

Article

In Silico and In Vitro Assessment of Antimicrobial and Antibiofilm Activity of Some 1,3-Oxazole-Based Compounds and Their Isosteric Analogues

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Abstract: In this paper, we report on the antimicrobial activity assessment of 49 compounds previously synthesized as derivatives of alanine or phenylalanine that incorporate a 4-(4-X-phenylsulfonyl)phenyl fragment (X = H, Cl, or Br), namely 21 acyclic compounds (6 × *N*-acyl- α -amino acids, 1 × *N*-acyl- α -amino acid ester, and 14 × *N*-acyl- α -amino ketones) and 28 pentatomic heterocycles from the oxazole-based compound class (6 × 4*H*-1,3-oxazol-5-ones, 16 × 5-aryl-1,3-oxazoles, and 6 × ethyl 1,3-oxazol-5-yl carbonates). Both in silico and in vitro qualitative and quantitative assays were used to investigate the antimicrobial potential of these derivatives against planktonic and biofilm-embedded microbial strains. Some of the tested compounds showed promising antimicrobial and antibiofilm activity depending on their chemical scaffold and lipophilic character.

Keywords: antimicrobial; antibiofilm; *N*-acyl- α -amino acid; 4*H*-1,3-oxazol-5-one; *N*-acyl- α -amino acid ester; *N*-acyl- α -amino ketone; 1,3-oxazole

1. Introduction

The rise of multidrug-resistant pathogenic microorganisms is a major health concern. In response, there is an urgent need for the identification of novel antimicrobial agents.

The development of new sulfonyl-group-containing analogs is a hot research topic in medicinal chemistry [1–7]. Among these compounds, numerous diaryl sulfones have been found to exhibit a variety of biological activities, including antimicrobial, antioxidant, antimycobacterial, antimalarial, anticancer, anti-inflammatory, and anti-HIV effects [8–17]. Further, some representatives of this class selectively block the 5-HT₆ receptors being developed as therapies for Alzheimer's disease [18,19]. Recently, Alsaedi et al. synthesized a series of pyrazolo [1,5-a]pyrimidine derivatives containing the phenylsulfonyl moiety and evaluated their antimicrobial activities. The results revealed that several sulfone analogues showed effects exceeding the activity of the reference drug. Unexpectedly, it was observed that derivatives containing one sulfone group were more effective against different bacterial and fungal strains than those containing two sulfone groups [20]. Moreover, some unsaturated 4*H*-1,3-oxazol-5-ones bearing the arylsulfonylphenyl moiety in their molecules exhibited good antifungal and antibiofilm potential [21]. Very recently, the results reported by Rashdan et al. highlighted a synthetic 1,2,3-triazole-containing sulfone derivative

structurally inspired by dapsone that exhibited outstanding antimicrobial properties against various bacterial strains [22].

In addition, the 1,3-oxazole is an important heterocyclic nucleus that is present in numerous active substances and displays a wide array of biological properties. Significant research has been conducted to synthesize 1,3-oxazole derivatives and to evaluate their pharmacological profile [23,24]. Therefore, over time, a large number of natural and synthetic 1,3-oxazole-based compounds, which have been associated with a wide spectrum of pharmacological activities, such as antimicrobial, antimalarial, antidiabetic, analgesic, anti-inflammatory, and antitumoral effects, have been reported [25–32]. It has been also shown that saturated 4*H*-1,3-oxazol-5-ones—the stable 5-oxo tautomers of 1,3-oxazol-5-ols—present antimicrobial, anticancer, antiviral, and trypanocidal actions [33–36].

N-acyl amino acids, in which the acyl moiety is derived from fatty acids, are similar to endogenous cannabinoids [37]. Among these *N*-fatty acyl- α -amino acids, *N*-arachidonoylserine has been reported to display antimicrobial and antibiofilm effects against methicillin-resistant *Staphylococcus aureus* (MRSA) strains. In addition, the staphylococcal-biofilm-associated virulence determinants are altered by this agent. *N*-arachidonoylserine is able to change the bacterial membrane potential and prevent biofilm formation without killing the bacterial cells [37,38]. It has been discovered that *N*-acyltyrosine derivatives act as bacterial metabolites that exhibit antibiotic effects against the *Bacillus subtilis* and an average inhibition of the *Pseudomonas aeruginosa* biofilm formation process [39]. *N*-acyl- α -amino acids exhibit other therapeutic effects, such as antihypertensive, mucolytic, anticancer, antianemic, antiulcer, and antioxidant actions [40–46].

Some representatives of the *N*-acyl- α -amino acid ester class have antibacterial, antileishmanial, antiproliferative, antidepressant, and monoamine oxidase inhibitory activities [47–52].

Several *N*-acyl- α -amino ketone derivatives show antiviral, antihypertensive, antithrombotic, and anti-inflammatory properties [53–58].

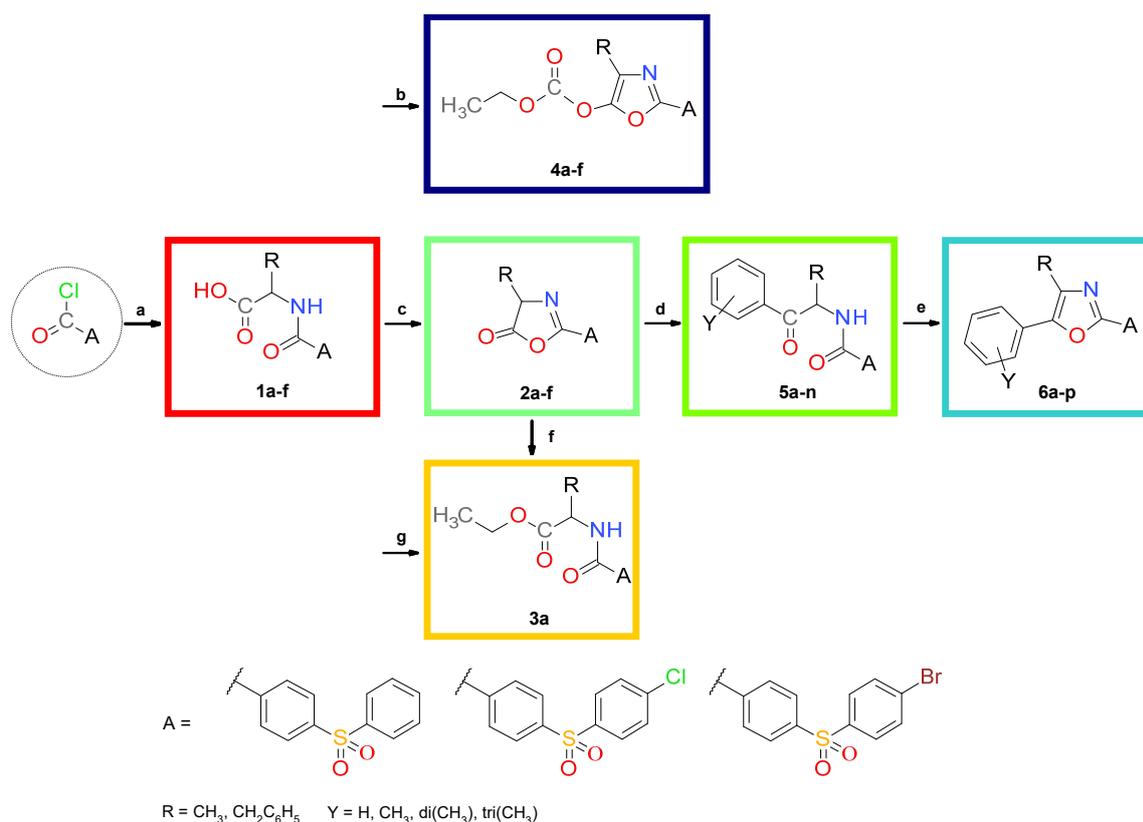
In an attempt to develop new antimicrobial agents, our group devoted considerable interest in the synthesis of compounds based on the 4-(4-*X*-phenylsulfonyl)phenyl fragment as the pharmacophore center [59–65]. In this study, the antimicrobial and antibiofilm effects of 49 compounds (**1a–f**, **2a–f**, **3a**, **4a–f**, **5a–n**, and **6a–p**) from the classes mentioned above were evaluated against different bacterial and fungal strains. In silico prediction of the antimicrobial, pharmacokinetic, and toxicological features of the tested compounds was also performed.

2. Results

2.1. Chemistry

Following the described synthesis procedures, a series of new *N*-acyl- α -amino acids (**1a–f**), 4*H*-1,3-oxazol-5-ones (**2a–f**), *N*-acyl- α -amino ketones (**5a–n**), 1,3-oxazoles (**4a–f**, **6a–p**), and one *N*-acyl- α -amino acid ester (**3a**) were synthesized [59–65]. The general structures of the compounds are presented in Scheme 1.

The chemical structures and purities (%) of the tested compounds **1a–f**, **2a–f**, **3a**, **4a–f**, **5a–n**, and **6a–p** are presented in Table 1.



Scheme 1. The general synthetic methodology as described in our previous papers [59–65]. Reagents and conditions: (a) alanine or phenylalanine/NaOH, CH_2Cl_2 , 0–5 °C, 30 min; (ii) room temperature (r.t.), 1 h; (iii) HCl; (b) $\text{ClCO}_2\text{C}_2\text{H}_5$ /N-methylmorpholine (NMM), CH_2Cl_2 , r.t., 24 h (molar ratio of **1a–f**/ $\text{ClCO}_2\text{C}_2\text{H}_5$ /NMM = 1:1.5:1.5); (c) $\text{ClCO}_2\text{C}_2\text{H}_5$ /NMM, CH_2Cl_2 , r.t., 30 min (molar ratio of **1a–f**/ $\text{ClCO}_2\text{C}_2\text{H}_5$ /NMM = 1:1:1); (d) benzene, toluene, *m*-xylene or mesitylene/anhyd AlCl_3 , r.t., 20 h; (e) POCl_3 , reflux, 4 h; (f) $\text{C}_2\text{H}_5\text{OH}$, reflux, 30 min; (g) $\text{C}_2\text{H}_5\text{OH}/\text{H}_2\text{SO}_4$, reflux, 12 h.

Table 1. The chemical structures and purities (%) of the tested compounds.

Compound	X	R	Purity (%)	Ref.	Compound	X	R	Y	Purity (%)	Ref.
1a	H	CH ₃	99.99	[59]	5a	H	CH ₃	H	96.55	[59]
1b	Cl	CH ₃	99.99	[60]	5b	Cl	CH ₃	H	98.02	[60]
1c	Br	CH ₃	99.99	[61]	5c	Br	CH ₃	H	97.46	[61]
1d	H	CH ₂ C ₆ H ₅	99.99	[62]	5d	H	CH ₃	4-CH ₃	95.10	[59]
1e	Cl	CH ₂ C ₆ H ₅	99.05	[63]	5e	Cl	CH ₃	4-CH ₃	96.65	[60]
1f	Br	CH ₂ C ₆ H ₅	99.63	[64]	5f	Br	CH ₃	4-CH ₃	92.28	[61]
2a	H	CH ₃	91.75	[59]	5g	H	CH ₃	2,4-(CH ₃) ₂	97.16	[59]
2b	Cl	CH ₃	92.92	[60]	5h	Cl	CH ₃	2,4-(CH ₃) ₂	91.10	[60]
2c	Br	CH ₃	90.78	[61]	5i	Br	CH ₃	2,4-(CH ₃) ₂	97.55	[61]
2d	H	CH ₂ C ₆ H ₅	92.03	[62]	5j	H	CH ₂ C ₆ H ₅	4-CH ₃	91.49	[62]
2e	Cl	CH ₂ C ₆ H ₅	91.49	[63]	5k	Cl	CH ₂ C ₆ H ₅	4-CH ₃	98.30	[63]
2f	Br	CH ₂ C ₆ H ₅	90.20	[64]	5l	Br	CH ₂ C ₆ H ₅	4-CH ₃	94.53	[64]
3a	H	CH ₃	94.41	[65]	5m	Cl	CH ₂ C ₆ H ₅	2,4-(CH ₃) ₂	90.83	[63]
4a	H	CH ₃	99.44	[65]	5n	Br	CH ₂ C ₆ H ₅	2,4-(CH ₃) ₂	92.40	[64]
4b	Cl	CH ₃	93.29	[65]	6a	H	CH ₃	H	94.90	[59]
4c	Br	CH ₃	97.80	[65]	6b	Cl	CH ₃	H	96.07	[60]
4d	H	CH ₂ C ₆ H ₅	98.98	[65]	6c	Br	CH ₃	H	98.79	[61]
4e	Cl	CH ₂ C ₆ H ₅	99.36	[65]	6d	H	CH ₃	4-CH ₃	99.50	[59]
4f	Br	CH ₂ C ₆ H ₅	98.81	[65]	6e	Cl	CH ₃	4-CH ₃	97.96	[60]
					6f	Br	CH ₃	4-CH ₃	97.66	[61]
					6g	H	CH ₃	2,4-(CH ₃) ₂	98.68	[59]
					6h	Cl	CH ₃	2,4-(CH ₃) ₂	97.51	[60]
					6i	Br	CH ₃	2,4-(CH ₃) ₂	96.80	[61]
					6j	Br	CH ₃	2,4,6-(CH ₃) ₃	90.58	[61]
					6k	Cl	CH ₂ C ₆ H ₅	H	95.13	[63]
					6l	H	CH ₂ C ₆ H ₅	4-CH ₃	93.31	[62]
					6m	Cl	CH ₂ C ₆ H ₅	4-CH ₃	97.57	[63]
					6n	Br	CH ₂ C ₆ H ₅	4-CH ₃	97.70	[64]
					6o	Cl	CH ₂ C ₆ H ₅	2,4-(CH ₃) ₂	99.15	[63]
					6p	Br	CH ₂ C ₆ H ₅	2,4-(CH ₃) ₂	99.90	[64]

2.2. Antimicrobial Activity Assessment

2.2.1. Qualitative Screening of Antimicrobial Activity

The qualitative screening tests showed a weak growth inhibitory effect of the tested compounds against the studied microorganisms with no clear inhibition zones, the limits of the growth inhibition not exceeding the compound solution deposition area on the agar layer. This could be explained by the poor diffusion into the agar media of the tested compounds. The growth inhibition zones were detected only for compounds **1a–f**; **2a–f**; **3a**; **4a–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g–k**; **6m**; and **6o,p**, which were further evaluated by the quantitative method to determine the minimal inhibitory concentration (MIC) values.

2.2.2. Quantitative Assay of Antimicrobial Activity

The broth dilution method was used to quantitatively assess the in vitro antimicrobial profile of compounds **1a–f**; **2a–f**; **3a**; **4a–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g–k**; **6m**; and **6o,p** against two Gram-positive bacteria (*S. epidermidis* 756 and *B. subtilis* ATCC 6683), two Gram-negative bacteria (*E. coli* ATCC 25922 and *P. aeruginosa* ATCC 27853), and one yeast strain (*C. albicans* 128).

Compounds **1a–c**; **4b–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g,h**; **6k**; **6m**; and **6o,p** did not inhibit the growth of any of the tested microbial strains up to a concentration of 225 µg/mL. In contrast, the MIC values of compounds **1d–f**; **2a–f**; **3a**; **4a**; and **6i,j** were much lower in the case of some microbial strains with MICs in the range of 56.2 to 14 µg/mL being recorded (Table 2). Compound **1e** exhibited a wide spectrum of antimicrobial activity, being active on both Gram-positive and -negative bacterial strains as well as against the fungal strain *C. albicans* 128, for which MICs ≤56.2 µg/mL were recorded. Compounds **1d,e**; **3a**; **4a**; and **6i,j** exhibited very good antifungal activity with MIC values of 14 µg/mL.

Table 2. The MIC and MBIC values (µg/mL) measured for compounds **1d–f**; **2a–f**; **3a**; **4a**; **6i,j**; and **6p** against the tested microbial strains.

Tested Compounds	<i>S. epidermidis</i> 756		<i>B. subtilis</i> ATCC 6683		<i>E. coli</i> ATCC 25922		<i>P. aeruginosa</i> ATCC 27853		<i>C. albicans</i> 128	
	MIC	MBIC	MIC	MBIC	MIC	MBIC	MIC	MBIC	MIC	MBIC
1d	>225	>225	>225	>225	28.1	225	>225	>225	14	112.5
1e	56.2	56.2	>225	>225	28.1	56.2	>225	14	14	112.5
1f	56.2	56.2	>225	>225	>225	>225	>225	28.1	>225	>225
2a	>225	>225	>225	>225	28.1	225	>225	>225	>225	>225
2b	>225	>225	>225	>225	28.1	56.2	>225	>225	>225	>225
2c	>225	>225	>225	>225	28.1	56.2	>225	>225	>225	>225
2d	56.2	56.2	>225	>225	>225	>225	>225	14	>225	>225
2e	>225	112.5	>225	56.2	28.1	56.2	>225	14	>225	>225
2f	>225	225	>225	>225	28.1	56.2	>225	14	>225	>225
3a	>225	112.5	>225	>225	>225	>225	14	14	14	112.5
4a	56.2	56.2	56.2	112.5	>225	>225	>225	14	14	112.5
6i	>225	>225	>225	>225	>225	>225	>225	>225	14	112.5
6j	>225	>225	>225	225	>225	>225	>225	14	14	112.5
6p	>225	112.5	>225	>225	>225	>225	>225	>225	>225	>225
Ciprofloxacin	0.15	0.15	<0.03	<0.03	0.012	0.012	0.15	0.15	- *	-
Fluconazole	-	-	-	-	-	-	-	-	<0.12	<0.12

* -, not tested.

2.2.3. Effects of the Compounds on Biofilm Formation

To further evaluate the effects of the analyzed compounds against microbial biofilms, crystal violet microtiter assay was performed for compounds **1a–f**; **2a–f**; **3a**; **4a–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g–k**; **6m**; and **6o,p**. The analysis showed that biofilm formation was not affected by compounds **1a–c**; **4b–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g,h**; **6k**; **6m**; and **6o**. However, decreased absorbance of the stained biomass was recorded for the biofilms grown in the presence of compounds **1d–f**; **2a–f**; **3a**; **4a**; **6i,j**; and **6p**, with the minimal biofilm inhibitory assay (MBIC) ranging from 14 to 225 µg/mL (Table 2). Compounds **1e**, **2d–f**, **3a**, **4a**, and **6j** demonstrated an MBIC value of 14 µg/mL against the *P. aeruginosa* ATCC

27853 biofilm. Compound **1e** inhibited the biofilm-forming capacity of the Gram-positive *S. epidermidis* 756 strain, the Gram-negative strains *E. coli* ATCC 25922 and *P. aeruginosa* ATCC 27853, and the fungal strain *C. albicans* 128.

Regarding the antimicrobial and antibiofilm activity of the standard antibiotic (ciprofloxacin) and antifungal (fluconazole) agents used as controls, the MIC values were much lower than those obtained for the tested compounds in all of the cases. This is somehow expected given the new compounds are not yet standardized in optimal formulations and their mechanisms of action could be different from those of the control drugs.

2.3. Prediction of the Biological Properties of the Compounds

2.3.1. In Silico Evaluation of the Molecular Mechanism of Action

Based on 2D structural descriptors, the PASS application was used to calculate the probability of the target compounds **1af**, **2a–f**, **3a**, **4a–f**, **5a–n**, and **6a–p** as being active (Pa) or inactive (Pi) on a large series of targets [66]. The analysis returned Pa values higher than the corresponding Pi values for 2097 pharmacological effects, of which 14 were directly related to antibacterial effects. The maximum and minimum predicted Pa values for the 49 compounds and the compounds with Pa values over 0.5 are displayed in Table 3.

Table 3. The probability (Pa) for the compounds to produce biological effects related to antibacterial action as predicted by the PASS application.

Target	Pa Max	Pa Min	Compounds with Pa > 0.5
Anti-infective	0.702	0.218	1a, 1b, 1c, 1d, 1e, 1f, 3a
Antimycobacterial	0.574	0.198	2c, 2f, 5c, 5f
Antituberculosis	0.526	0.199	5c, 5f
Antibiotic glycopeptide-like	0.403	0.083	0
Peptidoglycan glycosyltransferase inhibitor	0.323	0.212	0
Antibacterial	0.312	0.168	0
UDP-N-acetylmuramate-L-alanine ligase inhibitor (MurC)	0.225	0.116	0
Antibacterial, ophthalmic	0.164	0.122	0
Bacterial efflux pump inhibitor	0.119	0.118	0
Antiseptic	0.118	0.117	0
Antibiotic	0.106	0.106	0
Peptidoglycan beta-N-acetylmuramidase inhibitor	0.093	0.067	0
N-acetylmuramoyl-L-alanine amidase inhibitor	0.083	0.054	0
UDP-N-acetylmuramoylalanine-D-glutamate ligase inhibitor (MurD)	0.079	0.059	0
Bacterial leucyl aminopeptidase inhibitor	0.064	0.051	0

A total of 33 compounds presented Pa values over 0.3 for anti-infective effect, and 37 compounds were predicted to have antimycobacterial effects with Pa value above 0.3.

Figure 1 presents the plotted Pa values for anti-infective, antimycobacterial, and antibacterial effects based on the main chemical scaffold.

The prediction results indicated a clear correlation between the chemical scaffolds of the studied compounds and the potential to have an anti-effect, with *N*-acyl- α -amino acids (scaffold 1) emerging as the most promising class (Figure 1b). In the case of antibacterial effects, the prediction indicated 4*H*-1,3-oxazol-5-one ring (scaffold 2) as the most favorable core structure (Figure 1d). The potential of the compounds to produce antimycobacterial effects was not correlated with the chemical scaffolds (Figure 1c).

The PASS application can be used to indicate a probable mechanism of action [67] of new compounds. Compounds **1a–d**, **2a**, and **2d** had significant Pa values for the inhibition of peptidoglycan glycosyltransferase, a valuable target for new antimicrobial therapies [68]. UDP-N-acetylmuramate-L-alanine ligase inhibitor (MurC) is a member of the Mur enzymes family and, similar to peptidoglycan glycosyltransferase, is involved in synthesis of peptidoglycan [69]. Compounds **1a–c**; **1e,f**; **2d,e**; and **3a** presented small but significant Pa values towards this possible mechanism.

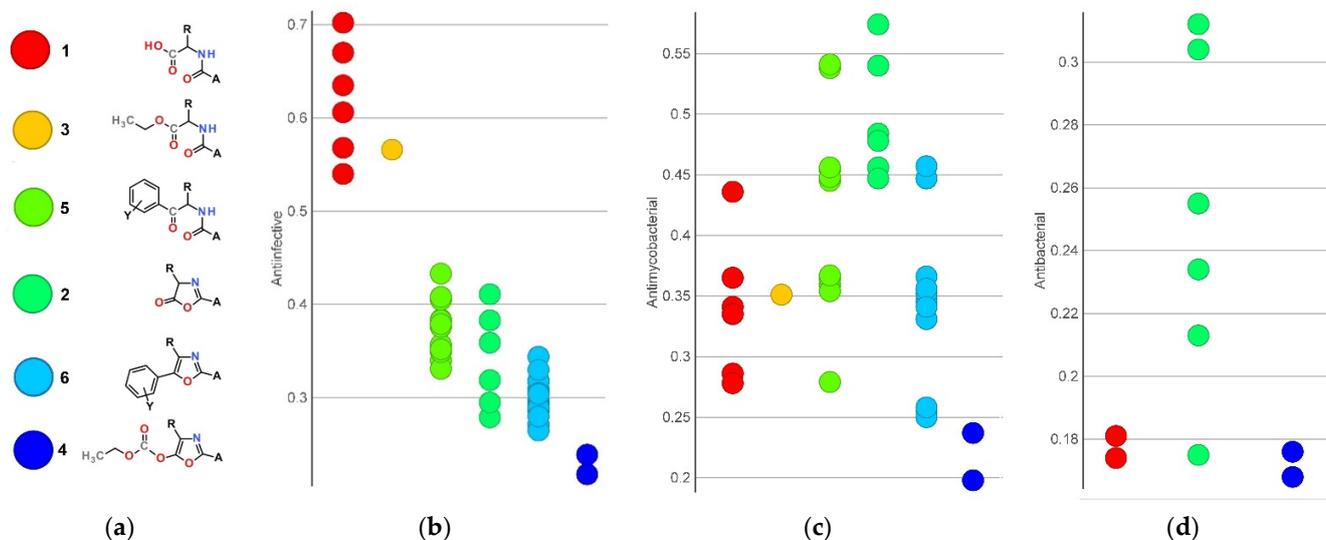


Figure 1. Pa values as predicted by the PASS application: (a) color codes for classification; (b) Pa values for anti-infective effect; (c) Pa values for antimycobacterial effect; (d) Pa values for antibacterial effect.

2.3.2. Structural Descriptors Analysis

DataWarrior v5.2.1 software [70] was used to calculate a series of structural descriptors, namely molecular weight (MW), logarithm of the octanol–water partition coefficient (cLogP), hydrogen bond donors count (HBD), hydrogen bond acceptor count (HBA), polar surface area (PSA), number of rotatable bonds (RB), and druglikeness (DLK). These descriptors are presented in Table 4 with the minimum and maximum values registered for the compounds.

Table 4. Descriptive statistics for the molecular descriptors.

Descriptor	Min	Max
MW	315.35	576.51
cLogP	1.29	7.61
HBD	0	5
HBA	0	2
RB	3	9
PSA	68.6	108.9
DLK	−19.7	4.9

In order to better understand the structure–activity relationships, each descriptor was plotted for both active and inactive compounds based on the MIC values presented in Table 2. The best difference of distribution of values was observed for cLogP (Figure 2). The results indicated that a lower lipophilic character was correlated with a higher antimicrobial effect. The chemical scaffold was also an important factor because some compounds were inactive despite a low cLogP value.

Four compounds with cLogP values in the range of 3.08–3.93 presented antimicrobial and antibiofilm effects towards *S. epidermidis*. Most of the inactive compounds had a cLogP value over 4, indicating that a high lipophilicity was detrimental. Except for the ethyl carbonate derivative **4a**, the active compounds on *S. epidermidis* were all derivatives of phenylalanine. In the case of *E. coli*, the analysis of the structure–activity relationships indicated two major factors: the presence of the *N*-acyl phenylalanine scaffold or its cyclic 4*H*-1,3-oxazol-5-one analogue and a cLogP value under 4.

Based on the ADMET predictions, compound **3a** had the best toxicological and pharmacokinetic profile of the new compounds. This compound was the ethyl ester of the derivative **1a**, indicating that the transformation of compounds **1d–f** could improve their toxicological profile and enhance their antibacterial properties.

3. Discussion

Over time, resistance of human pathogens to major antibiotics increases, and infectious agents that are resistant to most available antibiotics are rising globally [72]. An important strategy to help prevent and confront the resistance problem requires the discovery and development of new bioactive agents against both planktonic and adherent microorganisms. Medical devices and instruments are prone to microbial colonization and biofilm formation. Therefore, the discovery of agents that could prevent biofilm formation or adherence would be of great use. Very recently, we reported on the synthesis and antimicrobial and antibiofilm evaluation results of a series of compounds derived from valine [73,74].

In the present research, we examined the potential antimicrobial activity of some 1,3-oxazole derivatives and their isosteres sharing a 4-(4-X-phenylsulfonyl)phenyl moiety, which were synthesized using two other natural α -amino acids as raw materials, namely alanine and phenylalanine.

Preliminary qualitative antimicrobial screening revealed that compounds **1a–f**; **2a–f**; **3a**; **4a–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g–k**; **6m**; and **6o,p** exhibited inhibitory growth effects, although the growth inhibition zones were detected only in contact with the agar layer. These compounds were further evaluated in vitro to determine their effects on planktonic and adherent microbial growth. Compounds **1a–c**; **4b–f**; **5a,b**; **5d,e**; **5g,h**; **5j,k**; **5m,n**; **6g,h**; **6k**; **6m**; and **6o,p** did not exhibit growth inhibitory effects on any of the tested microbial strains up to a concentration of 225 $\mu\text{g}/\text{mL}$. Regarding the *S. epidermidis* 756 strain, among the different compounds, **1e,f**; **2d**; and **4a** were the most effective with an MIC of 56.2 $\mu\text{g}/\text{mL}$. The most active compound against *P. aeruginosa* ATCC 27853 was **3a**, the compounds **1d,e**; **3a**; **4a**; and **6i,j** were active against *C. albicans* 128 with a low MIC value of 14 $\mu\text{g}/\text{mL}$, and the compounds **1d,e**; **2a–c**; and **2e,f** proved to be active against *E. coli* ATCC 25922 with an MIC of 28.1 $\mu\text{g}/\text{mL}$. Regarding *B. subtilis* ATCC 6683, compound **4a** was found to be the best with an MIC of 56.2 $\mu\text{g}/\text{mL}$, while **3a** had an MIC of 14 $\mu\text{g}/\text{mL}$ against *P. aeruginosa* ATCC 27853.

The analyzed compounds affected the adherence and biofilm formation on inert surfaces at MBIC values in the range of 14–225 $\mu\text{g}/\text{mL}$. Our data demonstrated that compounds **1e**, **2d–f**, **4a**, and **6j** had a distinctly stronger effect on *P. aeruginosa* ATCC 27853 cells embedded in the biofilm (MBIC of 14 $\mu\text{g}/\text{mL}$) than on planktonic cells. We hypothesized that the tested compounds have a specific nonbactericidal mechanism that changes the bacterial cell surface rather than destroying the bacterial cell.

From the results obtained in the quantitative screening, it was observed that 2-[4-(4-X-phenylsulfonyl)benzamido]propanoic acids **1a–c** were inactive at the concentrations used in the assay. However, by intramolecular cyclodehydration, biologically active 4*H*-1,3-oxazol-5-ones **2a–c** were obtained, which displayed growth-inhibitory action with an MIC of 28.1 $\mu\text{g}/\text{mL}$ and had an antibiofilm effect on *E. coli* ATCC 25922 with MBIC values of 56.2 (**2b** and **2c**) and 225 (**2a**) $\mu\text{g}/\text{mL}$. The in silico prediction of the pharmacokinetic profile (ADME properties) indicated that compounds **2a–c** had good pharmacokinetic profiles. Compounds **1a–d**, **2a**, and **2d** were predicted to inhibit peptidoglycan glycosyltransferase.

By opening the 4*H*-1,3-oxazol-5-ones ring, the resulting *N*-acyl- α -amino ketones (**5a–i**) did not show antimicrobial properties up to a concentration of 225 $\mu\text{g}/\text{mL}$. Intramolecular cyclization of *N*-(1-aryl-1-oxopropan-2-yl)-4-(4-X-phenylsulfonyl)benzamides afforded the corresponding 1,3-oxazoles, which were inactive in the tested concentration range, with the exception of 2-[4-[(4-bromophenyl)sulfonyl]phenyl]-5-(2,4-dimethylphenyl)-4-methyl-1,3-oxazole **6i** and 2-[4-[(4-bromophenyl)sulfonyl]phenyl]-5-mesityl-4-methyl-1,3-oxazole **6j**, which showed antifungal action on *C. albicans* 128 (MIC = 14 $\mu\text{g}/\text{mL}$). Moreover, **6j** inhibited the formation of biofilm by *B. subtilis* ATCC 6683 (MBIC = 225 $\mu\text{g}/\text{mL}$) and

P. aeruginosa ATCC 27853 (MBIC = 14 µg/mL), and both 1,3-oxazoles presented antibiofilm effect against *C. albicans* 128 with an MBIC of 112.5 µg/mL. These properties were probably a consequence of the presence of a bromine atom in the *para* position of the arylsulfonylphenyl substituent linked to the C-2 and the *m*-xylyl or mesityl group grafted to the C-5 of the 1,3-oxazole ring. Esterification of 2-[4-(phenylsulfonyl)benzamido]propanoic acid **1a** or ethanolysis of 4-methyl-2-[4-(phenylsulfonyl)phenyl]-1,3-oxazol-5(4*H*)-one **2a** gave ethyl 2-[4-(phenylsulfonyl)benzamido]propanoate **3a**, which was active on *P. aeruginosa* ATCC 27853 and *C. albicans* 128 (MIC = 14 µg/mL). It exhibited an antibiofilm effect on *P. aeruginosa* ATCC 27853 (MBIC = 14 µg/mL), *S. epidermidis* 756, and *C. albicans* 128 (MBIC = 112.5 µg/mL). The conversion of *N*-acylated α -amino acid **1a** into its ester derivative **3a** led to the appearance of antimicrobial properties. Based on *in silico* predictions, the obtained product also had a good pharmacokinetic and toxicological profile. Derivatization of the *N*-acyl- α -amino acid **1a** also led to the ethyl [4-methyl-2-[4-(phenylsulfonyl)phenyl]-1,3-oxazol-5-yl] carbonate **4a**, which showed antimicrobial activity on *S. epidermidis* 756 and *B. subtilis* ATCC 6683 (MIC = 56.2 µg/mL) and on *C. albicans* 128 (MIC = 14 µg/mL). Compound **4a** was also capable of decreasing the biomass of *S. epidermidis* 756 (MBIC = 56.2 µg/mL), *P. aeruginosa* ATCC 27853 (MBIC = 14 µg/mL), *B. subtilis* ATCC 6683, and fungal strain *C. albicans* 128 (MBIC = 112.5 µg/mL).

All three *N*-acyl phenylalanines (**1d–f**) and all three corresponding 4*H*-1,3-oxazol-5-ones (**2d–f**) showed antimicrobial and antibiofilm activities. Thus, 3-phenyl-2-[4-(phenylsulfonyl)benzamido]propanoic acid **1d** was active on *E. coli* ATCC 25922 (MIC = 28.1 µg/mL and MBIC = 225 µg/mL) and *C. albicans* 128 (MIC = 14 µg/mL and MBIC = 112.5 µg/mL). In contrast, 4-benzyl-2-[4-(phenylsulfonyl)phenyl]-1,3-oxazol-5(4*H*)-one **2d**, which resulted from the cyclization of *N*-acyl- α -amino acid **1d**, had an inhibitory effect on Gram-positive bacterium *S. epidermidis* 756 (MIC = 56.2 µg/mL). Moreover, **2d** presented antibiofilm action against *S. epidermidis* 756 (MBIC = 56.2 µg/mL) and *P. aeruginosa* ATCC 27853 (MBIC = 14 µg/mL). The 2-[4-[(4-chlorophenyl)sulfonyl]benzamido]-3-phenylpropanoic acid **1e** showed a broad antimicrobial spectrum on Gram-positive bacterium *S. epidermidis* 756 (MIC = 56.2 µg/mL), Gram-negative bacterium *E. coli* ATCC 25922 (MIC = 28.1 µg/mL), and fungus *C. albicans* 128 (MIC = 14 µg/mL). Compound **1e** presented MBIC values of 56.2 µg/mL for *S. epidermidis* 756 and *E. coli* ATCC 25922, 14 µg/mL for *P. aeruginosa* ATCC 27853, and 112.5 µg/mL for *C. albicans* 128. These effects were probably a result of the presence of the phenylalanine fragment in the molecule and the chlorine atom in the *para* position of the arylsulfonylphenyl moiety. Intramolecular transformation of *N*-acyl- α -amino acid **1e** led to 4-benzyl-2-[4-[(4-chlorophenyl)sulfonyl]phenyl]-1,3-oxazol-5(4*H*)-one **2e**, which exhibited antimicrobial activity only on *E. coli* ATCC 25922 (MIC = 28.1 µg/mL) and antibiofilm effect on *S. epidermidis* 756 (MBIC = 112.5 µg/mL), *B. subtilis* ATCC 6683, *E. coli* ATCC 25922 (MBIC = 56.2 µg/mL), and *P. aeruginosa* ATCC 27853 (MBIC = 14 µg/mL). In addition, 2-[4-[(4-bromophenyl)sulfonyl]benzamido]-3-phenylpropanoic acid **1f** had an inhibitory action on *S. epidermidis* 756 (MIC = 56.2 µg/mL) and inhibited biofilm formation of *S. epidermidis* 756 (MBIC = 56.2 µg/mL) and *P. aeruginosa* ATCC 27853 (MBIC = 28.1 µg/mL). By intramolecular cyclization of this *N*-acyl- α -amino acid (**1f**) to the isosteric 4*H*-1,3-oxazol-5-one analogue **2f**, the antibacterial effect on *S. epidermidis* 756 disappeared, but the obtained compound inhibited the growth of *E. coli* ATCC 25922 (MIC = 28.1 µg/mL). Saturated azlactone **2f** also had antibiofilm activity against *S. epidermidis* 756 (MBIC = 225 µg/mL), *E. coli* ATCC 25922 (MBIC = 56.2 µg/mL), and *P. aeruginosa* ATCC 27853 (MBIC = 14 µg/mL). All *N*-acyl- α -amino ketones **5j–n** obtained by opening the ring of the 4-benzyl-2-[4-(4-*X*-phenylsulfonyl)phenyl]-1,3-oxazol-5(4*H*)-ones **2d–f** were devoid of antimicrobial action up to a concentration of 225 µg/mL. By cyclization of *N*-(1-aryl-1-oxo-3-phenylpropan-2-yl)-4-(4-*X*-phenylsulfonyl)benzamides, the five-membered heterocycles of the 1,3-oxazoles class (**6k–p**) were synthesized, from which only 4-benzyl-2-[4-(4-bromophenyl)sulfonyl]phenyl]-5-(2,4-dimethylphenyl)-1,3-oxazole **6p** affected adherence and biofilm formation of *S. epidermidis* 756 on inert surfaces with an MBIC of 112.5 µg/mL. In the case of this compound, we hypothesized that the antibiofilm activity may be correlated to the substitution with

bromine in the *para* position of the C-2-linked arylsulfonylphenyl fragment, and with the presence of the benzyl substituent bonded to the C-4, and the *m*-xylyl group grafted at position 5 of the 1,3-oxazole nucleus.

In the tested concentration range, {4-(benzyl/methyl)-2-[4-(4-X-phenylsulfonyl)phenyl]-1,3-oxazol-5-yl} ethyl carbonates **4b–f** were proved to be inactive on the studied strains. The results were confirmed by PASS analysis with small probabilities to produce anti-infective, antimycobacterial, or antibacterial effects.

Taken together, the antimicrobial activity results indicate that compounds **1d,e,f**; **2d,e,f**; **3a**; **4a**; and **6i,j** are the most promising candidates for further biological investigations and structural optimization as potential new anti-infective agents, as revealed by the lowest MIC and even MBIC values obtained.

From these compounds, the *in silico* assays predicted the anti-infective potential for **1e**, also exhibiting the broadest antimicrobial spectrum, and **2f**, which proved to successfully inhibit *P. aeruginosa* biofilm development. These compounds have also been predicted to have drug-like properties.

4. Materials and Methods

4.1. General Information

All solvents and reagents were purchased from commercial sources and used without further purification. The absorbance was measured on an Apollo LB 911 ELISA reader (Berthold Technologies GmbH & Co. KG, Waltham, MA, USA).

4.2. Chemistry

The tested compounds **1–6** were previously synthesized [59–65] according to the multiple-step strategy presented in Scheme 1. The *N*-acyl- α -amino acids **1a–f** were obtained by Schotten–Baumann-type *N*-acylation of α -amino acids (alanine or phenylalanine) with 4-(4-X-phenylsulfonyl)benzoyl chlorides (X = H, Cl, or Br). The intramolecular cyclodehydration of compounds **1a–f** using ethyl chloroformate in the presence of *N*-methylmorpholine (NMM) led to 4*H*-1,3-oxazol-5-ones **2a–f** when the molar ratio of **1a–f**/ClCO₂C₂H₅/NMM was 1:1:1 and the reaction time was 30 min and to ethyl 1,3-oxazol-5-yl carbonates **4a–f** when the molar ratio of the reactants was 1:1.5:1.5 and the reaction time was increased to 24 h. The carboxyl group of the *N*-acyl- α -amino acid **1a** was highlighted by its transformation into the corresponding ethyl ester **3a**, which was also obtained by *O*-acylation of the ethanol with 4*H*-1,3-oxazol-5-one **2a**. The Friedel–Crafts acylation, catalyzed by AlCl₃ of the aromatic hydrocarbons with the saturated azlactones **2a–f**, yielded *N*-acyl- α -amino ketones **5a–n**. These acyclic precursors underwent Robinson–Gabriel cyclization in the presence of phosphoryl trichloride with the formation of 5-aryl-1,3-oxazoles **6a–p**. The structures of some of the compounds were confirmed by an additional method. The compounds were purified by recrystallization from water (**1a–f**), cyclohexane (**2a–f** and **5b,c**), toluene (**3a**), ethanol (**4a–f**, **5a**, **5d–g**, **5i–n**, and **6a–p**), or ethanol–water (**5h**). Their purities were verified by RP-HPLC according to previously reported procedures [61–65], with the values ranging between 90.20 and 99.99% (Table 1). As shown in our previous works [59–65], all tested compounds were characterized using spectral methods (UV–vis, IR, MS, and ¹H- and ¹³C-NMR) and elemental analyses, confirming the purity of the compounds.

4.3. Antimicrobial Activity Assessment

4.3.1. Microbial Strains

The antimicrobial activity of the synthesized compounds was tested against two Gram-positive bacteria (*Staphylococcus epidermidis* 756 and *Bacillus subtilis* ATCC 6683), two Gram-negative bacteria (*Escherichia coli* ATCC 25922 and *Pseudomonas aeruginosa* ATCC 27853), and one yeast strain (*Candida albicans* 128).

4.3.2. Qualitative Screening of Antimicrobial Activity

The qualitative screening tests were performed using the agar diffusion method following the CLSI (Clinical and Laboratory Standards Institute) guidelines. The inoculums were prepared from 18–24 h microbial cultures obtained on tryptone soy broth (TSA) for bacteria and on yeast peptone glucose agar (YPGA) for yeast by direct colony suspension in sterile phosphate-buffered saline (PBS). The microbial suspensions turbidity was adjusted to 0.5 McFarland scale and used for the inoculation of the agar plates (Mueller–Hinton agar). Then, 5 μL of the solution of the tested compound at 450 $\mu\text{g}/\text{mL}$ concentration, prepared in DMSO, was placed on the agar surface. The negative (DMSO) and positive (standardized antibiotic discs of ciprofloxacin 5 μg and fluconazole 25 μg) controls were prepared. The Petri dishes were incubated at 37 $^{\circ}\text{C}$, and the diameters of inhibition growth zones were then measured.

4.3.3. Determination of the Minimal Inhibitory Concentration (MIC)

Quantitative analysis of the antimicrobial activity of the tested compounds was carried out using the broth microdilution method following the CLSI guidelines. Two-fold dilutions of the tested compounds were prepared in a liquid growth medium dispensed in a 96-well microplate. The range of final concentrations of the solutions in DMSO of all tested compounds was 1.7–225 $\mu\text{g}/\text{mL}$. Then, each well was inoculated with a microbial inoculum prepared in the same medium after dilution of the standardized microbial suspension adjusted to 0.5 McFarland scale. Binary serial dilutions for DMSO in the liquid growth medium were also prepared. The uninoculated media (MH broth or YPG) and inoculated media served as sterility controls and microbial growth controls. After mixing well, the inoculated 96-well microplates were incubated, without agitation, in aerobic conditions at 37 $^{\circ}\text{C}$ for 24 h. The MIC was measured as the lowest concentration of the tested compound showing no turbidity after 24 h, where turbidity was interpreted as visible bacterial growth. Ciprofloxacin, a broad-spectrum antibacterial agent, and antifungal fluconazole served as controls. The assays were performed in duplicate.

4.3.4. Determination of the Minimal Biofilm Inhibitory Concentration (MBIC)

The crystal violet assay was used to assess the biofilm's susceptibility to the tested compounds. After determination of the MIC values, the 96-well microplates were emptied, washed gently three times with phosphate-buffered saline (PBS) to remove the planktonic microbial cells, and then fixed with cold methanol for 5 min. The adherent cells in the plastic wells were further stained with 1% violet crystal solution for 30 min. The excess dye was removed by washing with distilled deionized water. In each well, 200 μL of 30% acetic acid was added. After 10 min of incubation to release the dye, the biofilm was assessed by measuring the absorbance at 492 nm using a plate-reading spectrophotometer. The MBIC value was determined as the lowest concentration of the tested compounds showing biofilm inhibition compared to the untreated control. The experiment was performed in duplicate.

4.4. Prediction of the Biological Properties of the Compounds

4.4.1. In Silico Evaluation of the Molecular Mechanisms of Action

The study was executed using the PASS (Prediction of Activity Spectra for Substances) software, a product that predicts the pharmacological potential of new compounds. The structures were introduced as SMILES, and the results were considered only if the P_a values were higher than the corresponding P_i values.

4.4.2. Predicted ADME-T Properties

The ADMETlab 2.0 online platform was used to evaluate the in silico ADMET profile for the 49 compounds. Several physicochemical, medicinal chemistry, and ADME properties were computed, together with toxicity endpoints and toxicophore-based assessment.

5. Conclusions

A total of 49 derivatives that incorporate a 4-(4-X-phenylsulfonyl)phenyl fragment into their structure and are designed based on the 1,3-oxazole scaffold and its isosteric analogues were investigated for their antimicrobial and antibiofilm activity. The compounds belonged to the following chemotypes: *N*-acyl- α -amino acids, 4*H*-1,3-oxazol-5-ones, *N*-acyl- α -amino acid esters, *N*-acyl- α -amino ketones, and 1,3-oxazoles classes. The assays revealed that the tested compounds **1d**, **e**, **3a**, and **4a** exhibited the best antimicrobial effects and could be considered as promising candidates for future biological investigations and structural optimization. Among the tested compounds, **1e** exhibited the most intense and broad spectrum of antimicrobial activity, including for the Gram-positive, Gram-negative, and fungal strains, which is probably correlated with the presence of the phenylalanine moiety in its structure and the chlorine atom in the *para* position of the arylsulfonylphenyl fragment. The predictive studies indicate the inhibition of peptidoglycan glycosyltransferase and, to a less extent, the inhibition of the UDP-*N*-acetylmuramate-*L*-alanine ligase as possible mechanisms of action.

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