Research Article

In Vitro Antibacterial Activity of Propyl-Propane-Thiosulfinate and Propyl-Propane-Thiosulfonate Derived from *Allium* spp. against Gram-Negative and Gram-Positive Multidrug-Resistant Bacteria Isolated from Human Samples

Antonio Sorlozano-Puerto (),¹ Maria Albertuz-Crespo,¹ Isaac Lopez-Machado,¹ Juan Jose Ariza-Romero,² Alberto Baños-Arjona,² Manuela Exposito-Ruiz,³ and Jose Gutierrez-Fernandez ()^{1,4}

¹Department of Microbiology, School of Medicine and PhD Program in Clinical Medicine and Public Health, University of Granada-ibs, Granada, Spain

²Department of Microbiology and Biotechnology, DMC Research Center, Granada, Spain

³Unit of Methodology of Research and Biostatistics, Virgen de las Nieves University Hospital-ibs, Granada, Spain

⁴Laboratory of Microbiology, Virgen de las Nieves University Hospital-ibs, Granada, Spain

Correspondence should be addressed to Antonio Sorlozano-Puerto; asp@ugr.es

Received 30 April 2018; Accepted 1 August 2018; Published 17 September 2018

Academic Editor: Taoufik Ghrairi

Copyright © 2018 Antonio Sorlozano-Puerto et al. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

Background. The aim of this study was to compare the in vitro antibacterial activity of two compounds derived from Alliaceae, PTS (propyl-propane-thiosulfinate), and PTSO (propyl-propane-thiosulfonate), with that of other antibiotics commonly used against bacteria isolated from humans. Materials and Methods. A total of 212 gram-negative bacilli and 267 gram-positive cocci isolated from human clinical samples and resistant to at least one group of antibiotics were selected. In order to determine the minimum inhibitory concentration (MIC) and minimum bactericidal concentration (MBC) to various antibiotics as well as PTS and PTSO, all isolates underwent broth microdilution assay. Results. PTS showed moderate activity against Enterobacteriaceae with MIC₅₀ (and MBC₅₀) and MIC₉₀ (and MBC₉₀) values of 256-512 mg/L, while PTSO showed greater activity with MIC₅₀ and MIC₉₀ values of 64-128 mg/L and MBC₅₀ and MBC₉₀ values of 128-512 mg/L. These data show the bactericidal activity of both compounds and indicate that PTSO was more active than PTS against this group of bacteria. Both compounds showed lower activity against P. aeruginosa (MIC₅₀ = 1024 mg/L, MIC₉₀ = 2048 mg/L, MBC₅₀ = 2048 mg/L, and MBC₉₀ = 2048 mg/L, for PTS; MIC₅₀ = 512 mg/L, $MIC_{90} = 1024 \text{ mg/L}, MBC_{50} = 512 \text{ mg/L}, \text{ and } MBC_{90} = 2048 \text{ mg/L}, \text{ for PTSO})$ compared to those obtained in others nonfermenting gram-negative bacilli (MIC₅₀ = 128 mg/L, MIC₉₀ = 512 mg/L, MBC₅₀ = 128 mg/L, and MBC₉₀ = 512 mg/L, for PTS; MIC₅₀ = 64 mg/L, MIC₉₀ = 256 mg/L, for PTSO) and also indicate the bactericidal activity of both compounds against these groups of bacteria. Finally, the activity against S. aureus, E. faecalis, and S. agalactiae was higher than that observed against enterobacteria, especially in the case of PTSO ($MIC_{50} = 8 \text{ mg/L}$, $MIC_{90} = 8 \text{ mg/L}$, $MBC_{50} = 32 \text{ mg/L}$, and MBC_{90} = 64 mg/L, in S. aureus; $MIC_{50} = 4 mg/L$, $MIC_{90} = 8 mg/L$, $MBC_{50} = 8 mg/L$, and $MBC_{90} = 16 mg/L$, in E. faecalis and S. agalactiae). Conclusion. PTS and PTSO have a significant broad spectrum antibacterial activity against multiresistant bacteria isolated from human clinical samples. Preliminary results in present work provide basic and useful information for development and potential use of these compounds in the treatment of human infections.

1. Introduction

The use of conventional antibiotics for the prevention of infectious diseases and as growth promoters in animal production has fostered the appearance of resistant bacteria and the transmission of these pathogens to humans [1]. In addition, the use and sometimes misuse of antibiotics in humans has increased the occurrence of infections (urinary



FIGURE 1: Chemical structure of propyl-propane-thiosulfinate (PTS).



FIGURE 2: Chemical structure of propyl-propane-thiosulfonate (PTSO).

tract infections, respiratory tract infections, skin and soft tissue infections, etc.) caused by multiresistant bacteria, which has reduced the therapeutic options and has made necessary the selection of new molecules with antibacterial properties [2]. Natural compounds obtained from vegetables with antibacterial properties could be considered an alternative to conventional antibiotics [3].

In recent years, the antibacterial properties of some compounds obtained from *Allium* plants such as garlic (*Allium sativum*) and onion (*Allium cepa*) have been described. These can inhibit the growth of a range of gram-positive and gram-negative bacteria, including both pathogenic and commensal bacteria in humans and animals [4, 5]. *Allium*derived products have been reported to be effective even against those strains that have become resistant to antibiotics [6].

Two of these Allium-derived compounds, propylpropane-thiosulfinate (PTS) (Figure 1) and propyl-propanethiosulfonate (PTSO) (Figure 2), are organosulphurate products obtained by decomposition of initial compounds naturally present in garlic bulbs as alliin and allicin. In several in vitro and in vivo studies against pathogenic bacteria from animals, both compounds have showed an antibiotic activity [3, 7, 8]. While the precise mechanism of action is not yet known, the main antibacterial effect of thiosulfinates (as allicin) has been reported to be due to (i) its accessibility resulting from high permeability through phospholipid membranes [9]; (ii) its chemical reaction with thiol groups of various enzymes such as the bacterial acetyl-CoA-forming system, consisting of acetate kinase and phosphotransacetyl-CoA synthetase, blocking acetate incorporation into fatty acids and inhibiting the formation of lipids [10]; and (iii) the inhibition of RNA polymerase and RNA synthesis [11].

Therefore, the aim of this study was to compare the *in vitro* antibacterial activity of the compounds derived from garlic PTS and PTSO with that of other antibiotics commonly used against gram-negative and gram-positive multidrug-resistant bacteria isolated from human clinical samples.

2. Material and Methods

2.1. Antibiotics, PTS and PTSO. All antibiotics were purchased from Sigma-Aldrich (Madrid, Spain) and each

antibiotic was dissolved according to the manufacturer's recommendations.

PTS and PTSO (95% purity) were supplied by DMC Research (Alhendín, Granada, Spain) and dissolved in polysorbate-80 to a final concentration of 50%. The biosynthesis of propyl-propane-thiosulfinate (PTS) and propylpropane-thiosulfonate (PTSO) is made from propiin, an amino acid derived from L-cysteine found in *Allium* species. The first step of the biosynthesis is the formation of a sulfenic acid, which is highly reactive and immediately produces PTS by a condensation reaction. In the last step, oxidation of PTS induces its dismutation in PTSO and propyl disulfide that can be oxidized and transformed to PTSO and that way the oxidation of PTS to PTSO is completed.

2.2. Bacterial Isolates. A total of 212 gram-negative bacilli and 267 gram-positive cocci isolated from clinical samples obtained from 479 different patients were selected. Identification and susceptibility studies were performed using WIDER system (Francisco Soria Melguizo, Madrid, Spain) or MicroScan system (Siemens Healthcare Diagnostics, Madrid, Spain). The susceptibility results obtained through these systems allowed the selection of isolates, based on the resistance presence to at least one group of antibiotics commonly used in the treatment of infections caused by these bacteria.

The presence of extended-spectrum beta-lactamaseproducing *Enterobacteriaceae* (ESBL) was confirmed by the diffusion method with disks containing cefotaxime (30 μ g), cefotaxime/clavulanic acid (30/10 μ g), ceftazidime (30 μ g), and ceftazidime/clavulanic acid (30/10 μ g). The resistance to methicillin was confirmed using the Mueller–Hinton agar diffusion procedure with 30 μ g cefoxitin disks. Both procedures were performed as recommended by the Clinical and Laboratory Standards Institute [12].

A total of 151 clinical isolates of *Enterobacteriaceae* (68 *Escherichia coli*, 33 *Klebsiella pneumoniae*, 6 *Klebsiella oxytoca*, 15 *Salmonella* spp., 17 *Yersinia enterocolitica*, 7 *Enterobacter cloacae*, 2 *Providencia stuartii*, 1 *Citrobacter amalonaticus*, 1 *Kluyvera cryocrescens*, and 1 *Proteus vulgaris*), 61 of nonfermenting gram-negative bacilli (40 *Pseudomonas aeruginosa*, 9 *Acinetobacter baumannii*, 7 *Aeromonas hydrophila*, 3 *Stenotrophomonas maltophilia*, 1 *Achromobacter xylosoxidans*, and 1 *Comamonas acidovorans*), 112 *Staphylococcus aureus* (all of them methicillin-resistant), 54 *Enterococcus faecalis* (all of them fluoroquinolone-resistant), and 101 *Streptococcus agalactiae* were selected. All isolates were stored at -40°C until the susceptibility study by microdilution.

2.3. In Vitro Antibacterial Assay. In order to determine the antibacterial susceptibilities, all 479 isolates underwent broth microdilution assay in Cation-Adjusted Mueller–Hinton Broth (CAMHB) following the guidelines of the CLSI [12]. Broth microdilution testing was performed with 96-well, round-bottom microtiter plates with a final concentration of the bacterial cell suspension equal to 1×10^5 colony forming units per milliliter (CFU/ml) in each well.

Each plate included negative controls (medium only) and 11 serial twofold dilutions of each antibiotic, PTS, or PTSO. The positive controls (only bacterial suspension without antibiotics) were added per well in a separate round-bottom plate.

concentration ranges (in mg/L) assayed The for Enterobacteriaceae for each antibiotic were the following: amoxicillin/clavulanate (0.25/0.125-256/128),piperacillin/tazobactam (0.5/4-512/4), cefuroxime (0.5-512), cefoxitin (0.5-512), cefotaxime (0.125-128), ceftazidime (0.5-512), cefepime (0.25-256), imipenem (0.016-16), meropenem (0.016-16), gentamicin (0.125-128), tobramycin (0.125-128), amikacin (0.5-512), ciprofloxacin (0.125-128), trimethoprim/sulfamethoxazole (0.06/1.1875-64/1216), and nitrofurantoin (1-1024). The concentration ranges assayed for nonfermenting gram-negative bacilli for each antibiotic were piperacillin/tazobactam (0.5/4-512/4), ceftazidime (0.5-512), cefepime (0.25-256), imipenem (0.125-128), meropenem (0.125-128), gentamicin (0.125-128), tobramycin (0.125-128), amikacin (0.5-512), and ciprofloxacin (0.125-128). The concentration ranges for staphylococci were gentamicin (0.25-256), tobramycin (0.25-256), erythromycin (0.06-64), clindamycin (0.06-64), levofloxacin (0.06-64), linezolid (0.03-32), vancomycin (0.015-16), teicoplanin (0.03-32), daptomycin (0.008-8), rifampicin (0.03-32), and trimethoprim/sulfamethoxazole (0.06/1.1875-64/1216). The concentration ranges for enterococci were ampicillin (0.03-32), levofloxacin (0.06-64), linezolid (0.008-8), vancomycin (0.06-64), teicoplanin (0.03-32), and daptomycin (0.008-8). Finally, the concentration ranges assayed for S. agalactiae for each antibiotic were ampicillin (0.004-4), erythromycin (0.06-64), clindamycin (0.06-64), levofloxacin (0.06-64), linezolid (0.008-8), vancomycin (0.008-8), and daptomycin (0.008-8).

The concentration ranges of PTS were 2-2048 mg/L in *Enterobacteriaceae*, nonfermenting gram-negative bacilli and *S. aureus*, and 4-4096 mg/L in *E. faecalis* and *S. agalactiae*. For PTSO, they were 2-2048 mg/L in *Enterobacteriaceae* and nonfermenting gram-negative bacilli and 0.125-128 mg/L in gram-positive cocci. Thus, the final concentration of polysorbate-80 in the wells was less than 1%.

The minimum inhibitory concentration (MIC) was defined as the lowest antibiotic concentration to completely inhibit the visible growth of a microorganism after overnight incubation and the isolates were considered to be susceptible, intermediate, or resistant, according to the recommendations of the CLSI [12]. A "susceptible" result indicates that the patient's organism should respond to therapy with that antibiotic using the dosage recommended normally for that type of infection and species. Conversely, a microorganism with a MIC interpreted as "resistant" should not be inhibited by the concentrations of the antibiotic achieved with the dosages normally used with that drug. An "intermediate" result indicates that a microorganism falls into a range of susceptibility in which the MIC approaches or exceeds the level of antibiotic that can ordinarily be achieved and for which clinical response is likely to be less than with a susceptible strain. MIC₅₀ and MIC₉₀ values were defined as the lowest concentration of the antibiotic at which 50 and 90% of the isolates were inhibited, respectively.

For minimum bactericidal concentration (MBC) testing, 100 μ l of broth from 1 to 5 wells containing no growth (which

showed no visible turbidity) was plated onto antibioticfree Columbia agar and incubated overnight at 37°C. The highest dilution that yielded no single bacterial colony on the agar plates was taken as MBC. Allium extracts were then considered as bacteriostatic or bactericidal depending on the MBC/MIC ratio which were, respectively, greater than 2 or between 2 and 1. MBC₅₀ and MBC₉₀ values were defined as the concentration of the antibiotic which kills 50 and 90% of the isolates, respectively.

Following the CLSI guidelines, we used the following strains as quality control in the procedures: *E. coli* ATCC 25922, *P. aeruginosa* ATCC 27853, *S. aureus* ATCC 29213, and *E. faecalis* ATCC 29212.

2.4. Statistical Analysis. Data analysis was performed using the software IBM SPSS Statistics v19. The Mann–Whitney U test was used to compare the distribution of MIC and MBC values of PTS and PTSO in the different groups of bacteria studied. A level of significance was considered with a p < 0.05.

3. Results

Tables 1 and 2 show the values (in mg/L) of the MIC_{50} , MIC_{90} , MBC_{50} , and MBC_{90} and percentages of resistance to the antibacterial agents tested of the 479 clinical isolates.

There was 59 ESBL-producing *Enterobacteriaceae* (42 *E. coli*, 12 *K. pneumoniae*, and 5 *K. oxytoca*). The presence of this resistance phenotype in 39.1% of *Enterobacteriaceae* was the main determinant of the high rates of resistance to beta-lactam antibiotics, whose range oscillated from 1.3% to meropenem (MIC₅₀ = 0.125 mg/L, MIC₉₀ = 1 mg/L) to 81.5% to cefuroxime (MIC₅₀ > 512 mg/L, MIC₉₀ > 512 mg/L).

ESBL-producing strains were more resistant to second to fourth-generation cephalosporins, such as cefuroxime $(MIC_{50} > 512 \text{ mg/L}, MIC_{90} > 512 \text{ mg/L}, 100\% \text{ resistant}),$ cefotaxime (MIC₅₀ = 128 mg/L, MIC₉₀ > 128 mg/L, and 100%resistant), ceftazidime (MIC₅₀ = 64 mg/L, MIC₉₀ = 256 mg/L, and 78.0% resistant), and cefepime (MIC $_{50}$ = 32 mg/L, MIC $_{90}$ = 128 mg/L, and 93.2% resistant) that combinations of betalactams with beta-lactamase inhibitors such as piperacillintazobactam (MIC₅₀ = 8/4 mg/L, MIC₉₀ = 256/4 mg/L, and 30.5% resistant) and amoxicillin/clavulanate (MIC₅₀ = 16/8mg/L, MIC₉₀ > 256/128 mg/L, and 52.5% resistant) or to carbapenems such as imipenem (MIC₅₀ = 0.5 mg/L, MIC₉₀ = 1 mg/L, and 100% susceptible) or meropenem (MIC₅₀ = 0.125mg/L, $MIC_{90} = 1$ mg/L, and 100% susceptible). Nevertheless, the absence of ESBL in Salmonella spp. and Yersinia spp. explains the lower number of isolates resistant to betalactam antibiotics in this group of enterobacteria (range 0-28.1%). Finally, in case of bacteria such as *Enterobacter* spp., Proteus spp., or Providencia spp., among others (remaining enterobacteria group in Table 1), high rates of resistance to beta-lactams were observed: 16.7% to meropenem (MIC₅₀ = 0.06 mg/L, MIC₉₀ = 1 mg/L) and 100% to cefuroxime $(MIC_{50} > 512 \text{ mg/L}, MIC_{90} > 512 \text{ mg/L}).$

Among the aminoglycosides, amikacin was the antibiotic with a higher rate of activity against enterobacteria (MIC_{50} = 16 mg/L, MIC_{90} > 512 mg/L, and 29.1% resistant), against 35.8% resistant to gentamicin (MIC_{50} = 4 mg/L, MIC_{90} = 128

TABLE 1: Activity *in vitro* of PTS, PTSO, and others antibacterial agents against gram-negative organisms.

Organisms (number of isolates)	MIC_{50}	MIC_{90}	MBC_{50}	MBC_{90}	% of resistant isolates
Futerohacteriaceae (n=151)	(III IIIg/L)	(III IIIg/ L)	(III IIIg/ L)	(III IIIg/ L)	
Amoxicillin/clavulanate	32/16	256/128	64/32	256/128	59.6
Piperacillin/tazobactam	8/4	256/4	32/4	512/4	31.8
Cefuroxime	>512	>512	>512	>512, 1	81.5
Cefoxitin	8	256	64	512	37.1
Cefotaxime	64	>128	128	>128	775
Ceftazidime	16	256	64	512	58.3
Cefenime	8	128	32	256	65.6
Iminenem	1	120	2	16	2.0
Meropenem	0.125	1	0.25	4	13
Gentamicin	0.125	1	32	-128	35.8
Tobramycin	4	128	32	>128	12 A
Amikacin	16	>512	52	>512	20.1
Ciprofloyacin	10 64	>312	128	>312	2 7 .1
Trimetheorim (culter atheware)	2/28	>120	64/1216	>120	52.0
Nitrofumeto/n	2/38	>04/1210	04/1210	>04/1210	55.0
DTS	52 256	250	128	512	43.0
PTS DTSO	250	512	200	512	-
FISU Eachanishia cali (n=69)	04	128	128	512	-
Amoviaillin (alexador ato	16/0	256/129	64/22	> 256/129	55.0
Din ana cillin (taraha stam	10/8	250/128	04/32	>256/128	55.9 26 F
Piperaciiin/tazobactam	8/4	128/4	32/4	256/4	26.5
Cefuroxime	>512	>512	>512	>512	95.6
Ceroxitin	8	128	32	256	33.8
Cetotaxime	128	>128	>128	>128	94.1
Ceftazidime	32	256	64	>512	75.0
Cetepime	16	128	64	256	80.9
Imipenem	0.5	1	2	4	0.0
Meropenem	0.06	I	0.125	4	0.0
Gentamicin	4	64	16	128	30.9
Tobramycin	4	64	16	128	33.8
Amikacin	8	32	32	128	14.7
Ciprofloxacin	64	128	128	>128	73.5
Trimethoprim/sulfamethoxazole	>64/1216	>64/1216	>64/1216	>64/1216	61.8
Nitrofurantoin	32	64	64	256	14.7
PTS	128	256	256	512	-
PTSO	64	128	128	512	-
Klebsiella spp. (n=39)					
Amoxicillin/clavulanate	32/16	64/32	64/32	256/128	82.1
Piperacillin/tazobactam	32/4	512/4	64/4	>512/4	59.0
Cefuroxime	>512	>512	>512	>512	97.4
Cefoxitin	32	512	64	>512	53.8
Cefotaxime	64	>128	128	>128	97.4
Ceftazidime	64	256	128	512	76.9
Cefepime	16	64	32	256	79.5
Imipenem	1	1	2	4	0.0
Meropenem	0.06	0.25	0.125	1	0.0
Gentamicin	64	>128	64	>128	66.7
Tobramycin	32	>128	32	>128	76.9

		TABLE 1: Contin	ued.		
Organisms (number of isolates)	MIC_{50}	MIC_{90}	MBC_{50}	MBC_{90}	% of resistant isolates
Amikacin	(III IIIg/L) 64	>512	128	>512	53.8
Ciprofloxacin	128	>128	>128	>128	87.2
Trimethoprim/sulfamethoyazole	>64/1216	>64/1216	>64/1216	>64/1216	79.5
Nitrofurantoin	64	128	128	256	61.5
PTS	256	512	256	512	01.5
PTSO	128	256	128	512	_
FSBI-producers (n=59)	120	250	120	512	
Amovicillin/clavulanate	16/8	>256/128	61/32	>256/128	52.5
Piperacillin/tazobactam	8/4	256/4	32/4	512/4	30.5
Cefurovime	≥512	>512	>512	>512/4	100
Cefovitin	8	64	32	128	25.4
Cefotavime	128	>128	>128	>120	100
Coffazidima	64	256	256	>5120	78.0
Cefenime	32	128	64	>256	93.2
Iminonom	0.5	120	2	/230	95.2
Moronom	0.5	1	0.25	4	0.0
Contomicin	0.125	1	0.25	4	0.0
Tahaanaa	4	128	16	>128	37.3
	4	128	32	>128	47.5
Amikacin Cinne flame sin	16	128	64	128	25.4
Trim ath annim (sulfam ath averals	04	>128	128	>128	/4.0
Nite (>64/1216	>64/1216	>64/1216	>64/1216	62.7
Nitrofurantoin	32 129	128	64	256	32.2
P1S	128	256	256	512	-
PISO	64	128	128	512	-
Saimonella spp. and Yersinia spp. (n=	:32)	256/120	(1/22	256/120	20.1
Amoxicillin/clavulanate	8/4	256/128	64/32	256/128	28.1
Piperacillin/tazobactam	2/4	128/4	16/4	128/4	12.5
Ceturoxime	4	>512	32	>512	25.0
Cefoxitin	8	64	64	128	12.5
Cefotaxime	1	1	8	64	12.5
Ceftazidime	4	4	16	64	0.0
Cetepime	2	32	16	64	0.0
Imipenem	1	1	16	16	0.0
Meropenem	1	1	4	8	0.0
Gentamicin	4	4	32	32	3.1
Tobramycin	4	4	32	32	6.3
Amikacin	16	128	128	256	34.4
Ciprofloxacin	1	128	8	128	37.5
Trimethoprim/sulfamethoxazole	2/38	2/38	16/304	>64/1216	3.1
Nitrofurantoin	256	512	512	1024	65.6
PTS	256	256	256	512	-
PTSO	64	128	64	128	-
Remaining enterobacteria (n=12)					
Amoxicillin/clavulanate	64/32	128/64	256/128	256/128	91.7
Piperacillin/tazobactam	4/4	64/4	8/4	256/4	25.0
Cefuroxime	>512	>512	>512	>512	100
Cefoxitin	256	>512	256	>512	66.7
Cefotaxime	64	>128	128	>128	91.7
Ceftazidime	8	128	32	512	58.3

Organisms (number of isolates) MIC ₉₉ (in mg/L) MIC ₉₉ (in mg/L) MBC ₉₁ (in mg/L) MBC ₉₁ (in mg/L) MBC ₉₉ (in mg/L) MBC ₉₁ (in mg/			TABLE 1: Contin	ued.		
Cefepime 8 64 8 236 66.7 Impenem 1 4 2 16 25.0 Meropenem 0.06 1 0.125 4 16.7 Gentamicin 4 32 32 >128 58.3 Amilacin 4 256 8 >512 16.7 Ciprofioxacin 1 >128 8 >128 50.0 Tirmethoprim/sulfamethoxazole 2/38 >64/1216 16/304 >64/1/216 50.0 Nitofurnatoin 64 128 128 256 - - PTS 128 256 126 256 - - PirosO 64 128 128 256 42.6 1mipenem 16 128 32 >128 32.5 Marcementing gram-negative badili (n=60) 128 32 >128 32.5 56 19.7 Opromoxia 8 128 32 >128 32.5 <	Organisms (number of isolates)	MIC ₅₀ (in mg/L)	MIC ₉₀ (in mg/L)	MBC ₅₀ (in mg/L)	MBC ₉₀ (in mg/L)	% of resistant isolates
Impenen I 4 2 16 25. Meropenem 0.06 1 0.125 4 167 Gentamicin 4 32 32 16 5128 58.3 Amikacin 1 5128 8 512 16.7 Ciprofloxacin 1 5128 8 512 50.0 Trimethoprim/sulfamethoxazole 2/38 >64/1216 16/304 >64/1216 50.0 Nitrofurantoin 64 128 256 256 256 - Nitrofurantoin 16/4 124 128 1024 32.8 - Cefapine 8 32 64 128 32.8 - - Cefapine 8 32 64 128 32.8 - - - Meropenen 16 128 32 128 32.9 - 128 - - Meropenen 16 128 32 128	Cefepime	8	64	8	256	66.7
Mcopenen 0.06 1 0.125 4 167 Gentamicin 4 32 32 >128 500 Tobramycin 4 32 32 >128 58.3 Anilacin 4 256 8 >512 16.7 Ciprofloxacin 1 178 184 >544/1216 16/304 >64/1216 50.0 Nitrofurantoin 64 >1024 128 >1024 8.3 32 PTS 128 256 2.6 - - 7 7 9.4 128 128 2.64 52.5 7 7 7 128 2.25 4 32 2.28 2.28 2.25 128 32 2.128 32.5	Imipenem	1	4	2	16	25.0
Gentamicin 4 32 32 >128 900 Tobramycin 8 32 16 >128 58.3 Amilacin 1 >128 8 >128 500 Tirmethoprin/sufamethoxazole 2/38 >64/1216 16/30 ><64/1216	Meropenem	0.06	1	0.125	4	16.7
Tobramycin 8 32 16 >128 8.8.3 Anilkacin 4 256 8 >512 167 Ciprofloxacin 1 >128 8 >512 500 Nitrofurantoin 64 >1024 128 >1024 8.3 PTS 128 256 256 256 - PTSO 64 512/4 128/ 526 256 Profermetting gram-negative bacilli (n=0) Noterimetting gram-negative bacilli (n=0) 32 64 512/4 34.4 Ceftazidime 8 128 64 512/4 32.8 52.5 Marpenen 16 128 32 5128 33.2 52.5 Meropenem 4 642 16 128 52.5 53.3 Gentamicin 4 5128 32 52.8 59.0 53.3 Tobramycin 32 5128 64 5128 53.5 55.7 Giprofloxacin 3	Gentamicin	4	32	32	>128	50.0
Analkacin 4 256 8 >512 16.7 Ciproloxacin 1 >128 8 >128 56.0 Nitrofurantoin 64 >1024 128 >1024 83.3 PTS 128 256 256 256 - Norterenting gram-negative bacilli (n=6) 128 256 - - Piperacilla/tazobactam 16/4 512/4 128/4 512/4 32.8 Cefepinie 8 32 64 256 42.6 Imigenem 16 128 32 >128 32.5 Gentamicin 4 >128 32 >128 32.9 Amikacin 8 128 32 >128 32.9 Ciproloxacin 32 >128 44 >128 32.9 Pirotamitinin 4 >128 32 256 197 Ciproloxacin 32 >128 64 >128 32.5 Pirota 16/	Tobramycin	8	32	16	>128	58.3
Ciprofloxacin 1 >128 8 >128 >640 Trimethoprim/sulfamethoxazole 2/38 >64/1216 16/304 >640/1216 50.0 Nitrofurantoin 128 >1024 128 >1024 83.3 PTS 128 256 256 256 - Nonfermenting gram-negative bacilli (n=0) 128 128 126 42.6 Caftazidime 8 128 64 512.4 32.4 Caftazidime 8 128 32 >128 32.5 Caftazidime 4 64 16 128 32.5 Gentamicin 4 128 32 >128 32.5 Gorbamycin 32 128 64 >128 32.5 Ciprofloxacin 32 128 64 >128 59.0 PTS 1024 2048 1024 2048 - PTSO 256 1024 264 128 32.5 Caftazi	Amikacin	4	256	8	>512	16.7
Trimethoprim/sulfamethoxazole 2/38 >64/1216 16/304 >64/1216 500 Nitrofurantoin 64 >1024 128 >1026 256 256 - PTS 64 128 128 256 256 - Nonfermenting gram-negative bacilli (net) 128 128 128 512/4 34.4 Cefegime 8 128 64 512/4 34.4 Cefegime 8 32 64 512 32.8 Gentamicin 16 128 32 >128 32.5 Gentamicin 4 5128 32 >128 39.3 Tobramycin 4 5128 32 218 39.3 Tobramycin 32 5128 64 5128 39.3 Tobramycin 32 5128 64 5128 39.0 PTS 1024 2048 1024 2048 - PTSO 1024 2048 1024 <	Ciprofloxacin	1	>128	8	>128	50.0
Nitrofurantoin 6.4 >1024 128 >024 256 2.56 . PTS 6.4 128 128 2.56 . . Prencillin/tazobactam 16/4 512/4 128/4 512/4 34.4 Ceftazidine 8 128 64 512 32.8 Cefepime 8 32 64 256 42.6 Inipenem 16 128 32 >128 32.7 Gentamicin 4 >128 32 >128 39.3 Tobranycin 4 >128 32 >128 39.3 Ciprofloxacin 32 >128 64 >128 39.3 PTS 1024 2048 124 204 39.7 PTS 1024 2048 124 2048 - PTS 1024 2048 128 32.5 128 Cefeprime 16 128 32 128 32.5	Trimethoprim/sulfamethoxazole	2/38	>64/1216	16/304	>64/1216	50.0
PTS 128 256 256 256 256 - PTSO 64 128 128 256 - Nonfermenting gram-negative bacill (n=0) 34.4 34.4 Cefazidime 8 128 64 512 32.8 Cefopime 8 32 64 525 52.5 Meropenem 16 128 32 5128 52.5 Gentamicin 4 64 16 128 52.5 Gentamicin 4 5128 32 5128 59.0 Tobranycin 32 512 2048 PTS 1024 2048 1.2 2048 PTSO 256 1024 2048 250 Perdomonas acreginosa (n=40) 64 128 128 57.5 Cefazidime 8 32 64 128 57.5 Cefazidime 8 32 128 57.5 <td>Nitrofurantoin</td> <td>64</td> <td>>1024</td> <td>128</td> <td>>1024</td> <td>83.3</td>	Nitrofurantoin	64	>1024	128	>1024	83.3
PTSO 64 128 128 256 - Nonfermenting gram-negative bacilli (n=61) 1 128 512/4 512/4 34.4 Cefazidime 8 128 64 512 32.8 Cefepine 8 32 64 526 42.6 Innipenem 16 128 32 5128 52.5 Meropenem 4 64 16 128 32. Tobramycin 4 5128 32 5128 64 Amilacin 8 128 32 256 19.7 Ciprofloxacin 32 5128 64 5128 59.0 PTSO 1024 2048 1024 2048 - PTSO 256 1024 512 2048 - PTSO 256 1024 512 2048 - Cefepine 8 62 64 128 32.5 Inipenem 16/4 256/4 128/4 32.5 Cefepine 8 32 64 128 32.5 Cefepine 8 32 64 128 32.5 Cefepine 16 128 32.5 55.5<	PTS	128	256	256	256	-
Nonfermenting gram-negative bacilli (n=6)Piperacillin/tazobactam16/4512/4128/4512/432.8Cefepine8326425642.6Inipenem1612832>12852.5Meropenem4641612839.3Tobramycin4>12832>12839.3Tobramycin4>12832>12859.0Ciprofloxacin32>12864>12859.0PTS1024204810242048-PTSO25610245122048-Peadomas acruginosa (n=40)PP256/4256/4256/4Cefegine16/4256/4128/4256/425.0Cefegine8326451232.5Inipenem16/4256/4128/425.627.5Cefegine8326451232.5Inipenem16/42583212857.5Gentamicin421816>12837.5Tobranycin412816>12837.5Ciprofloxacin32>1286451224.5PTS102420482048-15.5Gentamicin8323212862.5PTS102421826421824.5Cefepine65122048-PTSO5121024<	PTSO	64	128	128	256	-
Piperacillin/tazobactam 16/4 512/4 128/4 512/4 34.4 Cefepime 8 32 64 512 32.8 Cefepime 8 32 64 512 32.8 Imipenem 16 128 32 >128 32.5 Meropenem 4 64 16 128 32.2 Amikacin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 >128 39.3 PTS 1024 2048 122 2048 - PTS 1024 2048 122 2048 - PTSO 256 1024 128/4 256/4 25.0 Cefepime 8 32 64 128 32.5 Imipenem 16 128 32 128 57.5 Cefepime 8 32 <td< td=""><td>Nonfermenting gram-negative bacill</td><td>i (n=61)</td><td></td><td></td><td></td><td></td></td<>	Nonfermenting gram-negative bacill	i (n=61)				
Ceftazidime 8 128 64 512 32.8 Cefepine 8 32 64 256 42.6 Imipenem 16 128 32 >128 52.5 Gentamicin 4 64 16 128 32.5 Gentamicin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 256 197 Giprofloxacin 32 >128 64 >128 59.0 PTS 1024 2048 128 59.0 127 PTSO 1024 2048 128 25.7 59.0 PTSO 1024 2048 128 32.5 59.0 Cefazidime 8 64 64 22.8 25.0 Cefazidime 8 32 64 128 32.5 Imipenem 16 128 32.2 128 125.5 Gentamicin 4 64 32	Piperacillin/tazobactam	16/4	512/4	128/4	512/4	34.4
Cefepime 8 32 64 256 42.6 Imipenem 16 128 32 >128 52.5 Meropenem 4 64 16 128 32.5 Gentamicin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 >128 39.3 Ciprofloxacin 8 128 32 >128 59.0 PTS 024 2048 1024 2048 - PTSO 256 1024 2048 - - Precolimonas aeruginosa (m=40) 256/4 128/4 256/4 25.0 Cefepine 8 32 64 128 32.5 Imipenem 16/4 128 32 128 37.5 Cefepine 8 32 64 128 37.5 Gentamicin 4 428 16 >128 37.5 Gentamicin 4 128 128	Ceftazidime	8	128	64	512	32.8
Imipenem 16 128 32 >128 52.5 Meropenem 4 64 16 128 32.5 Gentamicin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 >128 39.3 Amikacin 8 128 32 256 19.7 Ciprofloxacin 32 >128 64 >128 59.0 PTS 1024 2048 1024 2048 - PTSO 256 1024 512 2048 - Preadomonas aeruginosa (n=40) 526 1024 254 25.0 25.6 Cefazidime 8 64 64 22.6 25.6 25.5 Imipenem 16 128 32 128 57.5 35.5 Gentamicin 4 54.8 32 12.8 37.5 15.2 Imipenem 16 128 32.2 128 64 512	Cefepime	8	32	64	256	42.6
Neropenem4641612852.5Gentamicin4>12832>12839.3Tobramycin4>12832>12839.3Amikacin81283225619.7Ciprofloxacin32>12864>12859.0PTS1024204810242048-PTSO25610245122048-Perdomas aeruginosa (n=40)256/4128/4256/425.0Ceftazidime8646425627.5Cefepime161283212857.5Meropenem4643212857.5Gentamicin4>12816>12837.5Tobranycin412816>12812.5Ciprofloxacin32>12864.321.512.5Gentamicin412816>12812.5TSO102420482048PTSO102420482048PTSO1024102420482048-PTSO102420482048PTSO102420482048PTSO102420482048PTSO102420482048PTSO102420482048PTSO102420482048- <t< td=""><td>Imipenem</td><td>16</td><td>128</td><td>32</td><td>>128</td><td>52.5</td></t<>	Imipenem	16	128	32	>128	52.5
Gentamicin 4 >128 32 >128 39.3 Tobramycin 4 >128 32 >128 279 Amikacin 8 128 32 >128 279 Amikacin 8 128 32 >128 64 >128 59.0 PTS 1024 2048 1024 2048 - - Predomonas aeruginosa (n=40) 1024 2048 1024 2048 - Pendomonas aeruginosa (n=40) 8 64 64 256 27.5 Cefeptime 8 32 64 128 32.5 Imipenem 16 128 32 128 57.5 Gentamicin 4 128 16 >128 37.5 Tobramycin 4 128 16 >128 12.5 Ciprofloxacin 32 >128 64 >128 62.5 PTS 1024 1024 128 2048 - </td <td>Meropenem</td> <td>4</td> <td>64</td> <td>16</td> <td>128</td> <td>52.5</td>	Meropenem	4	64	16	128	52.5
Tobramycin4>12832>128279Amikacin812832256197Ciprofloxacin32>12864>12859.0PTS1024204810242048-PTSO25610245122048-Preadimonas aeruginosa (n=40) $r128/4256/425.0Ceftazidime8646425627.5Ceftazidime8326412832.5Imipenem161283212857.5Gentamicin4643212837.5Tobramycin412816>12837.5Gentamicin8323212812.5Ciprofloxacin32>12864>12812.5PTS1024204820482048-PTS1024204820482048-PTSO51210251220451242.9Ciprofloxacin32>1286451242.9PTSO5121024512.4512.442.9Cefazidime66412825661.9Imipenem128/4512.4512.442.9Qerofloxacin4>1283231242.9Meropenem2168312.842.9Meropenem216832.842.9Meropenem216$	Gentamicin	4	>128	32	>128	39.3
Amikacin812832256197Ciprofloxacin32>12864>12859.0PTS1024204810242048-PTSO251024204810242048-PTSO251024128/42048-Perdomonas aeruginosa (n=40)256/4128/4256/425.0Ceftazidime8646425627.5-Ceftazidime8326412832.5-Imipenem161283212857.5-Gentamicin4512816>12837.5-Tobranycin412816>12837.5-Tobranycin412816>12817.5-PTS1024204820482048PTS1024204820482048PTS1024204820482048PTS1024204820482048PTS10242124512/4512/452.4-Ceftazidime81286412822.4-PTS12812816>12842.9-Meropenem2168212842.9-Giprofexacin4>12832212842.9Gentamicin1664128 <td< td=""><td>Tobramycin</td><td>4</td><td>>128</td><td>32</td><td>>128</td><td>27.9</td></td<>	Tobramycin	4	>128	32	>128	27.9
Ciprofloxacin 32 >128 64 >128 910 PTS 1024 2048 1024 2048 - PTSO 256 1024 512 2048 - PrsO 256 1024 512 2048 - Preacillin/tazobactam 16/4 256/4 128/4 256/4 25.0 Cefeazidime 8 64 64 256 27.5 Cefeazidime 8 64 64 256 27.5 Cefeazidime 8 32 64 128 32.5 Imipenem 16 128 32 128 57.5 Gentamicin 4 128 16 >128 37.5 Tobramycin 4 128 16 >128 62.5 PTS 1024 2048 2048 -4 24.9 PTSO 512 1024 2048 2048 -4 PTSO 128 128 <	Amikacin	8	128	32	256	19.7
PTS10242048102420481024PTSO25610245122048-Perdumonas aeruginosa (n=40) $=$ $=$ $=$ $=$ Piperacillin/tazobactam16/4256/4128/4256/425.0Ceftazidime8646425627.5Ceforpine8326412832.5Imipenem161283212857.5Meropenem4643212857.5Gentamicin412816>12837.5Tobranycin412816>12817.5Amikacin8323212862.5PTS1024204820482048-PTSO5121024204820482048-PTSO5121024204820482048-Preacillin/tazobactam128/4512/4512/452.4Cefepime166412825661.9Imipenem2>1286451242.9Cefepime166412825661.9Imipenem2>1286451242.9Queropenem21683232.4Preso166412825661.9Imipenem166412832.432.4Queropenem216832.4Queropenem21	Ciprofloxacin	32	>128	64	>128	59.0
PTSO 256 1024 512 2048 - Pseudomonas aeruginosa (n=40) 1 256/4 128/4 256/4 25.0 Piperacillin/tazobactam 16/4 256/4 256/6 27.5 Cefepine 8 64 64 256 27.5 Cefepine 8 32 64 128 32.5 Imipenem 16 128 32 128 57.5 Meropenem 4 64 32 128 57.5 Tobramycin 4 128 16 >128 37.5 Amikacin 8 32 32 128 16.5 PTS 1024 1024 2048 2048 -2.5 PTS 1024 2048 2048 -2.6 -2.7 PTSO 12 1024 2048 2048 -2.1 Cefrazidime 128/4 512/4 512/4 2.2.4 -2.1 Cefepine 16 64 <td>PTS</td> <td>1024</td> <td>2048</td> <td>1024</td> <td>2048</td> <td>-</td>	PTS	1024	2048	1024	2048	-
PeriodInternational problemInternational problemPiperacillin/tazobactam16/4256/4128/4256/425.0Ceftazidime8646425627.5Cefepime8326412832.5Imipenem161283212857.5Meropenem4643212837.5Tobramycin412816>12837.5Tobramycin412816>12817.5Amikacin8323212862.5PTS1024204820482048-PTSO51210245122048-Peracillin/tazobactam128/4512/4512/452.4Ceftazidime81286451242.9Ceftazidime166412825661.9Imipenem128/4512/4512/452.452.4Ceftazidime166412825661.9Imipenem2168>12842.9Cefepime166412825661.9Imipenem2168>12842.9Cefopine168>12842.9Gentamicin4>12832>12842.9Meropenem2168>12842.9Tobramycin4>12832>12842.9Tobramycin4>128 <td< td=""><td>PTSO</td><td>256</td><td>1024</td><td>512</td><td>2048</td><td>-</td></td<>	PTSO	256	1024	512	2048	-
Piperacillin/tazobactam16/4256/4128/4256/425.0Ceftazidime8646425627.5Cefepime8326412832.5Imipenem161283212857.5Meropenem4643212857.5Gentamicin4>12816>12837.5Tobramycin412816>12837.5Mikacin8323212812.5Ciprofloxacin32>12864>12862.5PTS1024204820482048-PTSO51210245122048-Remaining nonfermenting gram-negative bacilli (n=21)7724.4Piperacillin/tazobactam128/4512/4512/452.4Cefepime166412825661.9Imipenem2>12816>12842.9Meropenem2168>12842.9Meropenem4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Meropenem2168>12842.9Meropenem4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobr	Pseudomonas aeruginosa (n=40)					
Ceftazidime8646425627.5Ceftazidime8326412832.5Imipenem161283212857.5Meropenem4643212857.5Gentamicin4>12816>12837.5Tobramycin412816>12817.5Amikacin8323212862.5PTS1024204820482048-PTSO51210245122048-Remaining nonfermenting gram-negative bacilli (n=21)71.221.224.2Ceftazidime81286451242.9Ceftazidime166412825661.9Imipenem2>12816>12842.9Ceftazidime166412825661.9Imipenem2>12816>12842.9Meropenem2168>12842.9Gentamicin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12833.3Ciprofloxacin <td< td=""><td>Piperacillin/tazobactam</td><td>16/4</td><td>256/4</td><td>128/4</td><td>256/4</td><td>25.0</td></td<>	Piperacillin/tazobactam	16/4	256/4	128/4	256/4	25.0
Cefepime8326412832.5Imipenem161283212857.5Meropenem4643212857.5Gentamicin4>12816>12837.5Tobramycin412816>12817.5Amikacin8323212862.5PTS1024204820482048-PTSO512102420482048-Remaining nonfermenting gram-negative bacilli (n=21)712512/4512/452.4Ceftazidime128/4512/4512/452.424.9Ceftazidime166412825661.9Imipenem2>12816>12842.9Meropenem2168>12842.9Meropenem2168>12842.9Meropenem4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12833.3Ciprofloxacin4>12832>12832.Tobramycin4>12832>12832.Tobramycin4>12832>12832.Tobramycin4>12832>12832.P	Ceftazidime	8	64	64	256	27.5
Imipenem 16 128 32 128 575 Meropenem 4 64 32 128 575 Gentamicin 4 >128 16 >128 375 Tobramycin 4 128 16 >128 375 Amikacin 8 32 32 128 175 Amikacin 8 32 32 128 62.5 PTS 1024 2048 2048 2048 - PTSO 512 1024 512 2048 - Peracillin/tazobactam 128/4 512/4 512/4 52.4 Cefazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 </td <td>Cefepime</td> <td>8</td> <td>32</td> <td>64</td> <td>128</td> <td>32.5</td>	Cefepime	8	32	64	128	32.5
Meropenem4643212857.5Gentamicin4>12816>12837.5Tobramycin412816>12817.5Amikacin8323212812.5Ciprofloxacin32>12864>12862.5PTS1024204820482048-PTSO51210245122048-Remaining nonfermenting gram-negative bacilli (n=21)Piperacillin/tazobactam128/4512/4512/452.4Ceftazidime81286451242.9Cefepime166412825661.9Imipenem2>12816>12842.9Gentamicin4>12832>12842.9Gentamicin4>12832>12847.6Amikacin162566425633.3Ciprofloxacin4>12832>12852.4FTS128512128512-PTSO642566425633.3	Imipenem	16	128	32	128	57.5
Gentamicin4>12816>12837.5Tobramycin412816>12837.5Amikacin8323212812.5Ciprofloxacin32>12864>12862.5PTS1024204820482048-PTSO51210245122048-Remaining nonfermenting gram-negative bacilli (n=21)Piperacillin/tazobactam128/4512/4512/4>512/452.4Ceftazidime81286451242.9Cefepime166412825661.9Imipenem2>12816>12842.9Gentamicin4>12832>12842.9Gentamicin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12847.6Amikacin162566425633.3Ciprofloxacin4>12832>12852.4PTS128512128512-PTSO128512128512-Tobramycin4>12832>12852.4PTSO128512128512-PTSO128512128512-PTSO128512	Meropenem	4	64	32	128	57.5
Tobramycin412816>128175Amikacin8323212812.5Ciprofloxacin32>12864>12862.5PTS1024204820482048-PTSO51210245122048-Remaining nonfermenting gram-negative bacilli (n=21)Piperacillin/tazobactam128/4512/4512/452.4Ceftazidime81286451242.9Cefepime166412825661.9Imipenem2>12816>12842.9Gentamicin4>12832>12842.9Gentamicin4>12832>12842.9Tobramycin4>12832>12842.9Tobramycin4>12832>12847.6Amikacin162566425633.3Ciprofloxacin4>12832>12852.4PTS128512128512-PTS128512128512-PTS128512128512-PTS128512128512-PTS128512128512-PTS5664256512-PTS52664256512-PTS51252664256-	Gentamicin	4	>128	16	>128	37.5
Amikacin 8 32 32 128 12.5 Ciprofloxacin 32 >128 64 >128 62.5 PTS 1024 2048 2048 2048 - PTSO 512 1024 512 2048 - Remaining nonfermenting gram-negative bacilli (n=21) 512/4 >512/4 52.4 Ceftazidime 8 128 64 512 42.9 Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Gentamicin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 42.9 Tobramycin 4 >128 <td>Tobramycin</td> <td>4</td> <td>128</td> <td>16</td> <td>>128</td> <td>17.5</td>	Tobramycin	4	128	16	>128	17.5
Ciprofloxacin 32 >128 64 >128 62.5 PTS 1024 2048 2048 2048 62.5 PTSO 512 1024 512 2048 - Remaining nonfermenting gram-negative bacilli (n=21) 512/4 512/4 551/4 52.4 Piperacillin/tazobactam 128/4 512/4 512/4 551/4 52.4 Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin <td>Amikacin</td> <td>8</td> <td>32</td> <td>32</td> <td>128</td> <td>12.5</td>	Amikacin	8	32	32	128	12.5
PTS 1024 2048 2048 2048 2048 - PTSO 512 1024 512 2048 - Remaining nonfermenting gram-negative bacilli (n=21) - - - - Piperacillin/tazobactam 128/4 512/4 512/4 >512/4 52.4 Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64<	Ciprofloxacin	32	>128	64	>128	62.5
PTSO 512 1024 512 2048 - Remaining nonfermenting gram-negative bacilli (n=21) Piperacillin/tazobactam 128/4 512/4 512/4 >512/4 52.4 Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	PTS	1024	2048	2048	2048	-
Remaining nonfermenting gram-negative bacilli (n=21) Number of the field of	PTSO	512	1024	512	2048	-
Piperacillin/tazobactam 128/4 512/4 512/4 5512/4 52.4 Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 - <td>Remaining nonfermenting gram-neg</td> <td>ative bacilli (n=2</td> <td>1)</td> <td>012</td> <td>2010</td> <td></td>	Remaining nonfermenting gram-neg	ative bacilli (n=2	1)	012	2010	
Ceftazidime 8 128 64 512 42.9 Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 12 - PTSO 64 256 64 256 -	Piperacillin/tazobactam	128/4	-, 512/4	512/4	>512/4	52.4
Cefepime 16 64 128 256 61.9 Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 -	Ceftazidime	8	128	64	512	42.9
Imipenem 2 >128 16 >128 42.9 Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 42.9 Gentamicin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Cefenime	16	64	128	256	61.9
Important Important Important Important Important Important Meropenem 2 16 8 >128 42.9 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Imipenem	2	>128	16	>128	42.9
Millipentin 1 10 6 7120 7120 Gentamicin 4 >128 32 >128 42.9 Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Meropenem	2	16	8	>128	42.9
Tobramycin 4 >128 32 >128 47.6 Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Gentamicin	4	>128	32	>120	42.9
Amikacin 16 256 64 256 33.3 Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Tobramycin	4	>128	32	>128	476
Ciprofloxacin 4 >128 32 >128 52.4 PTS 128 512 128 512 - PTSO 64 256 64 256 -	Amikacin	16	256	64	256	22.2
PTS 128 512 52 7126 52.4 PTSO 64 256 64 256 -	Ciprofloyacin	10 Д	×128	32	×128	50.5 52 A
PTSO 64 256 64 256 -	PTS	128	512	128	512	-
	PTSO	64	256	64	256	-

MIC: minimum inhibitory concentration; MBC: minimum bactericidal concentration; % of resistant isolates: percentages of isolates intermediate or resistant according to the criteria published by the CLSI (2016).

BioMed Research International

TABLE 2: Activity in vitro of PTS, PTSO, and others antibacterial agents against gram-positive organisms.

Organisms (number of isolates)	MIC_{50}	MIC_{90}	MBC ₅₀	MBC_{90}	% of resistant isolates
Stathylococcus aureus methicillin-re	sistant (n=112)	(III IIIg/L)	(III IIIg/L)	(III IIIg/L)	
Gentamicin	4	256	16	>256	48.2
Tobramycin	64	>256	>256	>256	79.5
Ervthromycin	>64	>64	>64	>64	69.6
Clindamycin	>64	>64	>64	>64	49.1
Levofloxacin	8	32	32	>64	89.3
Linezolid	2.	4	4	8	0.0
Vancomycin	0.5	1	1	4	0.0
Teicoplanin	0.25	1	0.5	4	0.0
Daptomycin	0.25	0.5	0.5	2.	0.0
Rifampicin	<0.03	0.5	0.125	-	3.6
Trimethoprim/sulfamethoxazole	<0.06	0.5	0.5	2	3.6
PTS	<u>_</u> 0.00	128	512	1024	-
PTSO	8	8	32	64	-
Enterococcus faecalis (n=54)	0	Ū	52	01	
Ampicillin	1	2	2	8	0.0
Levoflovacin	32	2 64	>64	>64	100
Linezolid	2	2	4	8	0.0
Vancomycin	0.5	1	2	4	0.0
Teicoplanin	< 0.03	0.125	0.25	1	0.0
Daptomycin	20.05	4	4	8	0.0
PTS	128	128	2048	4096	-
PTSO	4	8	8	16	_
Streptococcus agalactiae (n=101)	Ŧ	0	0	10	
Ampicillin	0.06	0 125	0 125	0.5	0.0
Frythromycin	>64	>64	>64	>64	94.1
Clindamycin	>64	>64	>64	>64	85.1
Levofloxacin	0.5	1	2	8	6.9
Linezolid	1	2	2	4	0.0
Vancomycin	1	1	2	4	0.0
Daptomycin	0 125	0.5	0.5	2	0.0
PTS	64	128	512	2048	-
PTSO	4	8	8	16	-

MIC: minimum inhibitory concentration; MBC: minimum bactericidal concentration; % of resistant isolates: percentages of isolates intermediate or resistant according to the criteria published by the CLSI (2016).

mg/L) or 42.4% to tobramycin (MIC₅₀ = 4 mg/L, MIC₉₀ = 128 mg/L). The resistance to aminoglycosides was higher among *Klebsiella* spp. and the group "remaining enterobacteria" than *E. coli*, *Salmonella* spp., or *Yersinia* spp. In general, enterobacteria showed high resistance to fluoroquinolones (MIC₅₀ = 64 mg/L, MIC₉₀ > 128 mg/L, and 67.6% of resistant isolates to ciprofloxacin) and to trimethoprim-sulfamethoxazole (MIC₅₀ = 2/38 mg/L, MIC₉₀ > 64/1216 mg/L, and 53.0% resistant), except for *Salmonella* spp. and *Yersinia* spp., which showed the lowest rates (37.5% and 3.1% of resistant isolates to ciprofloxacin and trimethoprim-sulfamethoxazole, respectively). *E. coli* was the bacteria with a lower resistance to nitrofurantoin (MIC₅₀ = 32 mg/L, MIC₉₀ = 64 mg/L, and 14.7% resistant).

As previously mentioned, bacteria were selected for their detection of resistance to, at least, a group of antibiotics. However, a relevant characteristic of the 151 enterobacteria included in the study was the high frequency to coresistance to two or more of these groups (multidrug-resistant bacteria), as described in Table 3. Therefore, 74.0% of the isolates resistant to some beta-lactams antibiotics were also resistant to ciprofloxacin, 61.8% to trimethoprim-sulfamethoxazole, and 48.8% to some aminoglycoside. It should be noted that 22.8% of that resistant to beta-lactams was also resistant to all the other groups of antibiotics assayed.

The behaviour of PTS and PTSO against multidrugresistant enterobacteria was quite homogeneous, regardless the group analyzed (Table 1). The values of MIC_{50} and MIC_{90}

	IABLE J. MILAIYSIS UI CUI		roups of antibiotics.		
	Enterobacteria resistant to some beta-lactams antibiotics (n=123; 81.5%)	ESBL-producers enterobacteria (n=59; 39.1%)	Non ESBL-producers enterobacteria resistant to some beta-lactams antibiotics (n=64; 42.4%)	Non-fermenting gram-negative bacilli resistant to fluoroquinolones (n=36; 59.0%)	Methicillin-resistant Staphylococcus aureus (n=112)
Resistance to beta-lactams	1		1	75.0%	1
Resistance to aminoglycosides	48.8%	47.5%	50%	63.9%	79.5%
Resistance to fluoroquinolones	74.0%	74.6%	73.4%		89.3%
Resistance to TMX	61.8%	62.7%	60.9%		3.6%
Resistance to nitrofurantoin	41.5%	32.2%	50.0%		
Resistance to macrolides (erythromycin)	ı		1		69.6%
Resistance to lincosamides (clindamycin)	ı		1	·	49.1%
Resistance to rifampicin	ı	·	1		3.6%
Resistance to aminoglycosides & fluoroquinolones	46.3%	44.1%	50.0%		75.9%
Resistance to aminoglycosides & fluoroquinolones & TMX	41.5%	39.0%	45.3%	I	2.7%
Resistance to aminoglycosides & fluoroquinolones & TMX & nitrofurantoin	22.8%	22.0%	23.4%	I	ı
Resistance to beta-lactams & aminoglycosides	I		I	55.6%	
Resistance to aminoglycosides & fluoroquinolones & macrolides	ı	ı	I	I	60.7%
Resistance to aminoglycosides & fluoroquinolones & macrolides & lincosamides	·	·	ı	I	45.5%
TMX: Trimethoprim/sulfamethoxazole.					

of antibiotics.
groups
different
to
coresistance
of
Analysis
3:
TABLE

of PTS ranged from 128 to 256 mg/L and from 256 to 512 mg/L, while the MBC_{50} and MBC_{90} ranged from 256 mg/L and 256 to 512 mg/L, respectively. On the other hand, the values of MIC_{50} and MIC_{90} of PTSO ranged from 64 to 128 mg/L and 128 to 256 mg/L, while MBC_{50} y MBC_{90} ranged from 64 to 128 mg/L and from 128 to 512 mg/L, respectively. These data show the bactericidal activity of both compounds (MIC and MBC values were equal or differed in only one dilution) and indicate that PTSO was significantly more active than PTS against this group of bacteria (p<0.001).

Among the 61 nonfermenting gram-negative bacilli, the resistance to beta-lactams antibiotics ranged from 32.8% to ceftazidime (MIC₅₀ = 8 mg/L, MIC₉₀ = 128 mg/L) and 52.5% to imipenem (MIC₅₀ = 16 mg/L, MIC₉₀ = 128 mg/L) and meropenem (MIC₅₀ = 4 mg/L, MIC₉₀ = 64 mg/L). Carbapenems showed more activity against bacteria such as Acinetobacter spp., Aeromonas spp., and Stenotrophomonas spp. (MIC_{50} = 2 mg/L, MIC_{90} $\,>$ 128 mg/L, and 42.9% of isolates resistant to imipenem and $MIC_{50} = 2 \text{ mg/L}$, $MIC_{90} =$ 16 mg/L, and 42.9% of isolates resistant to meropenem), than against Pseudomonas spp. (MIC₅₀ = 16 mg/L, MIC₉₀ = 128 mg/L, and 57.5% of isolates resistant to imipenem and MIC_{50} = 4 mg/L, MIC₉₀ = 64 mg/L, and 57.5% of isolates resistant to meropenem). Among the aminoglycosides assayed, amikacin was the most active against both groups (MIC₅₀ = 8 mg/L, $MIC_{90} = 128$ mg/L, and 19.7% resistant). Finally, 59.0% of isolates were resistant to ciprofloxacin (MIC₅₀ = 32 mg/L, $MIC_{90} > 128 \text{ mg/L}$), which resulted in less active against P. aeruginosa isolates than against other bacteria of this group. As shown in Table 3, 75.0% of the isolates resistant to fluoroquinolones (ciprofloxacin) were also resistant to some beta-lactam antibiotic; 63.9% to some aminoglycoside and 55.6% showed resistance to these three groups of antibiotics.

Just as with the rest of antibiotics, when comparing the results obtained in P. aeruginosa with those obtained in others nonfermenting gram-negative bacilli, the behaviour, both of PTS and PTSO, was significantly different (Table 1). In the case of PTS, the results shown in *P. aeruginosa* were MIC_{50} = 1024 mg/L, MIC_{90} = 2048 mg/L, MBC_{50} = 2048 mg/L, and $MBC_{90} = 2048 \text{ mg/L}$, while in the rest of bacteria they showed more activity (MIC₅₀ = 128 mg/L, MIC₉₀ = 512 mg/L, $MBC_{50} = 128 \text{ mg/L}$, and $MBC_{90} = 512 \text{ mg/L}$) (p < 0.001). Likewise, the results for PTSO indicated less activity against *Pseudomonas* spp. (MIC₅₀ = 512 mg/L, MIC₉₀ = 1024 mg/L, $MBC_{50} = 512 \text{ mg/L}$, and $MBC_{90} = 2048 \text{ mg/L}$) than against the rest of isolates (MIC₅₀ = 64 mg/L, MIC₉₀ = 256 mg/L, $MBC_{50} = 64 \text{ mg/L}$, and $MBC_{90} = 256 \text{ mg/L}$) (p < 0.001). In any case, these data also indicate the bactericidal activity of both compounds, especially PTSO that showed significantly more activity than PTS (p < 0.001).

Concerning the gram-positive cocci, all the isolates were susceptible to vancomycin, teicoplanin (*S. agalactiae* was not tested), daptomycin, and linezolid. Besides, all the isolates of *E. faecalis* and *S. agalactiae* were also susceptible to ampicillin (Table 2).

All the isolates of *S. aureus* were resistant to methicillin (this was the selection criteria in this bacteria) and therefore to all beta-lactams antibiotics. High rates of resistance to fluoroquinolones (MIC₅₀ = 8 mg/L, MIC₉₀ = 32 mg/L, 89.3%

resistant to levofloxacin), to aminoglycosides (MIC₅₀ = 64mg/L, MIC₉₀ > 256 mg/L, 79.5% resistant to tobramycin), to macrolides (MIC₅₀ > 64 mg/L, MIC₉₀ > 64 mg/L, 69.6% resistant to erythromycin), or to lincosamides ($MIC_{50} > 64$ mg/L, MIC₉₀ > 64 mg/L, 49.1% resistant to clindamycin) were observed. In contrast, trimethoprim-sulfamethoxazole $(MIC_{50} < 0.06 \text{ mg/L}, MIC_{90} = 0.5 \text{ mg/L}, 3.6\% \text{ resistant})$ and rifampicin (MIC₅₀ < 0.03 mg/L, MIC₉₀ = 0.5 mg/L, and 3.6% resistant) showed the lowest rates of resistance. The 75.9% of these bacteria were resistant, both to aminoglycosides and fluoroquinolones, and 60.7% also showed resistance to macrolides and 45.5% also to clindamycin (Table 3). Finally, 100% of isolates of E. faecalis were resistant to levofloxacin $(MIC_{50} = 32 \text{ mg/L}, MIC_{90} = 64 \text{ mg/L})$ and resistance to any other antibiotic was not associated, whereas 86 out of 101 isolates of S. agalactiae were resistant to erythromycin and clindamycin.

PTSO showed significantly more activity than PTS in the three groups of gram-positive bacteria tested (p < 0.001, in all cases) and the values for MIC₅₀, MIC₉₀, MBC₅₀, and MBC₉₀ were, for both compounds, lower than those obtained against gram-negative bacteria (Table 2). However, MIC and MBC values in gram-positive bacteria differed significantly, especially for PTS (more than 2 dilutions), which indicates that these compounds could have a bacteriostatic but not a bactericidal effect against these bacteria at least at low concentrations.

4. Discussion

Organosulfur compounds obtained from Allium spp. such as PTS and PTSO have been proposed as an effective alternative to antibiotics to improve animal performance and prevent gastrointestinal disorders. This is due on the one hand to their greater stability in comparison to other natural compounds [13] and on the other hand to their activity against bacterial groups, such as Enterobacteriaceae, Staphylococcus spp., Enterococcus spp., Clostridium spp., Bacteroides spp., Lactobacillus spp., Bifidobacterium spp., or Campylobacter spp., among others [3, 4, 7]. Furthermore, it has been shown that feed supplementation with these compounds improves the digestion and absorption of nutrients in the gastrointestinal tract by modulating the intestinal microbiota and increases the villus height and mucosal thickness [7, 8]. Beyond its use in animals, it is possible that these molecules may as well be useful in the human clinical practice, due to the fact that alliaceous plants have been traditionally used for their antibacterial, antioxidant, and cardiovascular properties, as has been known for centuries [6].

To our knowledge, this is the first study to evaluate the activity of PTS and PTSO against a selection of gramnegative and gram-positive multiresistant bacteria isolated from human clinical samples. Antibiotic susceptibility tests were performed in accordance with the procedure outlined by CLSI in order to determine if a bacterium is susceptible or resistant to each of the antibiotic assayed. Although the cutoff points for PTS or PTSO are unknown, perform the assay under the same conditions as the other antibiotics allow us to make comparisons with them. Our results revealed that PTS showed moderate activity against *Enterobacteriaceae* with MIC_{50} (and MBC_{50}) and MIC_{90} (and MBC_{90}) values of 256-512 mg/L, while PTSO showed greater activity with MIC_{50} and MIC_{90} values of 64-128 mg/L and MBC_{50} and MBC_{90} values of 128-512 mg/L. These homogeneous results among the different groups of enterobacteria selected, regardless of the resistance shown to different antibiotics commonly used in clinical practice, reveal the bactericidal action of these compounds. According to these results, Ruiz et al. also proved a bactericidal effect against enterobacteria, such as *E. coli* and *Salmonella typhimurium* [3].

The activity against methicillin-resistant *S. aureus*, *E. faecalis*, and *S. agalactiae* was higher than that observed against enterobacteria, especially in the case of PTSO ($MIC_{50} = 8 \text{ mg/L}$, $MIC_{90} = 8 \text{ mg/L}$, $MBC_{50} = 32 \text{ mg/L}$, $MBC_{90} = 64 \text{ mg/L}$, in *S. aureus*; $MIC_{50} = 4 \text{ mg/L}$, $MIC_{90} = 8 \text{ mg/L}$, $MBC_{50} = 8 \text{ mg/L}$, and $MBC_{90} = 16 \text{ mg/L}$, in *E. faecalis* and *S. agalactiae*). The PTS activity against this group of bacteria was significantly lower, especially in the case of enterococci. Some authors have evaluated the potential of garlic allicin, a molecule structurally similar to PTS, to control oral pathogens, reporting inhibitory concentrations of 600 mg/L against *Streptococcus* spp. [14]. Other studies have reported a bacteriostatic effect of allicin against vancomycin resistant enterococci [15].

However, in contrast to the relatively good results obtained previously, both compounds showed lower activity against *P. aeruginosa* (MIC₅₀ = 1024 mg/L, MIC₉₀ = 2048 mg/L, MBC₅₀ = 2048 mg/L, MBC₅₀ = 2048 mg/L, for PTS; MIC₅₀ = 512 mg/L, MIC₉₀ = 1024 mg/L, MBC₅₀ = 512 mg/L, and MBC₉₀ = 2048 mg/L, for PTSO). It is possible that PTS and PTSO may be affected by active removal mechanisms when they come in contact with these bacteria. Further research is needed to determine with certainty the mechanisms involved in this increased resistance.

All these results are in agreement with the antibacterial effects of garlic previously described in the literature against bacterial isolates from animals and reference strains [3–6]. However, MBC determined in our experiment were much higher compared to Llana-Ruiz-Cabello et al. who demonstrated MBC lower than 5 mg/L in all cases [16]. The differences may be caused by different methodology.

In the present study, the values obtained for MIC and MBC in PTS and PTSO were very similar to those obtained in antibiotics such as nitrofurantoin, aminoglycosides, fluoroquinolones, and some beta-lactams. Based on the data obtained from MIC, the CLSI determines that a very large percentage of enterobacteria should be resistant to these antibiotics (as shown in Tables 1 and 2). It should therefore not be considered for clinical use. Likewise, we may think that the activity shown by PTS and PTSO should also not be considered for clinical use in humans considering the results obtained. However, due to the lack of susceptibility cut-off points for the compounds derived from garlic, no final conclusion can be drawn.

In correspondence with the need of discovering new potentially antibacterial natural products, the activity of these organosulfur compounds described in this study may be considered as promising. Furthermore, the use of naturally and potentially innocuous compounds that can be administered without high restrictions provided us with the possibility to discuss the viability of their application for the treatment of specific infectious pathologies, provided that adequate formulations are developed.

In our opinion, several therapeutic possibilities may be considered, i.e., superficial skin infections, such as acne, folliculitis or impetigo by topical use, the treatment of oral and gastrointestinal infections by oral administration, or even the treatment of urinary tract infections caused by multidrugresistant bacteria applied by intravesical instillation (in the same way that colistin is used). The concentration of the substance in the source of the infection should always be high enough to guarantee that it exceeds the values of MIC against the bacteria causing these processes.

It is clear that, in order to evaluate the real effectiveness of these substances, either in this or another situation, further testing would be necessary with a more diverse and larger group of bacteria. Furthermore, it would be necessary to establish suitable administration routes for the compounds and its efficacy *in vivo*. Finally, the concentrations that they achieve in the different tissues and fluids would also need to be known.

Lastly, PTS and PTSO are perceived as harmless since these compounds occur naturally in foods such as garlic or onion. Nevertheless, further studies on pharmacokinetic and toxicological characteristics are required before safe clinical use is considered. Some recent studies on cell lines and experimental animals reported low acute and subchronic oral toxicity in PTSO and a lack of genotoxicity, both *in vitro* and *in vivo* models [16–19].

5. Conclusion

Our results demonstrate that PTS, but mainly PTSO, have a significant broad spectrum antibacterial activity against a selection of gram-negative and gram-positive multiresistant bacteria isolated from human clinical samples. Further work is needed to demonstrate the effectiveness of these compounds *in vivo* models, although preliminary results in present work provide basic and useful information for development and its potential use in the treatment of human infections.

Data Availability

The data used to support the findings of this study are available from the corresponding author upon request.

Conflicts of Interest

The authors wish to confirm that there are no known conflicts of interest associated with this publication and there has been no significant financial support for this work that could have influenced its outcome.

References

 J. J. Dibner and J. D. Richards, "Antibiotic growth promoters in agriculture: history and mode of action," *Poultry Science*, vol. 84, no. 4, pp. 634–643, 2005.

- [2] M. F. Chellat, L. Raguž, and R. Riedl, "Targeting antibiotic resistance," *Angewandte Chemie International Edition*, vol. 55, no. 23, pp. 6600–6626, 2016.
- [3] R. Ruiz, M. P. García, A. Lara, and L. A. Rubio, "Garlic derivatives (PTS and PTS-O) differently affect the ecology of swine faecal microbiota in vitro," *Veterinary Microbiology*, vol. 144, no. 1-2, pp. 110–117, 2010.
- [4] P. S. Ruddock, M. Liao, B. C. Foster, L. Lawson, J. T. Arnason, and J.-A. R. Dillon, "Garlic natural health products exhibit variable constituent levels and antimicrobial activity against Neisseria gonorrhoeae, Staphylococcus aureus and Enterococcus faecalis," *Phytotherapy Research*, vol. 19, no. 4, pp. 327–334, 2005.
- [5] Z. M. Ross, E. A. O'Gara, D. J. Hill, H. V. Sleightholme, and D. J. Maslin, "Antimicrobial properties of garlic oil against human enteric bacteria: Evaluation of methodologies and comparisons with garlic oil sulfides and garlic powder," *Applied and Environmental Microbiology*, vol. 67, no. 1, pp. 475–480, 2001.
- [6] J. C. Harris, S. L. Cottrell, S. Plummer, and D. Lloyd, "Antimicrobial properties of Allium sativum (garlic)," *Applied Microbiology* and Biotechnology, vol. 57, no. 3, pp. 282–286, 2001.
- [7] M. J. Peinado, R. Ruiz, A. Echávarri, and L. A. Rubio, "Garlic derivative propyl propane thiosulfonate is effective against broiler enteropathogens in vivo," *Poultry Science*, vol. 91, no. 9, pp. 2148–2157, 2012.
- [8] M. J. Peinado, R. Ruiz, A. Echávarri, I. Aranda-Olmedo, and L. A. Rubio, "Garlic derivative PTS-O modulates intestinal microbiota composition and improves digestibility in growing broiler chickens," *Animal Feed Science and Technology*, vol. 181, no. 1-4, pp. 87–92, 2013.
- [9] T. Miron, A. Rabinkov, D. Mirelman, M. Wilchek, and L. Weiner, "The mode of action of allicin: Its ready permeability through phospholipid membranes may contribute to its biological activity," *Biochimica et Biophysica Acta*, vol. 1463, no. 1, pp. 20–30, 2000.
- [10] M. Focke, A. Feld, and H. K. Lichtenthaler, "Allicin, a naturally occurring antibiotic from garlic, specifically inhibits acetyl-CoA synthetase," *FEBS Letters*, vol. 261, no. 1, pp. 106–108, 1990.
- [11] R. S. Feldberg, S. C. Chang, A. N. Kotik et al., "In vitro mechanism of inhibition of bacterial cell growth by allicin," *Antimicrobial Agents and Chemotherapy*, vol. 32, no. 12, pp. 1763–1768, 1988.
- [12] CLSI, Performance Standards for Antimicrobial Susceptibility Testing, Clinical and Laboratory Standards Institute, Wayne, Pa, USA, 26th edition, 2016, CLSI supplement M100S.
- [13] P. Abad, F. J. Lara, N. Arroyo-Manzanares, A. Baños, E. Guillamón, and A. M. García-Campaña, "High-performance liquid chromatography method for the monitoring of the Allium derivative propyl propane thiosulfonate used as natural additive in animal feed," *Food Analytical Methods*, vol. 8, no. 4, pp. 916– 921, 2015.
- [14] G. Bachrach, A. Jamil, R. Naor, G. Tal, Z. Ludmer, and D. Steinberg, "Garlic allicin as a potential agent for controlling oral pathogens," *Journal of Medicinal Food*, vol. 14, no. 11, pp. 1338–1343, 2011.
- [15] D. Jonkers, J. Sluimer, and E. Stobberingh, "Effect of garlic on vancomycin-resistant enterococci," *Antimicrobial Agents and Chemotherapy*, vol. 43, article 3045, 1999.
- [16] M. Llana-Ruiz-Cabello, D. Gutiérrez-Praena, M. Puerto et al., "Acute toxicological studies of the main organosulfur compound derived from Allium sp. intended to be used in active

food packaging," *Food and Chemical Toxicology*, vol. 82, pp. 1–11, 2015.

- [17] P. Mellado-García, S. Maisanaba, M. Puerto et al., "Genotoxicity assessment of propyl thiosulfinate oxide, an organosulfur compound from Allium extract, intended to food active packaging," *Food and Chemical Toxicology*, vol. 86, pp. 365–373, 2015.
- [18] P. Mellado-García, M. Puerto, S. Pichardo et al., "Toxicological evaluation of an Allium-based commercial product in a 90day feeding study in Sprague-Dawley rats," *Food and Chemical Toxicology*, vol. 90, pp. 18–29, 2016.
- [19] P. Mellado-García, M. Puerto, A. I. Prieto et al., "Genotoxicity of a thiosulfonate compound derived from Allium sp. intended to be used in active food packaging: In vivo comet assay and micronucleus test," *Mutation Research - Genetic Toxicology and Environmental Mutagenesis*, vol. 800-801, pp. 1–11, 2016.