Synthesis and Antibacterial Activity of Novel Cyclic α-Aminophosphonates

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INTRODUCTION

α-Aminophosphonates I gained great interest of medicinal chemists due to their biological activities [1,2], metabolic stability and their weak toxicity towards mammalians. These organometallic compounds are phosphorus analogues of naturally occurring α-amino acids II (cf. Figure 1). The biological activity of α –aminophosphonates was reported in literatures as anticancer [3,4] and antibacterial [5]. Furthermore, these compounds are used as antifungal, anticancer, antiviral agents [6,7] and herbicides [8,9]. Therefore, α–aminophosphonates considered a promising pharmacophore core in the fields of drug design and discovery [10-12]. Moreover, nitrogen containing heterocyclic compounds such quinolines III, received much attention due their wide pharmacological applications [13]. The enhancement and synergistic effect of the quinoline moiety by modification of side chain at (C4 position) and/or with conjugation with other structural motifs [14,15]. For instance, hybridization of quinoline moiety with α-aminophosphonates was proved significant synergistic effect on biological activity [16]. Based on above-mentioned, the aim of the work presented in this study is to synthesize some new α –aminophosphonates bearing quinoline nucleus. Further aim is to screen the resulting hybrids as antibacterial agents against different bacterial strains. Moreover, correlate the structures with the biological activity.

FIGURE 1: Structures of α-Aminophosphonates I, α-Amino acids II and quinoline III.

MATERIALS AND METHODS

All 1H-NMR experiments (DMSO-d$_6$, and CDCl$_3$) were carried out with a 500 MHz Jeol at El-Mansoura University in addition, some samples were performed with 400 MHz Varian at the Main Chemical Warfare Laboratories, Egypt. Chemical shifts are reported in part per million (ppm) relative to the position respective to the solvent.
The mass and IR spectroscopy spectra were performed at Al-Azhar and El-Mansoura University respectively, Egypt [17,18]. Melting points were recorded on Stuart scientific melting point apparatus and are uncorrected. Antimicrobial activity for all synthesized compounds was screened at National Organization for drug and Control Research, Egypt. All reactions were followed by thin layer chromatography (TLC) on kiesel gel F254 precoated plates (Merck).

Starting materials, glutaraldehyde, 4,7-dichloroquinoline, amines, diphenyl phosphate and triphenyl phosphate were commercially available from sigma alderic. Solvents such as ethanol, acetonitrile, dichloromethane, hexane, dimethylformamide and diethyl ether were used without any further purification. All synthesized 4-aminquinoline derivatives were carried out as reported in literature [19-21].

Synthesis of α-Aminophosphonate Derivatives Using Triphenyl Phosphites

General Procedure for The Synthesis of α-Aminophosphonate Derivatives (4a-e): To solution of glutaraldehyde (0.24 mL, 2 mmol), dimamines (2 mmol), triphenyl phosphate (0.39 mL, 2 mmol) dissolved in acetonitrile (5 mL) Lewis acid catalyst LiClO4 (2.1 mg, 10% mol) was added. The reaction mixture was stirred at r.t and the completion of the reaction was monitored by TLC (48 hours). The formed precipitate was filtered off and dried to afford α-aminophosphonates 5 in good yields.

Diphenyl (α-Aminopiperidin-2-yl) Phosphonate 4a: Yield = (0.26g, 40%), reddish brown viscous oil, IR (KBr) cm⁻¹: 3340 (NH), 3070 (=CH), 2955, 2861 (-CH), 1629 (C=C, aromatic), 1537 (C=N), found: 332.33 [M⁺]

General Procedures for Synthesis of α-Aminophosphonate Derivatives Bearing Quinoline Moiety: (7a-e): Glutaraldehyde (0.24 mL, 2 mmol), diamines (2 mmol) and diphenyl phosphate (0.37 mL, 2mol) were dissolved in acetonitrile (5 mL) in presence of LiClO4 as Lewis acid catalyst (2.1 mg, 10% mol) was added. The reaction mixture was stirred at r.t and the completion of the reaction was monitored by TLC (48 hours). The desired product was filtered off and dried to afford α-aminophosphonates 9 with good yields.

Diphenyl (α-((7-Chloroquinolin-4-yl) Amino) Piperidin-2-yl) Phosphonate 7a: Yield = (0.44g, 45%), off white solids, m.p = 140-142 °C, IR (KBr) cm⁻¹: 3445 (NH), 3075 (=CH), 2926 (-CH), 1594 (C=C, aromatic), 1226 (P=O), 755 (POC). EIMS m/z: (C18H12ClN2O3P) calcd.: 408.43 [M⁺]

Diphenyl (α-(2-Aminophenyl) Piperidin-2-yl) Phosphonate 7b: Yield = (0.5g, 48%) , brown solids, m.p = 112-114 °C, IR (KBr) cm⁻¹: 3244 (NH), 3035(=CH), 2930 (-CH), 1612 (C=C, aromatic), 1546 (C=N) 1207 (P=O), 764 (POC). EIMS m/z: (C18H12ClN2O3P) calcd.: 493.92 [M⁺], found: 493.55.

Diphenyl (α-((7-Chloroquinolin-4-yl) Amino) Propyl) Phosphonate 7c: Yield = (0.46g, 43%), pale yellow solids, m.p = 104-106 °C, IR (KBr) cm⁻¹: 3421, 3247 (NH), 3109 (=CH), 2937 (-CH), 1615 (C=C, aromatic), 1580 (C=N) 1205 (P=O), 763 (POC). EIMS m/z: (C20H14ClN2O3P) calcd.: 536.00 [M⁺], found: 538.28.

Diphenyl (α-(2-(7-Chloroquinolin-4-yl) Amino) Ethyl) Phosphonate 7d: Yield = (0.4g, 48%), black viscous oil, IR (KBr) cm⁻¹: 3421, 3247 (NH), 3109 (=CH), 2926 (-CH), 1594 (C=C, aromatic), 1537 (C=N) 1365 (P=O), 700 (POC). EIMS m/z: (C24H18ClN2O3P) calcd.: 570.62 [M⁺], found: 570.66.
\( ^1\)H-NMR (DMSO \( d_6 \), 400MHz): \( \delta \) ppm 3.72 (br.m, 8H, 4CH\(_2\)), 6.05 (s, 1H, CH=\( N_\)\( P \)), 6.73-8.05 (m, 18H, Ar-H), 8.62 (s, 1H, Ar-H), 8.59 (d, 1H, CH=\( N_\)\( \equiv \)O) ppm 6.05 (s, 1H, C=\( N_\)\( P \)).

**Diphenyl(\( \alpha \)-4-((7-Chloroquinolin-4-yl) Amino) Phenyl) Piperidin-2-yl) Phosphonate 7e:** Yield (0.58g, 51%), pale yellow solids, m.p = 265-267° C, IR (KBr) cm\(^{-1}\): 3432, 3320 (NH), 3068 (\( \equiv \)C=CH), 2903 (\( \equiv \)CH). 1620 (C=C, aromatic), 1542 (C=N) 1230 (P=O), 792 (POC). EIMS m/z: \( (C_\text{13}H_{\text{18}}N_\text{3}O_\text{3}P) \) calcd.: 570.02 [M\(^+\)], found: 570.48. \(^1\)H-NMR (DMSO \( d_6 \), 400MHz) ppm \( \delta \): 1.16 (m, 2H, CH\(_2\)), 3.05 (br.m, 6H, 3CH\(_2\)), 5.91 (s, 1H, CH=\( N_\)\( P \)), 6.54-8.38 (m, 19H, Ar-H), 8.59 (d, 1H, CH=\( N_\)\( \equiv \)O), 10.20 (brs, 1H, NH).

**Biological Screening**

**Bacterial Strains**

The antibacterial activity of different chemical compounds was tested against a set of *E.coli* ATCC 11229, *Staphylococcus aureus* ATCC 25923 and human MDR pathogenic clinical strains of *Klebsiella pneumonia* (Kp1, Kp5, Kp Band Kp9) [14], and *E. coli* (E13, E15, E16 and E17) (unpublished data), and *Staphylococcus aureus* (SA4) (unpublished data) which were collected from Egyptian hospitals. Bacterial strains were recovered by culturing overnight at 37°C in Nutrient agar media (Oxoid, UK).

**Compounds Preparation for Antimicrobial Activity**

Compounds were dissolved in Dimethyl sulfoxide (DMSO), and then a final concentration of 100mg/ml was prepared using sterile distilled water. Control DMSO was used to check the effect of solvent on the growth of microorganisms.

**Agar Well Diffusion Method**

A volume of 100µl bacterial suspension (\( 10^6 \) CFU/ml) was spread on Müller-Hinton agar (Oxoid, UK). Wells were made on agar plates using sterile cork borer, and then 100 µl of each prepared compound was introduced into appropriately marked wells, then plates were incubated for 24 h at 37°C. DMSO was taken as a negative control. Antibacterial activity was evaluated by measuring the diameter of the inhibition zone in millimeters for each tested organism compared with the negative control [15].

**Result and Discussion**

**Chemistry**

A newly synthesized compounds of cyclic \( \alpha \)-aminophosphonates were performed by three one pot reaction of glutaraldehyde 1 (0.24 mL, 2 mmol), with various diamines 2 (2 mmol) and triphenyl phosphite 7 (0.39 mL, 2 mmol) in presence of (LiClO\(_4\)) in acetonitrile with equal molar ratio (1:1:1) to afford 5 in good yields as shown in scheme 1. In structure characterization of 5, the IR spectra showed the absorption bands of (NH\(_2\)) ranging from 3432 to 3291cm\(^{-1}\) and (\( \equiv \)P=O) from 1367 to 1226 cm\(^{-1}\). The \(^1\)H-NMR spectra showed the chemical shift of (\( \equiv \)C=\( N_\)\( P \)) as singlet peak ranging from 6: 6.00 to 6.08 ppm which affirmed the formation of \( \alpha \)-aminophosphate. It is noticeable that the absence of the aldehydic group (CHO) in \(^1\)H-NMR confirmed the formation of cyclic \( \alpha \)-aminophosphonates through intra molecular cyclization followed by water elimination [17,18]. The El mass spectra confirmed the formation of 5 throughout the coincidence of the molecular ion peak with expected mass.

The recommended mechanism for preparation of cyclic \( \alpha \)-aminophosphonates using LiClO\(_4\) as a catalyst illustrated in Scheme 2, the reaction starts with activation of carbonyl group of glutaraldehyde by Lewis acid catalyst (LiClO\(_4\)) followed by condensation of the carbonyl group of the starting dialdehyde with primary amines to afford the Schiff base B. Then the nitrogen of Schiff base that is formed in the first step of \( \alpha \)-aminophosphonates formation donates a pair of electron to make a coordinate bond with LiClO\(_4\). This makes nitrogen positively charged which induces partial positive charge on sp\(^3\) carbon followed by nucleophilic attack of nitrogen of Schiff base on carbon atom of the other carbonyl group of glutaraldehyde. Then the free pair of electrons of phosphorus attacks to the partially positively charged carbon B followed by elimination of phenol C and afford cyclic \( \alpha \)-aminophosphonates as depicted in Scheme 3. Synthesis of novel compounds of cyclic \( \alpha \)-aminophosphonate bearing guinoline moiety was accomplished by three one pot reaction of glutaraldehyde 1 (0.24 mL, 2 mmol), 4-aminoquinoline derivatives 6 (2 mmol) and diphenyl phosphite 7.
(0.37 mL, 2 mmol) in presence of Lewis acid catalyst (LiClO₄) and acetonitrile with equal molar ratios (1:1:1) to get the desired product with good yields.

Scheme 2: Suggested mechanism of α-aminophosphonate.

Scheme 3: Synthesis of α-aminophosphonate bearing 4.7 dichloroquinoline.

Scheme 4: A plausible mechanism of α-aminophosphonate bearing 4.7 dichloroquinoline.
In the structure elucidation of installation of 7, the IR spectra of synthesized compounds showed the absorption bands of \((\text{NH})\) ranging from 3432 to 3291 cm\(^{-1}\) and \((\text{P=O})\) from 1365 to 1205 cm\(^{-1}\). The \(^1\text{H}-\text{NMR}\) spectra showed the chemical shift of \((\text{CH}=\text{P})\) as singlet peak ranging from 5.91 to 6.17 ppm which confirmed the formation of \(\alpha\)-amino phosphonate. While the aromatic \((\text{CH}=\text{N})\) of quinoline moiety was reported ranging from 8.52 to 8.88 ppm. The absence of the aldehydic group \((\text{CHO})\) in \(^1\text{H}-\text{NMR}\) confirmed the formation of cyclic \(\alpha\)-amino phosphonates through intramolecular cyclization followed by water elimination. The EI mass spectrum affirmed the formation of 7 throughout the coincidence of the molecular ion peak with expected mass.A proposed mechanism of formation of 7 was established. A plausible mechanism for preparation of \(\alpha\)-amino phosphonates using LiClO\(_4\) as a catalyst showed in Scheme 4. First, the activation of carbonyl group of glutaraldehyde was achieved by Lewis acid catalyst, LiClO\(_4\), followed by condensation of the carbonyl group with derivatives of 4-aminquinoline 5 to form the Schiff base B. Then the free pair of electron on nitrogen of Schiff base attacked the partially positive carbon of the other carbonyl group of glutaraldehyde, which was activated by LiClO\(_4\) followed nucleophilic attack of phosphorus to the carbon of imine double bond to afford the target compound 7.

Table 1: Antimicrobial activity of cyclic \(\alpha\)-amino phosphonate using tri phenylphosphite against tested microorganisms.

| Chemical Formula | Bacterial Species (Inhibition zones mm ± SD*) | Gram-Negative Bacteria | Gram-Positive bacteria |
|------------------|-----------------------------------------------|------------------------|------------------------|
|                  | CIP-Resistant E. Coli                         | CIP-Resistant Klebsiella Pneumonia | E. coli ATCC 11229 | Staph. Aureus ATCC 25923 | MRSA Staph. aureus SA4 |
|                  | E15 | E16 | E17 | E13 | Kp1 | Kp8 | Kp9 | Kp5 |          |          |          |          |
| 4a                | 2 ± 0 | 2 ± 0 | 3 ± 0 | 2.2 ± 0 | 1.7 ± 0 | 2.7 ± 0 | 1.9 ± 0 | 3.6 ± 0 | 2.95 ± 0.05 | 2.5 ± 0 | 1.85 ± 0 |
| 4b                | 0 | 0 | 0 | 0 | 2.2 ± 0 | 1.15 ± 0.05 | 1.9 ± 0.1 | 0 | 1.85 ± 0.05 | 1.85 ± 0 | 1.35 ± 0.05 |
| 4c                | 0 | 0 | 0 | 0 | 2 ± 0 | 1.55 ± 0.05 | 2 ± 0 | 0 | 2.2 ± 0.05 | 0 | 0 |
| 4d                | 0 | 0 | 0 | 0 | 1.15 ± 0.05 | 0 | 0 | 1.55 ± 0.05 | 0 | 0 |
| 4e                | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1.2 ± 0 | 0 | 0 |
| 7a                | 1.2 ± 0 | 1.2 ± 0 | 1.35 ± 0.05 | 0 | 0 | 0 | 0 | 1.2 ± 0 | 1.85 ± 0.05 | 0 | 0 |
| 7b                | 3.1 ± 0.1 | 0 | 0 | 2.3 ± 0.1 | 0 | 0 | 0 | 0 | 2.2 ± 0.05 | 0 | 0 |
| 7c                | 1.5 ± 0 | 1.3 ± 0 | 1.5 ± 0 | 1.7 ± 0 | 1.5 ± 0 | 1.05 ± 0.05 | 1.25 ± 0.05 | 0 | 2.0 ± 0.05 | 0 | 0 |
| 7d                | 0 | 0 | 0 | 0 | 0 | 0 | 1.03 ± 0.05 | 1.5 ± 0.05 | 1.05 ± 0 | 0 | 0 |
| 7e                | 0 | 0 | 0 | 0 | 0 | 0 | 1.15 ± 0.05 | 1.8 ± 0.05 | 1.15 ± 0 | 1.8 ± 0.05 |

On the other hand, the latter group with quinoline moiety, compound diphenyl (1-(3-(7-chloroquinolin-4-yl) amino) propyl) piperidin-2-yl) phosphonate (7c) was the most active one against all tested pathogens (Figure 2). A comparison between cyclic \(\alpha\)-amino phosphonate using triphenylphosphite group and cyclic \(\alpha\)-amino phosphonate group bearing quinoline moiety showed that the former one was the most effective than the one bearing quinolone moiety against MDR bacteria. However, substitution 1.3 di amino propane connected with antimicrobial activity. Several studies stated that \(\alpha\)-amino-phosphonates have antibacterial and antifungal activities regardless of the moiety [19-22].

Biological Evaluation

In vitro, two groups of synthetic chemical compounds, cyclic \(\alpha\)-amino phosphonate using tri phenylphosphite and cyclic \(\alpha\)-amino phosphonate bearing quinolone moiety, were tested against MDR strains of E. coli and K. pneumonia that already resistant to ciprofloxacin [17], Staphylococcus aureus ATCC25923, MRSA S. aureus, and finally clinical Candida albicans using well diffusion method and the data analyzed as recommended by CLSI [18]. The cyclic \(\alpha\)-amino phosphonate using tri phenylphosphite compounds showed various antimicrobial activities against tested microorganisms, where diphenyl (1-aminopiperidin-2-yl) phosphonate (5a) was observed to record a broad-spectrum activity (Table 1). MDR E. coli 17 and K. pneumonia (with gyrase mutation at codon Asp 83 and Ser 87) were the most sensitive among the other tested bacteria. Based on our data, it has been observed that the substitution with other diamines as ethylene diamine, 1.3 di amino propane, phenylene diamine at (meta-position phenyl ring), 1.4 phenylene diamine didn’t exert the desired antimicrobial activity as compared with that have hydrazine diamine.
Conclusion

The formation of novel cyclic α-aminophosphonate was installed by three one pot reaction of glutaraldehyde, various diamine derivatives and di or triphenylphosphite in presence of LiClO₄ as a catalyst. It was reported that these newly synthesized compounds have potent activity against Klebsiella pneumonia and E. coli bacteria.

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Conflict of Interest

No conflict of interest.

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