Banciu, H.L. 2017. Hypersaline sapropels act as hotspots for microbial dark matter. *Scientific Reports*, 7(1): 6150. (AIS=1,35).

Citări:

Li J., Hu R., Guo Y., Chen S., Xie X., et al. 2019. Bioturbation of peanut worms *Sipunculus nudus* on the composition of prokaryotic communities in a tidal flat as revealed by 16S rRNA gene sequences. *MicrobiologyOpen*, 8(8):e00802.

Baricz A., Chiriac C.M., Andrei A.-Ș., Bulzu P.-A., Levei E.A., et al. 2019. Spatio-temporal insights into microbiology of the freshwater-to-hypersaline, oxic-hypoxic-euxinic waters of Ursu Lake. *Environmental Microbiology*, DOI: 10.1111/1462-2920.14909.

Alexe, M., Șerban, G., Baricz, A., Andrei, A.-Ș., Cristea, A., et al. 2018. Limnology and plankton diversity of salt lakes from Transylvanian Basin (Romania): A review. *Journal of Limnology*, 77(1): 17-34.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7x[4+(7x1,35)+3]=11,51$ puncte

5. Dina, N.E., Zhou, H., Colniță, A., Leopold, N., Szoke-Nagy, T., Coman, C., Haisch, C. 2017. Rapid single-cell detection and identification of pathogens by using surface-enhanced Raman spectroscopy. *Analyst*, 142(10): 1782-1789. (AIS=0,9)

Citări:

- Jabbar A.A., Alwan A.M., Zayer M.Q., Bohan A.J. 2020. Efficient single cell monitoring of pathogenic bacteria using bimetallic nanostructures embedded in gradient porous silicon. *Materials Chemistry and Physics*, 241:122359.
- Akanny E., Bonhommé A., Commun C., Doleans-Jordheim A., Farre C., et al. 2020. Surface-enhanced Raman spectroscopy using uncoated gold nanoparticles for bacteria discrimination. *Journal of Raman Spectroscopy*, DOI: 10.1002/jrs.5827.
- Zhang K., Hao C., Huo Y., Man B., Zhang C., et al. 2019. Label-free diagnosis of lung cancer with tissue-slice surface-enhanced Raman spectroscopy and statistical analysis. *Lasers in Medical Science*, 34(9):1849-1855.
- Rani A., Donovan N., Mantri N. 2019. Review: The future of plant pathogen diagnostics in a nursery production system. *Biosensors and Bioelectronics*, 145:111631.
- Ravindran V.B., Soni S.K., Ball A.S.A. 2019. Review on the current knowledge and prospects for the development of improved detection methods for soil-transmitted helminth ova for the safe reuse of wastewater and mitigation of public health risks. *Water (Switzerland)*, 11(6):1212.
- Hamm L., Gee A., Indrasekara A.S.S. 2019. Recent advancement in the surface-enhanced raman spectroscopy-based biosensors for infectious disease diagnosis. *Applied Sciences (Switzerland)*, 9(7):1448.
- Gong H., Zhang K., Dicko C., Bülow L., Ye L. 2019. Ag-Polymer Nanocomposites for Capture, Detection, and Destruction of Bacteria. *ACS Applied Nano Materials*, 2(3):1655-1663.
- Szöke-Nagy T., Porav A.S., Coman C., Cozar B.I., Dina N.E., Tripon C. 2019. Characterization of the Action of Antibiotics and Essential Oils against Bacteria by Surface-Enhanced Raman Spectroscopy and Scanning Electron Microscopy. *Analytical Letters*, 52(1):190-200.
- Andryukov B.G., Karpenko A.A., Matosova E.V., Lyapun I.N. 2019. Raman spectroscopy as a modern diagnostic technology for study and indication of infectious agents (Review). *Sovremennye Tehnologii v Medicine*, 11(4):161-173.
- Langer J., Jimenez De Aberasturi D., Aizpurua J., Alvarez-Puebla R.A., Auguie B., et al. 2019. Present and Future of Surface-Enhanced Raman Scattering. *ACS Nano*, DOI: 10.1021/acsnano.9b04224.
- Ankudze B., Asare B., Goffart S., Koistinen A., Nuutinen T., et al. 2019. Hydraulically pressed silver nanowire-cotton fibers as an active platform for filtering and surface-enhanced Raman scattering detection of bacteria from fluid. *Applied Surface Science*, 479:663-668.
- Huang C.-C., Hung Y.-S., Weng Y.-M., Chen W., Lai Y.-S. 2019. Sustainable development of carbon nanodots technology: Natural products as a carbon source and applications to food safety. *Trends in Food Science and Technology*, 86: 144-152.

- Yin L., Zhang Z., Liu Y., Gao Y., Gu J. 2019. Recent advances in single-cell analysis by mass spectrometry. *Analyst*, 144: 824-845.
- Chen X., Tang M., Liu Y., Huang J., Liu Z., et al. 2019. Surface-enhanced Raman scattering method for the identification of methicillin-resistant *Staphylococcus aureus* using positively charged silver nanoparticles. *Microchimica Acta*, 186: 102.
- Abdullah A., Dastider S.G., Jasim I., Shen Z., Yuksek N., et al. 2019. Microfluidic based impedance biosensor for pathogens detection in food products. *Electrophoresis*, 40: 508-520.
- Da Silva Neves M.M.P., Martín-Yerga D. 2019. Advanced nanoscale approaches to single-(bio)entity sensing and imaging. *Biosensors*, 8:E100.
- Nasseri, B., Soleimani, N., Rabiee, N., Kalbasi, A., Karimi, M., Hamblin, M.R. 2018. Point-of-care microfluidic devices for pathogen detection. *Biosensors and Bioelectronics*, 117:112-128.
- Wang, K., Li, S., Petersen, M., Wang, S., Lu, X. 2018. Detection and characterization of antibiotic-resistant bacteria using surface-enhanced raman spectroscopy. *Nanomaterials*, 8(10):762.
- Khansili, N., Rattu, G., Krishna, P.M. 2018. Label-free optical biosensors for food and biological sensor applications. *Sensors and Actuators, B: Chemical*, 265:35-49.
- Galvan, D.D., Yu, Q. Surface-Enhanced Raman Scattering for Rapid Detection and Characterization of Antibiotic-Resistant Bacteria. *Advanced Healthcare Materials*, 7(13):1701335.
- Pearson, B., Mills, A., Tucker, M., Gao, S., McLandsborough, L., He, L. 2018. Rationalizing and advancing the 3-MPBA SERS sandwich assay for rapid detection of bacteria in environmental and food matrices. *Food Microbiology*, 72:89-97.
- Colino, C.I., Millán, C.G., Lanao, J.M. Nanoparticles for signaling in biodiagnosis and treatment of infectious diseases. *International Journal of Molecular Sciences*, 19(6):1627.
- Zheng, X.-S., Jahn, I.J., Weber, K., Cialla-May, D., Popp, J. Label-free SERS in biological and biomedical applications: Recent progress, current challenges and opportunities. *Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy*, 197:56-77.
- Zhu, K., Hong, Z., Kang, S.-Z., Qin, L., Li, G., Li, X. 2018. Assembly of potassium niobate nanosheets/silver oxide composite films with good SERS performance towards crystal violet detection. *Journal of Physics and Chemistry of Solids*, 115: 69-74.
- Dina, N.E., Gherman, A.M.R., Chiş, V., Sârbu, C., Wieser, A., Bauer, D., Haisch, C. 2018. Characterization of Clinically Relevant Fungi via SERS Fingerprinting Assisted by Novel Chemometric Models. *Analytical Chemistry*, 90(4):2484-2492.
- Mosier-Boss, P.A. 2017. Review on SERS of bacteria. *Biosensors*, 7(4):51.
- Dina, N.E., Colniţă, A., Szöke-Nagy, T., Porav, A.S. 2017. A Critical Review on Ultrasensitive, Spectroscopic-based Methods for High-throughput Monitoring of Bacteria during Infection Treatment. *Critical Reviews in Analytical Chemistry*, 47(6):499-512.

Colniță, A., Dina, N.E., Leopold, N., Vodnar, D.C., Bogdan, D., Porav, S.A., David, L. 2017. Characterization and discrimination of gram-positive bacteria using raman spectroscopy with the aid of principal component analysis. *Nanomaterials*, 7(9):248.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 0,9) + 26] = 25,41$ puncte

6. Hegedűs, A., Mocan, A., Barbu-Tudoran, L., Coman, C., Dragoș, N. 2016. Molecular phylogeny of *Botryococcus braunii* strains (race A) – An integrative approach. *Algal Research*, 19(1):189-197. (AIS=1,3)

Citări:

Ali, I., Naqvi, S.R., Bahadar, A. 2018. Kinetic analysis of *Botryococcus braunii* pyrolysis using model-free and model fitting methods. (2018) *Fuel*, 214, pp. 369-380.

Nakamura, H., Shiozaki, T., Gonda, N., Furuya, K., Matsunaga, S., Okada, S. 2017. Utilization of ammonium by the hydrocarbon-producing microalga, *Botryococcus braunii* Showa. *Algal Research*, 25:445-451.

Gouveia, J.D., Ruiz, J., van den Broek, L.A.M., Hesselink, T., Peters, S., et al. 2017. *Botryococcus braunii* strains compared for biomass productivity, hydrocarbon and carbohydrate content. *Journal of Biotechnology*, 248:77-86.

Nakamura, H., Matsunaga, S., Kawagishi, H., Okada, S. 2016. Effects of 2-azahypoxanthine on extracellular terpene accumulations by the green microalga *Botryococcus braunii*, race B. *Algal Research*, 20:267-275.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 1,3) + 4] = 11,97$ puncte

7. Farkas, A., Crăciunaș, C., Chiriac, C., Szekeres, E., Coman, C., Butiuc-Keul, A., 2016. Exploring the role of coliform bacteria in class 1 integron carriage and biofilm formation during drinking water treatment. *Microbial Ecology*. DOI: 10.1007/s00248-016-0758-0. (AIS=1,0)

Citări:

Siedlecka A., Piekarska K. 2019. Antibiotic resistance in tap water during the summer season - Preliminary research. *E3S Web of Conferences*, 116:77.

Forbes S., Morgan N., Humphreys G.J., Amézquita A., Mistry H., McBain A.J. 2019. Loss of function in *Escherichia coli* exposed to environmentally relevant concentrations of benzalkonium chloride. *Applied and Environmental Microbiology*, 85(4):e02417-18.

Li, A., Chen, L., Zhang, Y., Tao, Y., Xie, H., et al. 2018. Occurrence and distribution of antibiotic resistance genes in the sediments of drinking water sources, urban rivers, and coastal areas in Zhuhai, China. *Environmental Science and Pollution Research*, 25:26209-26217.

Chen, X.P., Ali, L., Wu, L.-Y., Liu, C., Gang, et al. 2018. Biofilm formation plays a role in the formation of multidrug-resistant *Escherichia coli* toward nutrients in microcosm experiments. *Frontiers in Microbiology*, 9:367.

- Jiang, X., Yu, T., Liu, L., Li, Y., Zhang, K., et al. 2017. Examination of quaternary ammonium compound resistance in *Proteus mirabilis* isolated from cooked meat products in China. *Frontiers in Microbiology*, 8:2417.
- Jiang, X., Xu, Y., Li, Y., Zhang, K., Liu, L., et al. 2017. Characterization and horizontal transfer of qacH-associated class 1 integrons in *Escherichia coli* isolated from retail meats. *International Journal of Food Microbiology*, 258:12-17.
- Palacios, O.A., Zavala-Díaz de la Serna, F.J., de Lourdes Ballinas-Casarrubias, M., Espino-Valdés, M.S., Nevárez-Moorillón, G.V. 2017. Microbiological impact of the use of reclaimed wastewater in recreational parks. *International Journal of Environmental Research and Public Health*, 14(9):1009.
- Gillings, M.R. 2017. Lateral gene transfer, bacterial genome evolution, and the Anthropocene. *Annals of the New York Academy of Sciences*, 1389(1):20-36.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 1) + 8] = 13,3$ puncte

8. Andrei A.S., Robeson, M., Baricz, A., Coman, C., Muntean, V., Ionescu, A., Etiope, G., Alexe, M., Sicora, C., Podar, M., Banciu, H.L., 2015. Contrasting taxonomic and physiological stratification of microbial communities from two hypersaline meromictic lakes. *The ISME Journal*, DOI:10.1038/ismej.2015.60. (AI=3,3)

Citări:

- Čanković M., Žučko J., Radić I.D., Janeković I., Petrić I., et al. 2019. Microbial diversity and long-term geochemical trends in the euxinic zone of a marine, meromictic lake. *Systematic and Applied Microbiology*, 42(6):126016.
- Martin-Cuadrado A.-B., Senel E., Martínez-García M., Cifuentes A., Santos F., et al. 2019. Prokaryotic and viral community of the sulfate-rich crust from Peñahueca ephemeral lake, an astrobiology analogue. *Environmental Microbiology*, 21(10):3577-3600.
- Raffa C., Rizzo C., Strous M., De Domenico E., Sanfilippo M., et al. 2019. Prokaryotic dynamics in the meromictic coastal Lake Faro (Sicily, Italy). *Diversity*, 11(3):37.
- Miller E.R., Kearns P.J., Niccum B.A., Schwartz J.O', Ornstein A., Wolfe B.E. 2019. Establishment limitation constrains the abundance of lactic acid bacteria in the Napa cabbage phyllosphere. *Applied and Environmental Microbiology*, 85(13):e00269-19.
- Baricz A., Chiriac C.M., Andrei A.-Ș., Bulzu P.-A., Levei E.A., et al. 2019. Spatio-temporal insights into microbiology of the freshwater-to-hypersaline, oxic-hypoxic-euxinic waters of Ursu Lake. *Environmental Microbiology*, DOI: 10.1111/1462-2920.14909.
- Li W., Morgan-Kiss R.M. 2019. Influence of environmental drivers and potential interactions on the distribution of microbial communities from three permanently stratified Antarctic lakes. *Frontiers in Microbiology*, 10:1067.

- Danza F., Ravasi D., Storelli N., Roman S., Lüdin S., et al. 2019. Bacterial diversity in the water column of meromictic Lake Cadagno and evidence for seasonal dynamics. *PLoS ONE*, 13: e0209743.
- Matyugina E., Belkova N., Borzenko S., Lukyanov P., Kabilov M., et al. 2019. Structure and diversity dynamics of microbial communities at day and night: investigation of meromictic Lake Doroninskoe, Transbaikalia, Russia. *Journal of Oceanology and Limnology*, 36:1978-1992.
- Cristea, A., Baricz, A., Leopold, N., Floare, C.G., Borodi, G., et al. 2018. Polyhydroxybutyrate production by an extremely halotolerant *Halomonas elongata* strain isolated from the hypersaline meromictic Fără Fund Lake (Transylvanian Basin, Romania). *Journal of Applied Microbiology*, 125(5):1343-1357.
- Savvichev, A.S., Babenko, V.V., Lunina, O.N., Letarova, M.A., Boldyreva, D.I., et al. 2018. Sharp water column stratification with an extremely dense microbial population in a small meromictic lake, Trekhtzvetnoe. *Environmental Microbiology*, 20: 3784-3797.
- Wu, Y.-T., Yang, C.-Y., Chiang, P.-W., Tseng, C.-H., Chiu, H.-H., et al. 2018. Comprehensive insights into composition, metabolic potentials, and interactions among archaeal, bacterial, and viral assemblages in meromictic Lake Shunet in Siberia. *Frontiers in Microbiology*, 9:1763.
- Kalwasińska, A., Deja-Sikora, E., Burkowska-But, A., Szabó, A., Felföldi, T., et al. 2018. Changes in bacterial and archaeal communities during the concentration of brine at the graduation towers in Ciechocinek spa (Poland). *Extremophiles*, 22(2):233-246.
- Mora-Ruiz, M.D.R., Cifuentes, A., Font-Verdera, F., Pérez-Fernández, C., Farias, M.E., et al. 2018. Biogeographical patterns of bacterial and archaeal communities from distant hypersaline environments. *Systematic and Applied Microbiology*, 41:139-150.
- Liu, K., Ding, X., Tang, X., Wang, J., Li, W., et al. 2018. Macro and microelements drive diversity and composition of prokaryotic and fungal communities in hypersaline sediments and saline-alkaline soils. *Frontiers in Microbiology*, 9:352.
- Knelman, J.E., Graham, E.B., Prevéy, J.S., Robeson, M.S., Kelly, P., et al. 2018. Interspecific plant interactions reflected in soil bacterial community structure and nitrogen cycling in primary succession. *Frontiers in Microbiology*, 9:128.
- Alexe, M., Şerban, G., Baricz, A., Andrei, A.-Ş., Cristea, A., et al. 2018. Limnology and plankton diversity of salt lakes from Transylvanian Basin (Romania): A review. *Journal of Limnology*, 77(1):17-34.
- Lee, C.J.D., McMullan, P.E., O'Kane, C.J., Stevenson, A., Santos, I.C., et al. 2018. NaCl-saturated brines are thermodynamically moderate, rather than extreme, microbial habitats. *FEMS Microbiology Reviews*, 42(5):672-693.
- Andrei, A.-Ş., Baricz, A., Robeson, M.S., II, Păușan, M.R., Tămaș, T., et al. 2017. Hypersaline sapropels act as hotspots for microbial dark matter. *Scientific Reports*, 7(1):6150.
- Abedon, S.T. 2017. Why archaea are limited in their exploitation of other, living organisms. *Biocommunication of Archaea*, pp. 41-66. DOI: 10.1007/978-3-319-65536-9_4.

- Johansen, R.B., Johnston, P., Mieczkowski, P., Perry, G.L.W., Robeson, M.S., et al. 2017. Scattered far and wide: A broadly distributed temperate dune grass finds familiar fungal root associates in its invasive range. *Soil Biology and Biochemistry*, 112:177-190.
- Huggett, M.J., Kavazos, C.R.J., Bernasconi, R., Czarnik, R., Horwitz, P. 2017. Bacterioplankton assemblages in coastal ponds reflect the influence of hydrology and geomorphological setting. *FEMS Microbiology Ecology*, 93(6):fix067.
- Andrei, A.-Ș., Baricz, A., Păușan, M., Muntean, V., Sicora, C.I., et al. 2017. Spatial Distribution and Molecular Diversity of Archaeal Communities in the Extreme Hypersaline Meromictic Brâncoveanu Lake (Transylvanian Basin, Romania). *Geomicrobiology Journal*, 34(2):130-138.
- Mark Ibekwe, A., Ors, S., Ferreira, J.F.S., Liu, X., Suarez, D.L. 2017. Seasonal induced changes in spinach rhizosphere microbial community structure with varying salinity and drought. *Science of the Total Environment*, 579:1485-1495.
- Hengy, M.H., Horton, D.J., Uzarski, D.G., Learman, D.R. 2017. Microbial community diversity patterns are related to physical and chemical differences among temperate lakes near Beaver Island, MI. *PeerJ*, 2017(10): e3937.
- Johansen, R.B., Johnston, P., Mieczkowski, P., Perry, G.L.W., Robeson, M.S., et al. 2016. A native and an invasive dune grass share similar, patchily distributed, root-associated fungal communities. *Fungal Ecology*, 23:141-155.
- Baatar, B., Chiang, P.-W., Rogozin, D.Y., Wu, Y.-T., Tseng, C.-H., et al. 2016. Bacterial communities of three saline meromictic lakes in Central Asia. *PLoS ONE*, 11(3):e0150847.
- Utturkar, S.M., Cude, W.N., Robeson, M.S., Jr., Yang, Z.K., Klingeman, D.M., et al. 2016. Enrichment of root endophytic bacteria from *populus deltoides* and single-cell-genomics analysis. *Applied and Environmental Microbiology*, 82(18):5698-5708.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 3,3) + 25] = 36,47$ puncte

9. Baricz, A., Coman, C., Andrei, A.S., Muntean, V., Keresztes, Z.G., Păușan, M., Alexe, M., Banciu, H.L., 2014. Spatial and temporal distribution of archaeal diversity in meromictic, hypersaline Ocnei Lake (Transylvanian Basin, Romania). *Extremophiles*, 18:399-413. (AI=0,7)

Citări:

- Sayed A.M., Hassan M.H.A., Alhadrami H.A., Hassan H.M., Goodfellow M., Rateb M.E. 2020. Extreme environments: microbiology leading to specialized metabolites. *Journal of Applied Microbiology*, 128(3): 630-657.
- Pandya R.D., Singh S.P. 2020. Pigment production by an extreme halophilic archaea on *Halorubrum* sp. J4.2.2 from little Rann of Kutch, Gujarat, India. *Research Journal of Biotechnology*, 15(1): 88-100.

- Couto-Rodríguez R.L., Montalvo-Rodríguez R. 2019. Temporal analysis of the microbial community from the crystallizer ponds in Cabo Rojo, Puerto Rico, using metagenomics. *Genes*, 10(6):422.
- Baricz A., Chiriac C.M., Andrei A.-Ş., Bulzu P.-A., Levei E.A., et al. 2019. Spatio-temporal insights into microbiology of the freshwater-to-hypersaline, oxic-hypoxic-euxinic waters of Ursu Lake. *Environmental Microbiology*, DOI: 10.1111/1462-2920.14909.
- Vera-Gargallo, B., Ventosa, A. 2018. Metagenomic insights into the phylogenetic and metabolic diversity of the prokaryotic community dwelling in hypersaline soils from the odiel saltmarshes (SW Spain). *Genes*, 9(3):152.
- Alexe, M., Şerban, G., Baricz, A., Andrei, A.-Ş., Cristea, A., et al. 2018. Limnology and plankton diversity of salt lakes from Transylvanian Basin (Romania): A review. *Journal of Limnology*, 77(1):17-34.
- Lee, C.J.D., McMullan, P.E., O'Kane, C.J., Stevenson, A., Santos, I.C., et al. 2018. NaCl-saturated brines are thermodynamically moderate, rather than extreme, microbial habitats. *FEMS Microbiology Reviews*, 42(5):672-693.
- Abedon, S.T. 2017. Why archaea are limited in their exploitation of other, living organisms. *Biocommunication of Archaea*, pp. 41-66.
- Andrei, A.-Ş., Baricz, A., Păuşan, M., Muntean, V., Sicora, C.I., et al. 2017. Spatial Distribution and Molecular Diversity of Archaeal Communities in the Extreme Hypersaline Meromictic Brâncoveanu Lake (Transylvanian Basin, Romania). *Geomicrobiology Journal*, 34(2):130-138.
- Enache, M., Teodosiu, G., Itoh, T., Kamekura, M., Stan-Lotter, H. 2017. Halophilic microorganisms from man-made and natural hypersaline environments: Physiology, ecology, and biotechnological potential. *Adaption of Microbial Life to Environmental Extremes: Novel Research Results and Application*, Second Edition, pp. 201-226.
- Felföldi, T., Ramganes, S., Somogyi, B., Krett, G., Jurecska, L., et al. 2016. Winter Planktonic Microbial Communities in Highland Aquatic Habitats. *Geomicrobiology Journal*, 33(6):494-504.
- Kan, J., Clingenpeel, S., Dow, C.L., McDermott, T.R., Macur, R.E., Inskeep, W.P., Nealson, K.H. 2016. Geochemistry and mixing drive the spatial distribution of free-living archaea and bacteria in yellowstone lake. *Frontiers in Microbiology*, 7:210.
- Andrei, A.-S., Robeson, M.S., Baricz, A., Coman, C., Muntean, V., et al. 2015. Contrasting taxonomic stratification of microbial communities in two hypersaline meromictic lakes. *ISME Journal*, 9(12):2642-2656.
- Oren, A. 2015. Halophilic microbial communities and their environments. *Current Opinion in Biotechnology*, 33:119-124.
- Baricz, A., Cristea, A., Muntean, V., Teodosiu, G., Andrei, A.-Ş., et al. 2015. Culturable diversity of aerobic halophilic archaea (Fam. Halobacteriaceae) from hypersaline, meromictic Transylvanian lakes. *Extremophiles*, 19(2):525-537.

Najjari, A., Elshahed, M.S., Cherif, A., Youssef, N.H. 2015. Patterns and determinants of halophilic archaea (Class halobacteria) diversity in tunisian endorheic salt lakes and sebkhet systems. *Applied and Environmental Microbiology*, 81(13):4432-4441.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 0,7) + 14] = 16,03$ puncte

10. Hegedus, A., Mocan, A., Barbu-Tudoran, L., Coman, C., Drugă, B., Sicora, C., Dragoș, N., 2014. Morphological, biochemical, and phylogenetic assessments of eight *Botryococcus terribilis* strains collected from freshwaters of Transylvania. *Journal of Applied Phycology*, 27(2): 865-878. (AIS=0,6)

Citări:

Dragoș N., Chiriac C., Porav S., Szőke-Nagy T., Coman C., et al. 2019. *Desmodesmus tropicus* (Chlorophyta) in the Danube Delta—reassessing the phylogeny of the series Maximi. *European Journal of Phycology*, 54(3):300-314.

Baudelet, P.-H., Ricochon, G., Linder, M., Muniglia, L. 2017. A new insight into cell walls of Chlorophyta. *Algal Research*, 25:333-371.

Hegedűs, A., Mocan, A., Barbu-Tudoran, L., Coman, C., Dragoș, N. 2016. Molecular phylogeny of *Botryococcus braunii* strains (race A) – An integrative approach. *Algal Research*, 19(1):189-197.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 0,6) + 2] = 7,14$ puncte

11. Drugă, B., Welker, M., Sesărman, A., Hegedus, A., Coman, C., Sicora, C., Dragoș, N., 2013. Molecular characterization of microcystin-producing cyanobacteria from Romanian fresh waters. *European Journal of Phycology*, 48(3):287-294. (AIS=0,5)

Citări:

Drugă B., Buda D.-M., Szekeres E., Chiș C., Chiș I., Sicora C. 2019. The impact of cation concentration on *Microcystis* (cyanobacteria) scum formation. *Scientific Reports*, 9(1):3017.

Johansson E., Legrand C., Björnerås C., Godhe A., Mazur-Marzec H., et al. 2019. High diversity of microcystin chemotypes within a summer bloom of the cyanobacterium *Microcystis botrys*. *Toxins*, 11(12):698.

Xiao, M., Li, M., Reynolds, C.S. 2018. Colony formation in the cyanobacterium *Microcystis*. *Biological Reviews*, 93(3):1399-1420.

Singh, P., Singh, S.S., Aboal, M., Mishra, A.K. 2015. Decoding cyanobacterial phylogeny and molecular evolution using an evonumeric approach. *Protoplasma*, 252(2):519-535.

Punctaj articol conform formulei (2) din ordinul 6.129/2016: $0,7 \times [4 + (7 \times 0,7) + 3] = 8,33$ puncte

Articole în reviste indexate BDI ca autor principal

1. Chiriac, C., Baricz, A., Coman, C. 2018. Draft genome sequence of *Janthinobacterium* sp. strain ROICE36, a putative secondary metabolite-synthesizing bacterium isolated from Antarctic snow. *Genome Announcements*, 6(15): e01553-17.

Citări:

Dieser, M., Smith, H.J., Ramaraj, T., Foreman, C.M. 2019. *Janthinobacterium* CG23_2: Comparative genome analysis reveals enhanced environmental sensing and transcriptional regulation for adaptation to life in an antarctic supraglacial stream. *Microorganisms*, 7(10):454.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 2 puncte

2. Țugui, C.G., Vlădăreanu, I., Baricz, A., Coman, C., 2015. Detection of beta-lactamase resistance genes in a hospital chlorinated wastewater treatment system. *Studia Univ. Babeș-Bolyai, Biologia*, 60(2):33-38

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

3. Szekeres, E., Dragoș, N., Porav, S., Baricz, A., Chiriac, C. M., Szöke-Nagy, T., Coman, C. 2015. Evaluation of bio-resources: Monitoring *Arthrospira* growth in supplemented brackish water. *Studia Univ. Babeș-Bolyai, Biologia*, 60(2):45-48.

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

4. Chiriac, C. M., Barbu-Tudoran, L., Baricz, A., Szekeres, E., Szöke-Nagy, T., Dragoș, N., Coman, C. 2015. Bacterial diversity in a microbial mat colonizing a man-made geothermal spring from Romania, *Studia Univ. Babeș-Bolyai, Biologia*, 60(1): 5-22.

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

5. Coman, C., Bercea, V., Sicora, C. 2012. The study of PS II photochemical activity in *Synechococcus* sp. PCC 7002 under different light intensities. *Annals RSCB*, XVII:53-60.

Citări:

1. Bemal, S., Anil, A.C. 2016. Genetic and ecophysiological traits of *Synechococcus* strains isolated from coastal and open ocean waters of the Arabian Sea. *FEMS Microbiology Ecology*, 92(11): fiw162.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 2 puncte

6. Dragoș, N., Bercea, V., Bica, A., Drugă B., Nicoară, A., Coman, C., 2010. Astaxanthin production from a new strain of *Haematococcus pluvialis* grown in batch culture. *Annals RSCB*, XV:353-361.

Citări:

Khoo K.S., Lee S.Y., Ooi C.W., Fu X., Miao X., et al. 2019. Recent advances in biorefinery of astaxanthin from *Haematococcus pluvialis*. *Bioresource Technology*, 288:121606.

Jung S.M., Jeon J.Y., Park T.H., Yoon J.H., Lee K.S., Shin H.W. 2019. Enhancing production-extraction and antioxidant activity of astaxanthin from *Haematococcus pluvialis*. *Journal of Environmental Biology*, 40(5):924-931.

- Hong, M.E., Choi, H.I., Kwak, H.S., Hwang, S.-W., Sung, Y.J., et al. 2018. Rapid selection of astaxanthin-hyperproducing *Haematococcus* mutant via azide-based colorimetric assay combined with oil-based astaxanthin extraction. *Bioresource Technology*, 267:175-181.
- Liu, J.H., Zhang, L., Zha, D.C., Chen, L.Q., Chen, X.X., Qi, Z.M. 2018. Biosorption of malachite green onto *Haematococcus pluvialis* observed through synchrotron Fourier-transform infrared microspectroscopy. *Letters in Applied Microbiology*, 67(4):348-353.
- Jackson, B.A., Bahri, P.A., Moheimani, N.R. 2017. Repetitive non-destructive milking of hydrocarbons from *Botryococcus braunii*. *Renewable and Sustainable Energy Reviews*, 79:1229-1240.
- Panis, G., Carreon, J.R. 2016. Commercial astaxanthin production derived by green alga *Haematococcus pluvialis*: A microalgae process model and a techno-economic assessment all through production line. *Algal Research*, 18:175-190.
- Shah, M.M.R., Liang, Y., Cheng, J.J., Daroch, M. 2016. Astaxanthin-producing green microalga *Haematococcus pluvialis*: From single cell to high value commercial products. *Frontiers in Plant Science*, 7:531.
- Vanags, J., Kunga, L., Dubencovs, K., Galvanauskas, V., Grigs, O. 2015. Influence of light intensity and temperature on cultivation of microalgae *Desmodesmus communis* in flasks and laboratory-scale stirred tank photobioreactor. *Latvian Journal of Physics and Technical Sciences*, 52: 59-70.
- Santhanam, R. 2015. Nutritional freshwater life. *Nutritional Freshwater Life*, pp. 1-298.
- Hudek, K., Davis, L.C., Ibbini, J., Erickson, L. 2014. Commercial products from algae. *Algal Biorefineries*, 1:275-295.
- Sahandi, J. 2011. Reproduction of Persian Gulf anemone fish (*Amphiprion clarkii*) in captive system. *AACL Bioflux*, 4(5):704-708.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 12 puncte

7. Bercea, V., Coman, C., 2008. The photochemical activity of PS II and PS I photosystems in green alga *Mougeotia sp.* in the presence of ascorbate and hydrogen peroxide in state 2 transition. Stud.Univ. Babeş-Bolyai, Biologia, LIII:75-89

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

8. Coman, C., Drugă, B., Bica, A., Nicoară, A., Dragoş, N., 2008. A molecular approach to diversity estimation of cyanobacteria from Marghita and Roşiori thermomineral drillings (Bihor county). Stud. Univ. Babeş-Bolyai, Biologia, LIII:71-80.

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

9. Coman, C., Drugă, B., Bica, A., Nicoară, A., Dragoş, N., 2007. Diversitatea moleculară a cianobacteriilor asociate forajului termomineral de la Marghita (jud. Bihor). Analele SNBC, XII:272-281.

Citări: 0.

Punctaj articol conform formulei (3) din ordinul 6.129/2016: 1 punct

Articole în reviste indexate BDI ca si contributor

1. Szöke-Nagy, T., Hegedüs, A., Baricz, A., Chiriac, C. M., Szekeres, E., Coman, C., Dragoş, N., 2015. Identification, isolation and bioinformatic analysis of squalene synthase-like cDNA fragments in *Botryococcus terribilis* AICB 870 strain, *Studia Univ. Babeş-Bolyai, Biologia*, 60(1):23-37.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

2. Mituleţu, M., Bercea, V., Coman, C., Drugă, B., Sicora, C., 2013. Induction of photosynthetic state transitions in the cyanobacterium *Microcystis aeruginosa* AICB 702. *Annals RSCB*, XVIII:56-66.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

3. Mituleţu, M., Drugă, B., Hegedüs, A., Coman, C., Dragoş, N., 2013. Phylogenetic analysis of *Microcystis* strains (Cyanobacteria) based on the 16S-23S ITS and cpcBA-IGS markers, *Annals RSCB*, XVIII:22-31.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

4. Bica, A., Barbu-Tudoran, L., Drugă, B., Coman, C., Nicoară, A., Szöke-Nagy, T., Dragoş, N., 2012. *Desmodesmus communis* (Chlorophyta) from Romanian freshwaters: coenobial morphology and molecular taxonomy based on the ITS2 of new isolates. *Annals RSCB*, XVII:16-28.

Citari:

Lortou U., Gkelis S. 2019. Polyphasic taxonomy of green algae strains isolated from Mediterranean freshwaters. *Journal of Biological Research (Greece)*, 26(1):11.

Akgül, F., Kizilkaya, İ.T., Akgül, R., Erduğan, H. 2017. Morphological and molecular characterization of scenedesmus-like species from Ergene River Basin (Thrace, Turkey). *Turkish Journal of Fisheries and Aquatic Sciences*, 17(3):609-619.

Baudelet, P.-H., Ricochon, G., Linder, M., Muniglia, L. 2017. A new insight into cell walls of Chlorophyta. *Algal Research*, 25:333-371.

Hegewald, E., Braband, A. 2017. A taxonomic revision of *desmodesmus* serie *desmodesmus* (Sphaeropleales, Scenedesmaceae). *Fottea*, 17(2):191-208.

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 3,5 puncte

5. Bercea, V., Coman, C., Sicora, C., Dragoş, N., 2011. The photochemical PS II activity in cyanobacterial strains belonging to the nostocales group in anaerobiosis conditions. *Stud. Univ. Babeş-Bolyai, Biologia*, 56:13-25.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

6. Dragoş, N., Mocan, A., Sălăjean, C., Nicoară, A., Bica, A., Drugă, B., Coman, C., Bercea, V., 2010. The effects of temperature on growth and lipid fatty acids composition in

cyanobacterium *Synechocystis* sp. strain AICB 51. Stud. Univ. Babeş-Bolyai, Biologia, LV:51-59.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

7. Drugă B., Bica Adriana, Coman C., Nicoară Ana, Dragoş N., 2009. New Primer Combination for Sequencing the Cyanobacterial 16S rRNA Gene. Analele SRBC, XIV:33-38.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

8. Bica, A., Andrei, C., Drugă, B., Coman, C., Nicoara, A., Dragoş, N., 2007. Filogenia algei verzi *Botryococcus braunii* Kützing pe baza secvenţelor ADNr 18S și 16S. Analele SNBC, XII:282-291

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

9. Bica, A., Coman, C., Drugă, B., - Dragoş, N., 2007. Predicţia exprimării genei *ftsZ* la cianobacterii pe baza preferenţialităţii de utilizare a codonilor, Analele SNBC, XII:209-217.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

10. Drugă, B., Sofronie, I., Văsar, I., Coman, C., Bica, A., Nicoara, A., Dragoş, N., 2008. The molecular diversity of cyanobacterial mats associated with thermal springs. Stud. Univ. Babeş-Bolyai, Biologia, LIII:59-69.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

11. Drugă, B., Bica, A., Coman, C., Nicoara, A., Bercea, V., Dragos, N., 2007. Markerii moleculari în filogenia și taxonomia tulpinilor cianobacteriene din genul *Microcystis*. Analele SNBC, XII:263-271.

Citari: 0

Punctaj articol conform formulei (4) din ordinul 6.129/2016: 0,7 puncte

Brevete omologate, ca autor principal:

Coman, C., Drugă, B., Hegedus, A., Mitulețu, M., Sicora, C., 2013. Tulpină de *Anabaenopsis* sp. și metodă îmbunătățită de producere a hidrogenului dizolvat prin cultivarea acesteia. Nr. brevet: 130066/27.02.2015.

Punctaj conform formulei (3) din ordinul 6.129/2016: 1 punct

Capitole în volume la alte edituri internaționale

Coman, C., Drugă, B., Bica, A., Barbu-Tudoran, L., Dragoș N., 2012. A microbial mat developed around a man-made geothermal spring from Romania: structure and cyanobacterial composition. In: Noffke, N., Chafetz, H., (eds.), *Microbial mats in siliciclastic depositional systems through time*, SEPM Special Publication No. 101, SEPM (Society for Sedimentary Geology), ISBN 978-1-56576-314-2, pp. 47–53.

Citari: 0

Punctaj conform formulei (11) din ordinul 6.129/2016: 4 puncte

Capitole/volume la edituri naționale

1. Coman, C., Soran, M.-L., Farkas, A., 2016. Prelevarea probelor de mediu pentru investigarea reziduurilor de antibiotice, a diversității microbiene și a rezistenței la antibiotice. În: Coman, C. (Ed.), 2016. *Ghid metodologic de monitorizare a antibioticelor și a rezistenței la antibiotice în mediul înconjurător/Methodological guide for monitoring antibiotics and antibiotic resistance in the environment*. Accent, ISBN 978-606-561-165-8, pp. 25-41.

Citari: 0

Punctaj conform formulei (12) din ordinul 6.129/2016: 3,33 puncte

2. Coman, C., Soran, M.-L., Farkas, A., 2016. Environmental sampling for investigations of antibiotic residues, microbial diversity and antibiotic resistance. In: Coman, C. (Ed.), 2016. *Ghid metodologic de monitorizare a antibioticelor și a rezistenței la antibiotice în mediul înconjurător/Methodological guide for monitoring antibiotics and antibiotic resistance in the environment*. Accent, ISBN 978-606-561-165-8, pp. 201-214.

Citari: 0

Punctaj conform formulei (12) din ordinul 6.129/2016: 3,33 puncte

Editor/redactor/coordonator cărți la edituri naționale

Coman, C. (Ed.), 2016. Ghid metodologic de monitorizare a antibioticelor și a rezistenței la antibiotice în mediul înconjurător/Methodological guide for monitoring antibiotics and antibiotic resistance in the environment. Accent, ISBN 978-606-561-165-8, pp. 1-366.

Citari: 0.

Punctaj conform formulei (15) din ordinul 6.129/2016: 20 puncte

Data,

Semnătura,