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Research Article

Isolation and identification of the pyrethroid insecticide deltamethrin degrading bacteria from insects

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Abstract

Many studies have showed that the pesticide residues in the environment increase day by day because of their continuous use. Pesticides can degrade chemically, physically and biologically. Biodegradation is an eco-friendly, inexpensive and highly effective approach compared to other methods. Bacteria are the most commonly used biological agents in biodegradation studies. Widespread use of pyrethroid pesticides such as deltamethrin causes pollution of environment. A total of 14 bacterial isolates were isolated from insects (*Poecilimon tauricola, Locusta migratoria, Gryllus bimaculatus* and *Forficula auricularia*) living in pesticide contaminated environments. These bacterial isolates were identified and characterized as *Pseudomonas aeruginosa, Stenotrophomonas maltophilia, Bacillus atrophaeus, Acinetobacter lwoffii, Rhodococcus coprophilus, Brevundimonas vesicularis, Pseudomonas syringae, Yersinia frederiksenii, Bacillus licheniformis, Enterobacter intermedius and Serratia marcescens based on biochemical and morphological properties and fatty acid profiles. As a result, these bacterial isolates can be used for the remove of deltamethrin at various environments.*

Keywords: Bacteria, Biodegradation, Deltamethrin, Isolation

Böceklerden Piretroid İnsektisit Deltametrin Yıkıcı Bakterilerin İzolasyonu ve Tanımlanması

Öz

Birçok çalışma, sürekli kullanımları nedeniyle ortamdaki pestisit kalıntılarının her geçen gün arttığını göstermiştir. Pestisitler kimyasal, fiziksel ve biyolojik olarak parçanabilirler. Biyodegradasyon, diğer yöntemlere kıyasla çevre dostu, ucuz ve oldukça etkili bir yaklaşımdır. Biyodegradasyon çalışmalarında bakteriler en sık kullanılan biyolojik ajanlardır. Deltametrin gibi piretroid pestisitlerin yaygın kullanımı çevrenin kirlenmesine neden olmaktadır. Pestisit kontamine ortamlarda yaşayan böceklerden (*Poecilimon tauricola, Locusta migratoria, Gryllus bimaculatus* ve *Forficula auricularia*) toplam 14 bakteri izolatı izole edilmiştir. Bu bakteri izolatları, biyokimyasal ve morfolojik özellikleri ve yağ asidi profillerine dayanarak *Pseudomonas aeruginosa, Stenotrophomonas maltophilia, Bacillus atrophaeus, Acinetobacter lwoffii, Rhodococcus coprophilus, Brevundimonas vesicularis, Pseudomonas syringae, Yersinia frederiksenii, Bacillus licheniformis, Enterobacter intermedius ve Serratia marcescens* olarak tanımlanmış ve karakterize edilmiştir. Sonuç olarak, bu bakteri izolatları çeşitli ortamlarda deltametrinin parçaalanması için kullanılabilir.

Anahtar Kelimeler: Bakteriler, Biyodegradasyon, Deltametrin, İzolasyon

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1. Introduction

Natural or synthetic pesticides (organochlorine, organophosphate, carbamate, pyrethroids) are widely used to control unwanted pests. Pyrethroids account for about one-fifth of the global agrochemical market. Pyrethroids have potent neurotoxic activity against insects and low toxicity to animals. With the permanent use of pyrethroid worldwide, its residue has become a problem to animals, including humans. Pyrethroid insecticides, e.g., cyphenothrin, fenvalerate, esfenvalerate, deltamethrin, cypermethrin, cyhalothrin, fluvarinate, tralomethrin, cycloprothrin, acrinathrinallethrin, imiprothrin, permethrin and fenpropathrin are used in agriculture, animal health, home, and garden pest control throughout the world (Cycoń and Piotrowska-Seget, 2016; Zhang et al., 2016; Hao et al., 2018).

Pesticides are degraded into simpler and often less toxic chemicals in various ways such as chemical reactions, photodegradation and biodegradation. Biodegradation is an environment friendly, cheap and high efficiency approach compared to other methods. Bacteria and fungi with high enzyme (transferases, isomerases, ligases and hydrolases especially esterases, peroxidases and oxygenases) activity are used in biodegradation studies (Ortiz-Hernández et al., 2013; Ozdal et al., 2017).

Deltamethrin ($C_{22}H_{19}Br_2NO_3$) is a broad-spectrum insecticide belonging to pyrethroids (Figure 1). Deltamethrin is widely used in agriculture because of its low cost, persistence, stability and low toxicity to mammals. It is used for the control of pests such as mosquitoes, cockroaches, flies, ants and fleas due to effective at very low concentrations (Hao et al., 2018; Lu et al., 2019)

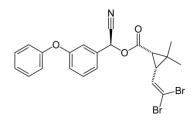


Figure 1. Chemical structure of deltamethrin

Microflora in the digestive tract of insect species is being investigated. The nutrient-rich digestive tract of insects is an appropriate growth environment for these microorganisms. The bacterial flora in the digestive tract of the insect has a very variable and broad enzymatic potential. Insect gut bacterial isolates have been demonstrated to break down many compounds such as pesticide (Ozdal et al., 2016a, b). Insect intestines provide a suitable medium for gene transfer between bacteria. Microorganisms can adapt to new environments by acquiring different features with horizontal gene transfer, conjugative plasmid and simple mutations to different environmental conditions (Pietri et al., 2018; Ramakrishnan et al., 2019). In this context, it is highly possible to isolate pesticide resistant microorganisms from insect intestines.

In many insect groups, resistance to pesticides occurs as a result of the use of pesticides. The intestinal flora of insects, which are observed to be resistant to pesticides, is very rich in bacteria that can be used in the biodegradation of pesticides. The purpose of this study was to isolate the bacteria capable of degrading deltamethrin from different insects.

2. Materials and Methods

2.1. Chemicals

Deltamethrin and other chemicals used in the study were of analytical purity and were obtained from Sigma and the media were obtained from Merck and Difco.

2.2. Preparation of media and solutions used in the study

Carbon-free mineral salt medium (MSM) was used for isolation of deltamethrin degrading bacteria. The medium contained 2.0 g of $(NH_4)_2SO_4$, 0.2 g of MgSO₄ 7H₂O, 0.01 g of CaCl₂ 2H₂O, 0.001 g of FeSO₄ 7H₂O, 1.5 g of Na₂H-PO₄ 12H₂O, and 1.5 g of KH₂PO₄ per litre of deionized water (Cycoń et al., 2014). The final pH value was adjusted to 7.2. After autoclaving (121 °C, 15 min) and cooling, the medium was supplemented with 100 mg/L deltamethrin.

2.3. Insects used in the study

The insects belonging to Orthoptera and Dermaptera were collected from different regions during the spring-summer period and species identification Prof. Dr. Orhan Erman. Bacteria that can use deltamethrin as a carbon source were isolated from insects.

2.4. Isolation of Deltamethrin Degrading Bacteria

The insect samples were subjected to surface sterilization with 70% ethyl alcohol for 3 minutes and after alcohol removal with sterile physiological water (SFS), homogenized by crushing in a sterile mortar with SFS (Okay et al., 2013). Serial dilutions of homogenate were prepared and 0.1 mL of liquid was inoculated to liquid minimal medium containing 100 mg/L of deltamethrin. After one week, 1 ml of each culture was re-inoculated into new deltamethrin-MSM medium and further incubated at 30°C and 150 rpm for 7 days. This

subculture was repeated under the same culture conditions, and then an aliquot (0.2 ml) from each culture was applied to solid deltamethrin-MSM for isolation of single colonies. Colonies of different character were isolated by transferring to Tryptic Soy Agar plates and stored on slant agar at $+ 4 \circ C$.

2.5. Identification of isolates

Deltamethrin degrading bacteria were identified using morphological, cultural, biochemical properties (Gram, cell shape, endospores, movement, catalase, oxidase) (Harley and Prescott, 2002) and fatty acid profiles (Kotan et al., 2006).

3. Results and Discussion

There is a close relationship between bacteria and other living things. Therefore, insect microflora enables us to find new and biotechnological microorganisms. *Serratia marcescens* MO-1 isolated from grasshopper (*Poecilimon tauricola*) has both chitinase activity (Okay et al., 2013) and the ability to produce prodigiosin pigment (Kurbanoglu et al., 2015) which has antimicrobial and anticancer properties. Also, *Pseudomonas aeruginosa* OG1 isolated from cockroaches (*Blatta orientalis*) can produce pyocyanin, a pigment of biotechnological importance (Ozdal, 2019). Insects can change their ecological and physiological properties thanks to symbiotic bacteria (Pietri and Liang, 2018). Kikuchi et al., (2012) indicated that bacteria of the genus *Burkholderia* develop resistance against the fenitrothion (organophosphate pesticide) in the bean bug (*Riptortus pedestris*). Chlorpyrifos and fipronil resistant strains of diamond back moth (*Plutella xylostella*) have higher levels of Lactobacillales, Pseudomonadales and Xanthomonadales compared to susceptible insects (Xia et al., 2013). *Stenotrophomonas maltophilia* OG-2, isolated from the intestine of the cockroach (*Blatta orientalis*), can degrade both α -endosulfan (Ozdal et al., 2017) and synthetic pyrethroid α -cypermethrin (Gur et al., 2014).

In this study, insects belonging to Orthoptera and Dermaptera were collected from the areas where insecticides were used (Table 1). As a result of isolations, 14 bacteria were isolated on solid medium containing deltamethrin. Table 1 lists the insect species and isolate groups from which the isolates were obtained.

Strain	Insect Name	Place of	Family	Order	
Group		collection			
DPT	Poecilimon tauricola (Ramme 1951)	Erzurum	Tettigoniidae	Orthoptera	
DLM	Locusta migratoria (Linnaeus 1758)	İzmir	Acrididae	Orthoptera	
DGB	Gryllus bimaculatus (De Geer 1773)	Antalya	Gryllidae	Orthoptera	
DFA	Forficula auricularia (Linnaeus 1758)	Samsun	Forficulidae	Dermaptera	

 Table 1 Insect species from which isolates were obtained

A total of 14 bacterial isolates were isolated from medium containing deltamethrin based on visible colony differences. Among all these 14 deltamethrin degrading bacterial isolates, 3 were Gram-positive rods and 11 were Gram-negative rods. Total 2 isolates indicated positive results for endospore. All the isolates were catalase positive. Of these isolates, 3 were oxidase positive, 10 were motile and 3 were urease positive (Table 2).

Table 2 Cultural, morphological and biochemical properties of isolates								
Isolate code	Gram	Cell shape	Endospor	Motile	Catalase	Oxidase	Urease	
DGB1	-	Rod	-	+	+	+	-	
DLM1	+	Rod	-	-	+	-	+	
DPT1	+	Rod	+	+	+	-	-	
DPT2	-	Rod	-	+	+	-	-	
DLM2	-	Rod	-	-	+	-	+	
DFA1	+	Rod	+	+	+	-	-	
DPT3	-	Rod	-	+	+	-	-	
DPT4	-	Rod	-	+	+	-	-	
DPT5	-	Rod	-	+	+	+	-	
DLM3	-	Rod	-	+	+	-	+	
DFA2	-	Rod	-	+	+	-	-	
DGB2	-	Rod	-	+	+	+	-	
DGB3	-	Rod	-	-	+	-	-	
DLM4	-	Rod	-	-	+	-	-	

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Fatty acids	Bacteria													
·	DGB1	1 DGB2	DGB3	DFA1		DPT1	DPT2	DPT3	DPT4	DPT5	DLM1	DLM2	DLM3	DLM4
10:0		0.32			0.62				0.30	0.26		0.47	0.21	
10:0 3OH		3.14			0.16		3.3		10.12	3.18			2.2	
12:0		4.3	6.37		2.67		4.9	5.7	1.13	4.2		6.4	5.09	9.8
12:0 2OH		4.9					4.7	0.4	0.42	4.7			4.29	
12:0 3OH	2.4		4.92		2.82		4.3		2.3			0.4		3.2
13:0 iso					0.41									
14:0 iso				1.1	0.67	1.1								
14:0	3.05	0.63			3.80			6.3	4.7	0.58	1.9	1.7	0.65	
14:0 3ÔH								5.57						
15:0 iso				19.1	33.67	12.47								
15:0 Antesio				40.8	12.10	50.34								
15:0	3.38							2.1			2.8	0.2		
16:0 iso				4.7	1.10	3.3								
16:1 w7c								29.76	12.1		6.2			
16:1 w9c					3.58									
16:1 w11c				2.2										
16:0	20.7	24.8	12.5	5.1	9.05	2.74	26.4	29.99	27.3	25.2	23.2	37.2	25.66	19.44
16:0 10-methyl											14.4			
17:1 w8c	2.2		0.93											
17:0 cyclo		0.85					2.3	3.2	13.7	1		2	0.48	
17:0	1.1	0.13						1.9	0.44	0.15	1.9	0.1		
17:1 iso w10c														
17:0 iso				5.9	3.11	5.7	0.3					0.1		
17:0 antesio				15.8	0.41	16.1								
17:0 10-methyl											2.4			
17:0 w Cyclo 7-8												29.1		
18:1 w7c	52.7	42.4	5.85		0.84		13.7	11.7	14.2	41.3		2.1	44.14	3.45
18:1 w9c			37.21		1.62						2.3			21.67
18:0	0.4	0.38	2.33		0.22		2.3	0.46	0.3	0.44		1.1	0.64	
18:0 10-methyl											6.7			
19:0												0.7		
16:1 w7c/15:0 iso	6.8	16.66	26.45		10.96		30.6			17.43		3.6	18.29	39.6
20H														
17:1 w8c	2.2	0.22								0.20	0.8			
14:0 3OH/16:1 iso												9.4		

Table 3 Fatty acid profiles of bacteria isolated from insects

According to the results of MIS analysis, fatty acid profiles of the isolates are summarized in Table 3. As a result of MIS analysis, 8 different genera and 13 different species of bacterial isolates were identified. Isolates based on morphological, biochemical and fatty acid data were identified as *Acinetobacter lwoffii* (DLM4), *Pseudomonas aeruginosa* (DPT5, DBG2), *Stenotrophomonas maltophilia* (DFA2), *Bacillus licheniformis* (DFA1), DLM1), *Bacillus atrophaeus* (DPT1), *Pseudomonas syringae* (DPT2), *Yersinia frederiksenii* (DLM2), *Enterobacter intermedius* (DPT3), *Serratia marcescens* (DPT4) and *Flavimonas oryzihabitans* (DLM3). The genera of the isolated bacteria were mainly identified as *Pseudomonas* and *Bacillus*. The highest bacterial diversity was observed in *Poecilimon tauricola* (5) and *Locusta migratoria* (4).

Table 4 Deltamethrin degrading microorganisms isolated from different environments						
Strain	Source	Reference				
Streptomyces aureus HP-S-01	Activated sludge	Chen et al., 2011				
Bacillus cereus Y1	Deltametrin contaninated soil	Zhang et al., 2016				
Lysinibacillus fusiformis ZJ6	Soil	Hao et al., 2018				
Acinetobacter calcoaceticus MCm5	Pyrethroid contaninated soil	Akbar et al., 2015a				
Brevibacillus parabrevis FCm9	-					
Sphingomonas sp. RCm6						
Bacillus megaterium JCm2	Pyrethroid contaninated soil	Akbar et al., 2015b				
Rhodococcus sp. JCm5	-					
Ochrobactrum anthropi JCm1						
Pseudomonas aeruginosa JQ-41	Pyrethroid contaninated soil	Song et al., 2015				
Serratia marcescens DeI-1, DeI-2	Deltamethrin treated soil	Cycoń et al., 2014				
Acinetobacter baumannii ZH-14	Sewage sludge	Zhan et al., 2018				

Many different bacteria have been isolated and characterized with their ability to degradation various pesticides. In previous studies, bacteria capable of degrading deltamethrin were mostly isolated from agricultural areas where intensive pesticides were used. However, the potential of these microorganisms to degrade deltamethrin has been confirmed for some bacteria of the genera *Acinetobacter*, *Bacillus*, *Brevibacillus*, *Pseudomonas*, *Serratia*, *Rhodococcus* (Table 4). Song et al., (2015) studied the deltamethrin biodegradation with *Pseudomonas aeruginosa* JQ-41 strain isolated from the pyrethroid contaminated soil. In another study, *Acinetobacter calcoaceticus* MCm5 was used in biodegradation of deltamethrin (Akbar et al., 2015a). Similar bacteria were isolated

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in this study. When Table 4 is analyzed, it is seen that the bacteria used in deltamethrin degradation are generally isolated from soil and sludge. All bacteria obtained from this study were isolated from insect flora. In addition, it has been determined that new species may be effective in deltamethrin biodegradation.

The bacteria isolated in this study can undoubtedly be used in biodegradation studies. As seen in Table 5, different strains of the species isolated in this study have been reported to have been used for the degradation of many different pesticides. It has been determined that insects are important source for the isolation of bacteria that break down pesticides.

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Table 5 Compariso	n of the use	e of the isola	ated species	s in the de	egradation of	f different pesticides
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Bacteria	Pesticide	References		
Acinetobacter lwoffii	Endosulfan	Ozdal et al., 2016b		
	Atrazine	Yang et al., 2017		
Pseudomonas aeruginosa	Endosulfan	Ozdal et al., 2016b		
	Fenvalerate	Fulekar, 2009		
	Acephate, dimethoate, parathion, chlorpyrifos, malathion	Ramu and Seetharaman, 2014		
	chlorpyrifos and dichlorvos	Gaonkar et al., 2019		
Stenotrophomonas maltophilia	α -endosulfan, α -cypermethrin	Gur et al., 2014; Ozdal et al., 2017		
	Diazinon	Pourbabaee et al., 2018		
Bacillus licheniformis	Fenvalerate	Tang et al., 2018		
	β-cyclodextrin, β-cypermethrin	Zhao et al., 2015		
Bacillus atrophaeus	α-Endosulfan	Ozdal et al., 2016b		
Pseudomonas syringae				
Yersinia frederiksenii	Permethrin	Lee et al., 2004		
Enterobacter intermedius				
Serratia marcescens	DDT	Neerja, Grewal et al., 2016		
Flavimonas oryzihabitans	DDT	Barragan-Huerta et al., 2007		

4. Conlusion

Strains of *Pseudomonas aeruginosa*, *Stenotrophomonas maltophilia*, *Bacillus atrophaeus*, *Acinetobacter lwoffii*, *Rhodococcus coprophilus*, *Brevundimonas vesicularis*, *Pseudomonas syringae*, *Yersinia frederiksenii*, *Bacillus licheniformis*, *Enterobacter intermedius* and *Serratia marcescens*, able to use deltamethrin as the only carbon source, were isolated from *Poecilimon tauricola*, *Locusta migratoria*, *Gryllus bimaculatus* and *Forficula auricularia*. Pesticide resistant insect microbiota has been shown to be a rich source for isolation of microbes that can degradation pesticides and a promising tool for biotechnological discovery in bioemediation programs. In order to find new biocatalysts in the degradation of pesticides, isolation can be made from insects that can live in pesticide environments. As a result, it can be said that isolated deltamethrin degrading microorganisms can be used in the treatment studies in the dirty areas of this insecticide. However, optimization studies are also needed to make biodegradation highly efficient and feasible.

References

- Akbar, S., Sultan, S., & Kertesz, M. (2015a). Determination of cypermethrin degradation potential of soil bacteria along with plant growth-promoting characteristics. *Current Microbiology*, 70(1), 75-84.
- Akbar, S., Sultan, S., & Kertesz, M. (2015b). Bacterial community analysis of cypermethrin enrichment cultures and bioremediation of cypermethrin contaminated soils. *Journal of Basic Microbiology*, 55(7), 819-829.
- Barragan-Huerta, B. E., Costa-Pérez, C., Peralta-Cruz, J., Barrera-Cortés, J., Esparza-García, F., & Rodríguez-Vázquez, R. (2007). Biodegradation of organochlorine pesticides by bacteria grown in microniches of the porous structure of green bean coffee. *International Biodeterioration & Biodegradation*, 59(3), 239-244.
- Chen, S., Lai, K., Li, Y., Hu, M., Zhang, Y., & Zeng, Y. (2011). Biodegradation of deltamethrin and its hydrolysis product 3phenoxybenzaldehyde by a newly isolated *Streptomyces aureus* strain HP-S-01. *Applied Microbiology and Biotechnology*, 90(4), 1471-1483.
- Cycoń, M., & Piotrowska-Seget, Z. (2016). Pyrethroid-degrading microorganisms and their potential for the bioremediation of contaminated soils: a review. *Frontiers in Microbiology*, 7, 1463.
- Cycoń, M., Żmijowska, A., & Piotrowska-Seget, Z. (2014). Enhancement of deltamethrin degradation by soil bioaugmentation with two different strains of *Serratia marcescens*. International Journal of Environmental Science and Technology, 11(5), 1305-1316.
- Fulekar, M. H. (2009). Bioremediation of fenvalerate by *Pseudomonas aeruginosa* in a scale up bioreactor. *Romanian Biotechnological Letters*, 14, 4900-4905.
- Gaonkar, O., Nambi, I. M., & Suresh Kumar, G. (2019). Biodegradation kinetics of dichlorvos and chlorpyrifos by enriched bacterial cultures from an agricultural soil. *Bioremediation Journal*, 23(4), 259-276.
- Gür, Ö., Özdal, M., & Algur, Ö. F. (2014). Biodegradation of the synthetic pyrethroid insecticide α-cypermethrin by *Stenotrophomonas maltophilia* OG2. *Turkish Journal of Biology*, 38(5), 684-689.
- Hao, X., Zhang, X., Duan, B., Huo, S., Lin, W., Xia, X., & Liu, K. (2018). Screening and genome sequencing of deltamethrindegrading bacterium ZJ6. *Current Microbiology*, 75(11), 1468-1476.

Harley, J. P., & Prescott L.M. 2002. Laboratory Exercises in Microbiology. McGraw-Hill Pub. 5th edition. *e-ISSN: 2148-2683*

- Kikuchi, Y., Hayatsu, M., Hosokawa, T., Nagayama, A., Tago, K., & Fukatsu, T. (2012). Symbiont-mediated insecticide resistance. *Proceedings of the National Academy of Sciences*, 109(22), 8618-8622.
- Kotan, R., Sahin, F., & Ala, A. (2006). Identification and pathogenicity of bacteria isolated from pome fruit trees in the Eastern Anatolia region of Turkey. *Journal of Plant Diseases and Protection*, 8-13.
- Kurbanoglu, E. B., Ozdal, M., Ozdal, O. G., & Algur, O. F. (2015). Enhanced production of prodigiosin by Serratia marcescens MO-1 using ram horn peptone. *Brazilian Journal of Microbiology*, 46(2), 631-637.
- Lee, S., Gan, J., Kim, J. S., Kabashima, J. N., & Crowley, D. E. (2004). Microbial transformation of pyrethroid insecticides in aqueous and sediment phases. *Environmental Toxicology and Chemistry: An International Journal*, 23(1), 1-6.
- Lu, Q., Sun, Y., Ares, I., Anadón, A., Martínez, M., Martínez-Larrañaga, M. R., Yuan, X., Wang, M.A., & Martínez, M. A. (2019). Deltamethrin toxicity: A review of oxidative stress and metabolism. *Environmental Research*, 170, 260-281.
- Neerja, Grewal, J., Bhattacharya, A., Kumar, S., Singh, D. K., & Khare, S. K. (2016). Biodegradation of 1, 1, 1-trichloro-2, 2-bis (4chlorophenyl) ethane (DDT) by using *Serratia marcescens* NCIM 2919. *Journal of Environmental Science and Health, Part B*, 51(12), 809-816.
- Okay, S., Özdal, M., & Kurbanoğlu, E. B. (2013). Characterization, antifungal activity, and cell immobilization of a chitinase from *Serratia marcescens* MO-1. *Turkish Journal of Biology*, *37*(6), 639-644.
- Ortiz-Hernández, M. L., Sánchez-Salinas, E., Dantán-González, E., & Castrejón-Godínez, M. L. (2013). Pesticide biodegradation: mechanisms, genetics and strategies to enhance the process. *Biodegradation-Life of Science*, 251-287.
- Ozdal, M., Ozdal, O. G., & Algur, O. F. (2016a). Isolation and characterization of α-endosulfan degrading bacteria from the microflora of cockroaches. *Polish Journal of Microbiology*, 65(1), 63-68.
- Ozdal, Ö. G., Özdal, M., Algur, Ö. F., & Sezen, A. (2016b). Isolation and identification of α-Endosulfan degrading bacteria from insect microflora. *Turkish Journal of Agriculture-Food Science and Technology*, 4(4), 248-254.
- Ozdal, M. (2019). A new strategy for the efficient production of pyocyanin, a versatile pigment, in *Pseudomonas aeruginosa* OG1 via toluene addition. *3 Biotech*, 9(10), 374.
- Ozdal, M., Ozdal, O. G., Algur, O. F., & Kurbanoglu, E. B. (2017). Biodegradation of α-endosulfan via hydrolysis pathway by *Stenotrophomonas maltophilia* OG2. 3 Biotech, 7(2), 113.
- Pietri, J. E., & Liang, D. (2018). The Links Between Insect Symbionts and Insecticide Resistance: Causal Relationships and Physiological Tradeoffs. *Annals of the Entomological Society of America*, 111(3), 92-97.
- Pietri, J. E., Tiffany, C., & Liang, D. (2018). Disruption of the microbiota affects physiological and evolutionary aspects of insecticide resistance in the German cockroach, an important urban pest. PloS One, 13(12), e0207985.
- Pourbabaee, A. A., Soleymani, S., Farahbakhsh, M., & Torabi, E. (2018). Biodegradation of diazinon by the Stenotrophomonas maltophilia PS: pesticide dissipation kinetics and breakdown characterization using FTIR. International Journal of Environmental Science and Technology, 15(5), 1073-1084.
- Ramakrishnan, B., Venkateswarlu, K., Sethunathan, N., & Megharaj, M. (2019). Local applications but global implications: Can pesticides drive microorganisms to develop antimicrobial resistance? *Science of The Total Environment*, 654, 177-189.
- Ramu, S., & Seetharaman, B. (2014). Biodegradation of acephate and methamidophos by a soil bacterium *Pseudomonas aeruginosa* strain Is-6. *Journal of Environmental Science and Health, Part B*, 49(1), 23-34.
- Song, H., Zhou, Z., Liu, Y., Deng, S., & Xu, H. (2015). Kinetics and mechanism of fenpropathrin biodegradation by a newly isolated *Pseudomonas aeruginosa* sp. strain JQ-41. *Current Microbiology*, 71(3), 326-332.
- Tang, J., Liu, B., Shi, Y., Zeng, C. Y., Chen, T. T., Zeng, L., & Zhang, Q. (2018). Isolation, identification, and fenvalerate-degrading potential of Bacillus licheniformis CY-012. *Biotechnology & Biotechnological Equipment*, 32(3), 574-582.
- Xia, X., Zheng, D., Zhong, H., Qin, B., Gurr, G. M., Vasseur, L., Lin H., Bai J., He, W., & You, M. (2013). DNA sequencing reveals the midgut microbiota of diamondback moth, *Cimex xylostella* (L.) and a possible relationship with insecticide resistance. PLoS One. 8(7): e68852.
- Yang, F., Jiang, Q., Zhu, M., Zhao, L., & Zhang, Y. (2017). Effects of biochars and MWNTs on biodegradation behavior of atrazine by Acinetobacter lwoffii DNS32. Science of the Total Environment, 577, 54-60.
- Zhan, H., Wang, H., Liao, L., Feng, Y., Fan, X., Zhang, L., & Chen, S. (2018). Kinetics and novel degradation pathway of permethrin in *Acinetobacter baumannii* ZH-14. *Frontiers in Microbiology*, *9*, 98.
- Zhang, H., Zhang, Y., Hou, Z., Wang, X., Wang, J., Lu, Z., Zhao, X., Sun, F., & Pan, H. (2016). Biodegradation potential of deltamethrin by the *Bacillus cereus* strain Y1 in both culture and contaminated soil. *International Biodeterioration & Biodegradation*, 106, 53-59.
- Zhao, J., Chi, Y., Liu, F., Jia, D., & Yao, K. (2015). Effects of two surfactants and beta-cyclodextrin on beta-cypermethrin degradation by *Bacillus licheniformis* B-1. *Journal of Agricultural and Food Chemistry*, *63*(50), 10729-10735.