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Novel Quaternary Ammonium Derivatives of 4-Pyrrolidino Pyridine: Synthesis, Structural, Thermal, and Antibacterial Studies

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Abstract: Six novel quaternary ammonium derivatives of 4-pyrrolidino pyridine were prepared and isolated via a facile one-pot synthesis and a simple purification procedure. The purity and the molecular structure of the 4-pyrrolidino pyridine derivatives were confirmed with 
$^1$H and $^{13}$C NMR spectroscopy and powder X-ray diffraction techniques. The crystal structures of the compounds were characterized by single crystal X-ray diffraction (SCXRD) and their thermal properties were studied by Differential Scanning Calorimetry (DSC) analyses. The antibacterial properties of the title compounds against five bacterial strains were evaluated using Kirby–Bauer disk diffusion susceptibility test. The compounds crystallize in the monoclinic or orthorhombic crystal systems (space groups: $P_2_1/c$, $P_2_1/n$, or $P_2_1_2_1_2_1$) and their crystal structures are stabilized by a combination of intra- and intermolecular halogen bonding interactions, short contacts and $\pi$-$\pi$ interactions. Above interactions, they contribute to the thermal stability and lack of phase transition effects up to 350 °C. Two of the compounds possess antibacterial effect against $E$. coli or $S$. aureus bacterial strains—similar or better than the kanamycin reference.

Keywords: single crystal XRD; NMR; powder XRD; DSC; quaternary ammonium salts; antibacterial activity

1. Introduction

The rapidly increasing bacterial resistance against regularly used antibiotics and antimicrobial agents motivates the necessity for discovery of new antibacterial compounds. The quaternary ammonium salts (QAS) are promising candidates in this regard. They are heterocyclic compounds, composed of positive nitrogen “head” and lipophilic “tail”. This structural peculiarity determines QAS’s amphiphilic properties and contributes to their broad range of applications namely as cationic surfactants [1,2], phase-transfer catalysts [3,4], fabric softeners [5], ionic liquids [6,7], lithium battery binders [8,9], etc. Compounds such as benzalkonium chloride, cetalkonium chloride, etc., are an integral part of different household, industrial, and clinical non-selective disinfectants because of their strong bacteriostatic and bactericidal effect. In general, the basic mechanisms of biological activity of antibiotics involve metabolic pathway inhibition, interference with the synthesis of essential nucleic acids and cell wall components or disruption of the bacterial membrane [10]. The last mechanism is considered to be the main feature of QAS’s antibacterial action based on the electrostatic and/or hydrophobic interactions with the negatively charged cell membrane. Such interactions destabilize the intracellular matrix of the bacteria, leading to leakage of essential fluids and cell death [11]. On the other hand, principal microbial mechanisms of drug resistance are based on gene mutations and horizontal
gene-transfer (HGT) [12,13]. The mutations, for instance, can increase the bacterial ability to produce antibiotic targeting enzymes, e.g., β-lactamases that reduce the concentration of the drug or may also intensify the over-expression of specific protein transporters (“efflux pumps”) that can expel the antibacterial agent out of the cell [14–16]. The “efflux pumps”, in particular, are one of the main reasons for reduced activity of quaternary ammonium salt based antibacterial drugs. Another reason is the structure of the bacterial membrane as QASs are less active against Gram-negative than Gram-positive bacteria due to the presence of additional outer lipid layer.

Several authors [17–19] point out, that in order to be effective antibacterial agents, the optimal length of the QAS’s carbon linkers/chains should be around 12 to 16 carbon atoms. Thus, to exploit/enhance the biological properties and counteract the mechanisms of resistance, QASs are incorporated (encapsulated) in different types of materials and are subject to various studies. Kang et al. [20] work up cotton fibers with a mixture of QASs and silver particles that possess synergistic antimicrobial properties against \textit{P. aeruginosa}. Liu et al. [21] prepared terpene based polyurethane coatings containing chemically bonded quaternary ammonium salts that exhibit significant antibacterial activity against \textit{S. aureus} and \textit{E. coli}. Several studies focus on chitosan–QAS antifungal properties against phytopathogenic fungal species [22–24]. The 4-pyrrolidino pyridine (4PP) moiety as a target of quaternization has been acknowledged in recent years. The 4PP quaternary ammonium derivatives are part of diverse scientific studies encompassing various applications as: potent acetyl- and butyrylcholinesterase inhibitors [25]; catalysts for alkene isomerization–hydroboration reactions [26]; asymmetric non-enzymatic catalysis of acyl-transfer reactions [27]; catalysts for 1,2- and 1,4-addition reactions [28] and even as part of different metal-organic complexes [29], etc.

In general, the quaternization process, pioneered by Menshutkin, is best described as $\text{S}_2\text{N}_2$ reaction, where tertiary amine reacts with alkyl (aryl) substituted halide to form quaternary ammonium salt [30]. The role of the reaction media is important and polar aprotic solvents are favored because they contribute to the formation of solvate ions [31–33]. Different modifications of the reaction [34,35] were made through the years in order to increase the yield, reactivity, and to facilitate the purification process. The antimicrobial activity research involving the 4-pyrrolidino pyridine fragment—in particular—is a continuation of our previous reported paper that hints the potential antimicrobial properties of such chemical species [36].

Herein, we report on a facile one-step synthesis of quaternary ammonium derivatives of 4-pyrrolidino pyridine, in which the long aliphatic “tail” is replaced by compact aryl-containing substituents, their solution, solid state characterization, thermal properties, and antimicrobial activity against Gram-positive and Gram-negative bacterial strains.

2. Materials and Methods

2.1. Synthesis and Crystallization

All starting reagents were purchased from AlfaAesar and were used without additional purification. The general one-step procedure for the synthesis (Scheme 1) of the quaternary ammonium compounds involved dissolution of the starting compounds in polar aprotic solvent (acetonitrile (ACN)) with constant stirring at temperature near the boiling point of the solvent. The reactions were performed in inert atmosphere (argon) because of the possible hydrolysis of the halogen compounds. Due to the excellent solubility of the starting reagents in the reaction media, the end of quaternization was marked with the formation of microcrystalline precipitate. In some cases, even large single crystals appeared directly from the acetonitrile solution after cooling down to room temperature (~20 °C). The purification procedure consisted of several washes with solvent and/or recrystallization from methanol or methanol/acetone. The purity and the molecular structure of the synthesized compounds was confirmed with $^1$H and $^{13}$C NMR spectroscopy and powder X-ray diffraction techniques.
2.1.1. Synthesis of 1-(2-(3-Nitrophenyl)-2-Oxoethyl)-4-(Pyrrolidin-1-yl)Pyridin-1-Ium Bromide, Compound 1

4-pyrrolidino pyridine (74 mg, 0.5 mmol) and 2-bromo-1-(3-nitrophenyl) ethan-1-one (122 mg, 0.5 mmol) were dissolved in 5 mL acetonitrile. Crystalline precipitate formed almost immediately after dissolution and was left stirring for 6 h at 80 °C. The product was filtered and washed with acetonitrile. Crystals were extracted for SCXRD. Compound 1: yield = 61%, light brown needles (ACN); mp 262-264 °C; 1H NMR (DMSO-d$_6$, 600 MHz) $\delta$ 2.202 (4H, $m$, CH$_2$-3 and CH$_2$-4 Pir), 3.523 (4H, $m$, CH$_2$-2 and CH$_2$-5 Pir), 3.893 (3H, $s$, OCH$_3$-4 Ar), 4.014 (3H, $s$, OCH$_3$-2 Ar), $\delta$ 5.657 (2H, $s$, CH$_2$-CO), 6.715 (1H, $dd$, $J$ = 8.9 Hz, 2.3 Hz, CH-5 Ar), 6.774 (1H, $s$, CH-2), 6.931 (2H, $d$, $J$ = 7.7 Hz, CH-3 and CH-5 Pyr), $\delta$ 7.846 (1H, $d$, $J$ = 8.9 Hz, CH-6 Ar), $\delta$ 8.144 (2H, $d$, $J$ = 7.7 Hz, CH-2 and CH-6 Pyr); $^{13}$C NMR (DMSO-d$_6$, 151 MHz) $\delta$ 25.17 (CH$_2$-2 and CH$_2$-4 Pir), 48.69 (CH$_2$-2 and CH$_2$-5 Pir), 56.29 (OCH$_3$-4 Ar), 56.71 (OCH$_3$-2 Ar), 66.41 (CH$_2$-CO), 98.79 (CH-3 Ar), 107.48 (CH-5 Ar), 108.06 (CH-3 and CH-5 Pyr), 116.85 (C$_q$-1 Ar), 132.73 (CH-6 Ar), 143.66 (CH-2 and CH-6 Pyr), 153.45 (C$_q$-4 Pyr), 162.39 (C$_q$-2 Ar), 166.05 (C$_q$-4 Ar), 190.35 (C=O).

2.1.2. Synthesis of 1-(2-(3,4-Dihydro-2H-Benzo[B][1,4]Dioxepin-7-Yl)-2-Oxoethyl)-4-(Pyrrolidin-1-Yl)Pyridin-1-Ium Bromide, Compound 2

4-pyrrolidino pyridine (74 mg, 0.5 mmol) and 2-bromo-1-(3,4-dihydro-2H-benzo[b][1,4]dioxepin-7-yl)ethan-1-one (136 mg, 0.5 mmol) were dissolved in 5 mL acetonitrile and the solution was stirred for 24 h at 80 °C. After that the clear reaction mixture was left to cool down to 20 °C and after 1 to 2 h, colorless crystals started to form. Single crystals were extracted for SCXRD and the rest were filtered and washed with acetonitrile. Compound 2: yield 71%, colorless blocks (ACN); mp 204-207 °C; 1H NMR (DMSO-d$_6$, 600 MHz) $\delta$ 2.026 (4H, $m$, CH$_2$-3 and CH$_2$-4 Pir), 2.188 (2H, $m$, C–CH$_2$–C Ar), 3.529 (4H, $m$, CH$_2$-2 and CH$_2$-5 Pir), 4.227 (2H, $m$, $^1$CH$_2$-O Ar), 4.290 (2H, $m$, $^1$CH$_2$-O Ar), 5.890 (2H, $s$, CH$_2$-CO), 6.961 (2H, $d$, $J$ = 7.7 Hz, CH-3 and CH-5 Pyr), 7.159 (1H, $d$, $J$ = 8.3 Hz, CH-5 Ar), 7.607 (1H, $d$, $J$ = 2.2 Hz, CH-2 Ar), 7.631 (1H, $dd$, $J$ = 8.3 Hz, 2.2 Hz, CH-6 Ar), 8.144 (2H, $d$, $J$ = 7.7 Hz, CH-2 and CH-6 Pyr); $^{13}$C NMR (DMSO-d$_6$, 151 MHz) $\delta$ 25.06 (CH$_2$-3 and CH$_2$-4 Pir), 30.94 (C–CH$_2$–C Ar), 48.76 (CH$_2$-2 and CH$_2$-5 Pir), 62.57 (CH$_2$-CO), 70.89 (CH$_2$-O Ar), 80.31 (CH-3 and CH-5 Pyr), 122.09 (CH-2 Ar), 122.31 (CH-5 Ar), 124.28 (CH$_3$ Ar), 129.39 (C$_q$-1 Ar), 143.56 (CH-2 and CH-6 Pyr), 150.84 (C$_q$-3 Ar), 153.47 (C$_q$-4 Pyr), 156.34 (C$_q$-4 Ar), 191.20 (C=O).
2.1.3. Synthesis of 1-(2-(2,4-Dimethoxyphenyl)-2-Oxoethyl)-4-(Pyrrolidin-1-yl)Pyridin-1-Ium Bromide, Compound 3

4-pyrrolidino pyridine (74 mg, 0.5 mmol) and 2-bromo-1-(2,4-dimethoxyphenyl)ethan-1-one (130 mg, 0.5 mmol) were dissolved in 5 mL acetonitrile and the solution was stirred for 24 h at 80 °C and remained colorless. The reaction mixture was left to cool down and slowly evaporate at 25 °C, and after 72 h crystals started to form. Single crystals were extracted for SCXRD. Compound 3: colorless plates (ACN); yield: 74%; mp 288–291 °C;

2.1.4. Synthesis of 1-(2-Oxo-2-Phenylethyl)-4-(Pyrrolidin-1-yl) Pyridin-1-Ium Bromide, Compound 4

4-pyrrolidino pyridine (74 mg, 0.5 mmol) and 2-bromacetophenone (100 mg, 0.5 mmol) were dissolved in 5 mL ACN. After constant stirring at 80 °C in inert (argon) atmosphere for 24 h, the solution turned orange and was left to cool down and slowly crystalize at room temperature. Large orange crystals, formed within a week, were filtered and washed with acetonitrile. Compound 4: orange blocks (ACN) yield: 70%; mp 266-271 °C;

2.1.5. Synthesis of 1-(2-(2,4-Dimethoxyphenyl)-2-Oxoethyl)-4-(Pyrrolidin-1-yl)Pyridin-1-Ium Bromide, Compound 5

4-pyrrolidino pyridine (74mg, 0.5 mmol) and 1-[[1,1'-biphenyl]-4-yl]-2-bromoethan-1-one (138 mg, 0.5 mmol) were dissolved in 5 mL acetonitrile. The solution was stirred for 24 h at 80 °C resulting in a pinkish white precipitate that was filtered and washed with acetonitrile. Orange single crystals suitable for SCXRD were obtained after recrystallization from methanol. Compound 5: orange blocks (methanol); yield 69%; mp 288-291 °C;

2.1.6. Synthesis of 1-Butyl-4-(Pyrrolidin-1-yl) Pyridin-1-Ium Iodide, Compound 6

4-pyrrolidino pyridine (74 mg, 0.5 mmol) and butyl iodide (92 mg, 57ul, 0.5 mmol) were mixed in 5 mL ACN. The reaction was performed with constant stirring at temperature near the boiling point of the acetonitrile and in inert (argon) atmosphere for 24 h. The solution was vacuum evaporated
and the precipitate recrystallized from methanol/aceton solution resulting in light yellow crystals. Compound 6: light yellow blocks (methanol/aceton); yield: 65%; mp 150–152 °C; $^1$H NMR (DMSO-$d_6$, 600 MHz) $\delta$ 0.871 (3H, $t$, $J = 7.4$ Hz, CH$_3$ Bu), 1.211 (2H, $m$, CH$_2$-3 Bu), 1.172 (2H, $m$, CH$_2$-2 Bu), 1.982 (4H, $m$, CH$_2$-3 and CH$_2$-4 Pir), 3.469 (4H, $m$, CH$_2$-2 and CH$_2$-5 Pir), 4.143 (2H, $m$, CH$_2$-1 Bu), 6.873 (2H, dd, $J = 7.7$ Hz, 1.8 Hz, CH-3 and CH-5 Pyr), 8.284 (2H, dd, $J = 7.7$ Hz, 1.8 Hz, CH-2 and CH-6 Pyr); $^{13}$C NMR (DMSO-$d_6$, 151MHz) $\delta$ 13.86 (CH$_3$ Bu), 19.14 (CH$_2$-3 Bu), 25.18 (CH$_2$-3 and CH$_2$-4 Pir), 32.82 (CH$_2$-2 Bu), 48.76 (CH$_2$-2 and CH$_2$-5 Pir), 56.93 (CH$_2$-1 Bu), 108.77 (CH-3 and CH-5 Pyr), 142.45 (CH-2 and CH-6 Pyr), 153.34 (C$_q$-4 Pyr).

2.2. Characterization Methods

2.2.1. Single Crystal X-ray Diffraction (SCXRD)

Single crystals of the quaternary ammonium compounds with suitable size and diffracting quality were mounted on glass capillaries or nylon loops. The diffraction peak intensities and coordinates were collected on Agilent SupernovaDual diffractometer equipped with Atlas CCD detector using micro-focus MoKα radiation ($\lambda = 0.71073$ Å). Data were processed with CrvysAlisPro software [37]. The structures were solved with direct methods and refined by the full-matrix least-squares method on $F^2$ with ShelxS and ShelxL programs [38]. All non-hydrogen atoms were located successfully from Fourier map and were refined anisotropically. Hydrogen atoms were placed on calculated positions riding on the parent atom ($U_{eq} = 1.2$ for C–H aromatic = 0.93 Å and C–H methylenic = 0.97 Å). Complete crystallographic data for the structures of compounds 1–6 reported in this paper have been deposited in the CIF format with the Cambridge Crystallographic Data Center as 1992663, 1992664, 1992661, 1992662, 1992660, and 1992665. These data can be obtained free of charge via http://www.ccdc.cam.ac.uk/conts/retrieving.html (or from the CCDC, 12 Union Road, Cambridge CB2 1EZ, UK; Fax: +441223336033; E-mail: deposit@ccdc.cam.ac.uk).

2.2.2. Powder X-ray Diffraction (PXRD)

Powder samples of the synthesized quaternary ammonium compounds were analyzed on Empyrean Powder X-ray diffractometer (Malvern Panalytical, Netherlands) in the range 2°–50° 2θ using Cu radiation ($\lambda = 1.5406$ Å) and PIXcel3D detector. The diffraction patterns of the precipitates were compared with those generated from the SCXRD experiment to confirm the presence or absence of additional phases.

2.2.3. Nuclear Magnetic Resonance (NMR) Spectroscopy

The NMR spectra were recorded on a Bruker Avance II+ 600 spectrometer (Rheinstetten, Germany), $^1$H at 600 MHz and $^{13}$C at 151 MHz, in DMSO-$d_6$; the chemical shifts were quoted in ppm in $\delta$-values against the solvent peak and the coupling constants were calculated in Hz. The spectra were processed with Topspin 2.1 program. For simplicity, pyridyl ring nuclei are depicted as Pir, pyrroolidine unit as Pir, and N$^+$-substituents, respectively, as Bu (butyl), Ph (phenyl), BiPh (biphenyl), and Ar (3-NO$_2$C$_6$H$_4$, 2,4-(OMe)$_2$C$_6$H$_3$ and 3,4-(O–CH$_2$CH$_2$CH$_2$–O)–C$_6$H$_3$). The carbon resonances of compounds 1–3 were extracted from 2D heterocorrelations HSQC and HMBC.

2.2.4. Differential Scanning Calorimetry (DSC)

DSC analyzes were performed on Discovery DSC 250 (TA instruments). Samples weighing between 1 and 5 mg were heated in aluminum pans from 20 to 350 °C (10 °C·min$^{-1}$) in argon (flow rate 10 mL·min$^{-1}$). Melting points of the synthesized compounds were determined from the DSC experiments.
2.2.5. Disk-Diffusion Method (Kirby-Bauer Test)

The susceptibility of the selected bacterial strains (Klebsiella pneumoniae subspecies ozaenae- (CNCTC 1/35 SZU), Pseudomonas aeruginosa (CNCTC Ps 41/65), Escherichia coli, Staphylococcus aureus—ATCC 6538 and Bacillus subtilis) against the synthesized quaternary ammonium compounds (1–6) were evaluated using the Kirby-Bauer test. Solutions of compounds 1–6 with concentration of 1 mg/mL in 2% DMSO were prepared. A volume, corresponding to 30 µL of the tested compound was pipetted on sterile cellulose discs (6 mm in diameter). Warm (50–55 °C, ~25 mL) Mueller-Hinton agar (containing ~1.5% agar) medium was poured into 90 or 60 mm sterile Petri dishes. The prepared dishes were left to cool down and solidify for 6 h. Bacterial suspensions in liquid growth medium (Lysogeny broth) with concentrations of ~1.10⁶ cfu·mL⁻¹ (0.5 McFarland standard) were prepared. Small amounts (~0.1–0.2 mL) of each bacterial suspension were pipetted on the solid agar dishes and gently spread across the surface using cell spreader. After what dried cellulose discs with studied compounds were placed on the surface of the solidified growth medium. The plates were then incubated at 37 °C for 12 to 18 h. The diameters of the areas around the discs in which no growth is observed were measured.

3. Results and Discussion

3.1. Single Crystal XRD Studies

As part of the facile purification procedure, colorless (2 and 3), light brown (1) and yellow to orange (4, 5, and 6) single crystals of the title compounds were obtained by slow evaporation method from hot methanol, acetonitrile or methanol-acetone solutions. The 4-pyrrolidino pyridine derivatives crystallize as bromine (1–5) or iodine (6) salts in the monoclinic (S.G. P2₁/c—1, 2 and P2₁/n—3, 5, and 6) or in the orthorhombic (S.G. P2₁2₁2₁, 4) crystal system, bearing one molecule in the asymmetric unit (ASU) and four molecules in the unit cell (Table S1). ORTEP [39] view of the molecules present in the ASU of compounds 1–6 with appropriate labeling scheme are shown on Figure 1. The values for selected bond lengths and angles determined from the SCXRD experiment (Table S2) are in agreement with those of other similar structures in the Cambridge structural database (CCDC, ref. codes: QAWPAG, AGEPOR, EKOJOF [40–42]).

The reported quaternary ammonium bromine salts 1–5 can be described as build-up of two fragments—4-pyrrolidino pyridine and different aromatic substituents (3-NO₂C₆H₄ for 1, 3,4-(O–CH₂CH₂CH₂O)–C₆H₃ for 2, 2,4-(OMe)₂C₆H₃ for 3, phenyl for 4 and biphenyl for 5)—connected via 2-oxoethyl (–CH₂–O) bridge. On the other hand, compound 6 is a 4-pyrrolidino pyridine iodine salt where the aromatic tertiary nitrogen atom is quaternized by an aliphatic butyl substituent. The flexibility of the carbon atoms in the pyrrolidino heterocycle (“half-chair” conformation), leads to a high probability of positional disorder. Indeed, atoms C2 and C3 in the pyrrolidine ring of 1, 2, and 5 are disordered over two position (as shown on Figure 1a,b,e) with occupancy ratio between the different disordered parts: 60%/40% for 1, 58%/42% for 2, and 70%/30% for 5. Other than that, the two cyclic fragments constructing the QAS molecule are relatively planar with RMSD of the 4-pyrrolidino pyridine fragment for all structures varying between 0.073 Å (in 5) and 0.143 Å (in 6) and those of the aromatic substituents between 0.005 Å (for phenyl ring in 4) and 0.326 Å (for 3,4-(O–CH₂CH₂CH₂O)–C₆H₃ substituent in 2). The angle between the mean planes of the two fragments varies significantly with values between 22.09° (for 5) and 91.00° (for 2). The pronounced variations in the above mentioned angles along with the occurrence of single bond between C11 and C12 indicate a good rotational flexibility of the aromatic substituents that can be a crucial part in the antimicrobial activity mechanism. A close inspection of the molecular features of compounds 1–6, reveals the lack of strong proton donating groups (such as NH₂ or OH), required for hydrogen bonding interactions. On the other hand, strong proton accepting groups as C=O (for 1–5) and NO₂ (for 1) or moderately strong ones as C–O–C in compound 2 and –O–CH₃ in 3 are present. Therefore, the formation of typical hydrogen bonding interactions is impossible. Thus, the only possibilities for intra- or intermolecular interactions stabilizing the crystal structures depends on the presence of
halogen (bromine or iodine) anions, e.g., formation of “strong” halogen bonds or weak C–H···O and π–π interactions involving CH₂ or CH groups from the aliphatic or aromatic systems. Indeed, the three dimensional packing of the molecules in the crystal structures of 1–5 discloses a network of strong (H···X distance of 2.69–3.28 Å) halogen bonding interactions and short contacts of C–H_methylenic···X and C–H_aromatic···X type (X = Br⁻ and I⁻, Table S3). Along with the above mentioned H···X interactions, the more common C–H···O interactions involving C–H_methylenic and C–H_aromatic hydrogen atoms and C–O–C and C=O functional groups in 2 and 3 forms R²(8) and R²(10) motifs [43,44] respectively (Figure 2a,b). Compound 6, bearing only 4-pyrrolidino pyridine and butyl fragments, stabilize its crystal structure through a chain of C–H_methylenic···I and C–H_aromatic···I intermolecular interactions forming C₂₁(6) motifs (Figure 2c) [43,44]. In addition, compounds 2 and 3 contain acetonitrile or water molecules (e.g., crystal solvates or hydrates), occupying the cavities between the QASs molecules. The water molecule is involved in hydrogen bonding interaction with Br and thus further stabilize the crystal structure of 3. The solvent acetonitrile in 2 is also involved in the crystal structure stabilization through C–H···X interactions with the Br anion (Figure 2a,b and Table S3). Stacking π–π interactions can be found on three of the reported QASs—1, 3, and 5. Compound 1 has two nearly parallel stacked nitrobenzene rings (the angle between the mean planes of the benzene rings is 0°). The 2,4-dimethoxy benzene and pyridine ring for 3 and the phenyl ring (C18-C19-C20-C21-C22-C23) and the pyridine ring for 5 are stacked with mean planes angles between the rings of 4.015° and 7.155° respectively. The centroid to centroid distances of the stacked rings for 1, 3, and 5 are 3.699 Å, 3.783 Å, and 3.882 Å and the shift distances of centroids are 1.520 Å, 1.345 Å, and 1.838 Å, respectively.

Figure 1. ORTEP [39] views of the molecules in the ASU of the crystal structures of (a) compound 1, (b) compound 2, (c) compound 3, (d) compound 4, (e) compound 5, and (f) compound 6. The thermal ellipsoids are drawn with 50% probability level and all atoms are given with an appropriate labeling scheme.
Figure 2. Halogen bonding interactions and short contacts for compounds 2—(a), 3—(b) and 6—(c). The three-dimensional packing of the molecules of 2, 3 and 5 reveals specific pairing of the halogen bonding interactions between the QASs molecules, forming $R_2^2(8)$, $R_2^2(10)$ and $C_2^1(6)$ motifs [43,44], respectively. The hydrogen atoms are given in light green for better visualization.
3.2. DSC Studies

The DSC data show that the crystal structure is conserved up to the melting point for compounds 1, 4, 5, and 6 (Figure 3 and Supplementary Materials). As mentioned, compounds 2 and 3 are crystal solvates and crystal hydrates with acetonitrile and water molecules present in the crystal structures. The DSC thermograms reveal that the water and acetonitrile molecules evaporate with broad endo effects covering the 67 to 97 °C and 100 to 112 °C temperature regions, respectively (Figure 3). The room temperature PXRD of compound 2 (Figure S2) showed a diffraction peak at ~5.5° 2θ that could not be related to the diffraction of the starting compounds or the generated pattern from single crystal data. Based on the DSC data we heated (e.g., dried) compound 2 for 1h at 90 °C and collected PXRD again. The observed diffracting pattern was completely different from the initial one but replicated the peak at 5.5° 2θ. Thus, the peak at 5.5° 2θ observed in the initial PXRD may be associated with the crystal phase without acetonitrile (most likely obtained during the grinding of the crystals). The PXRD results for 2 correlate well with the 1H-NMR studies which disclose only one molecular structure in solution and high purity of the compound. The melting points of 1–6 are spread in the 150–300 °C region. Interestingly, compounds 1–6 behave differently after they are melted. Some, such as 1, 3, 4, and 5, start to decompose immediately or slightly after (interval of 10–15 °C) they melt while the decomposition of 2 and 6 occurs more than 120 °C after the melting. The choice of a simple and cost-effective synthesis and purification procedures giving good yields and allowing the isolation of sufficiently pure compounds is supported by the DSC data revealing a sharp and narrow endo effects related to the melting points and confirmed by the NMR and PXRD data not showing traces of the starting organic compounds.

Figure 3. Combined graphical representation of the results from the DSC analyzes for compounds 1-6 with appropriate color and symbol scheme. The maxima of the endo effects, corresponding to the melting temperatures (T_m) for all compounds, are denoted.

3.3. Antibacterial Susceptibility Testing and Activity

In our previous work [36] the antibacterial activity of 1-(2-(naphthalen-2-yl)-2-oxoethyl)-4-(pyrrolidin-1-yl) pyridin-1-ium bromide (CCDC 1875089) was checked against Klebsiella pneumoniae, Escherichia coli, Staphylococcus aureus and Bacillus subtilis and showed inhibitory activity only against S. aureus (inhibition zone of 16 mm). Based on the exhibited specific activity, we decided to perform some additional random screening with different substituents
to check if the activity is solely due to the 4-pyrrolidino pyridine moiety or is a combination between the different substituents.

Compounds 1–6 have been screened for antibacterial activities using disk diffusion method by measuring the inhibition zone in mm. Kanamycin (30 μg) was used as standard drug for antibacterial activity. The initial investigation of the antibacterial activity showed that only two of them, namely 3 and 5, exhibit comparatively good inhibitory effect against Escherichia coli and Staphylococcus aureus, respectively. Based on the screening the antibacterial experiments for 3 and 5 against E. coli and S. aureus, respectively, were performed three times with consistent antibacterial effect (Figure 4). The measured inhibition zone of 3 and 5 (given as average of several measurements, Table S4) are comparable to the kanamycin reference with values of 18 mm vs 21 mm and 14 mm vs 16 mm, respectively. The factors controlling antimicrobial activity of the substituted pyridinium salts are the hydrophobicity, adsorbability, surface activity, position of the substituents and electron density of the ammonium nitrogen atom. In order to design novel effective antimicrobial agents, we need to understand the structure activity relation of the different factors. Here all six compounds have an identical 4-pyrrolidino pyridine moiety containing the quaternary nitrogen atom and thus the differences of the observed effects may be explained by the different substituent and the different hydrophobicity of the compounds. The logP values for compounds 1–6 are 2.21 (±1.61), 2.37 (±1.78), 3.87 (±1.78), 2.14 (±2.1), 3.81 (±2.21), and 2.09 (±2.38), respectively [45]. Indeed, the values for compound 3 (as hydrate) and 5 differ from the remaining trend (~2.2) suggestion an easier passage through the cell membrane. Interestingly the overlay of the molecules present in the ASU of 3 and 5 did not reveal drastic differences. The main variation occurs in the respective orientation of the 2,4-dimethoxyphenyl and 1,1’-biphenyl]-4-yl aromatic rings (Figure 5). The overlay of 3 with compounds 1, 2, 4, and 6 also discloses a highly conserved geometry of the 4-pyrrolidino pyridine moiety and small variation in the rotational orientation of the substituents. Thus, the bacterial specificity to E. coli and S. aureus should be explained by the different substituents: 2,4-dimethoxyphenyl and 1,1’-biphenyl]-4-yl. One should note that the specificity is not Gram-positive or Gram-negative related because compounds 3 and 5 are not inhibiting the growth of the other G+ (Bacillus subtilis) and G− (Klebsiella pneumoniae subsp. ozaenae, Pseudomonas aeruginosa) bacteria in the test set. Thus, compounds 3 and 5 are suitable for targeting of a specific bacterial species.

Figure 4. Observed antibacterial effect (experiments were repeated three times) corresponding to 30 μg of compound 5 against S. aureus (a) and compound 3 against E. coli (b), compared to 30 μg kanamycin (reference). The diameter of the area around the discs, containing the tested compounds, in which no visible growth is observed (zone of inhibition) was measured using ImageJ software [46].
Figure 5. Overlay of the molecules present in the ASU of compounds 3 (red) and 5 (blue) in two orientations (rotated at 90°) showing the conserved geometry of the 4-pyrrolidino pyridine moiety and the rotational freedom around the C11–C12 bond; hydrogen atoms have been omitted for clarity.

4. Conclusions

We demonstrated a facile synthesis and purification procedure of quaternary ammonium derivatives of 4-pyrrolidino pyridine containing different aromatic substituents. In all cases, the purification of the compounds by simple washing with solvent and/or recrystallization (methanol, acetonitrile or methanol/acetone) produces reasonable yields (60–74%). The compounds were screened for antimicrobial activity against Gram-positive and Gram-negative bacterial strains (five in total) and two of the compounds, 1-(2-(2,4-dimethoxyphenyl)-2-oxoethyl)-4-(pyrrolidin-1-yl)pyridin-1-ium bromide and 1-(2-[[1,1’-biphenyl]-4-yl]-2-oxoethyl)-4-(pyrrolidin-1-yl) pyridin-1-ium bromide, showed inhibition activity similar or better than the kanamycin control. Single crystal analysis revealed that in all compounds, H···X halogen along with π–π interactions, are responsible for the stabilization of the crystal structures. The 4-pyrrolidino pyridine moiety displays a highly conserved structural geometry and the observed differences in the antimicrobial activity arise from the different hydrophobicity while the specificity is due to the aromatic substituents. The results show that the observed activities are not solely linked to the 4-pyrrolidino-pyridinium fragment and that additional synthesis of various wisely designed compounds need to be undertaken.

Supplementary Materials: The following are available online at http://www.mdpi.com/2073-4352/10/5/339/s1; Figures S1, S3, S4, S5, and S6: Comparison of the Powder X-ray diffraction pattern (blue) versus Single Crystal X-ray diffraction pattern (generated via Mercury software, red) for compounds 1, 3, 4, 5, and 6, respectively; Figure S2: Powder X-ray diffraction patterns of compound 2 as crystallized at room temperature (blue) versus Single Crystal X-ray diffraction pattern (generated via Mercury software, red) for compound 2, starting compound 1 (green), starting compound 2 (yellow) and after heating at 90 °C (magenta); Figures S7, S8, S9, S10, S11, and S12: 1H NMR spectra of compounds 1–6, respectively; Figures S13, S14, S15, S16, S17, and S18: 1H–1H NOESY spectra of compounds 1–6, respectively; Figure S19: 1H–1H COSY spectrum of compound 2; Figures S20, S21, S22, S23, S24, and S25: 1H–13C HSQC spectrum of compounds 1–6, respectively; Figures S26, S27, S28, S29, S30, and S31: 1H–13C HMBC spectrum of compounds 1–6, respectively; Figures S32, S33, and S34: 13C NMR spectrum of compounds 4–6, respectively; Figures S35, S36, S37, S38, S39, and S40: Observed thermal effects from the DSC analysis of compounds 1–6, respectively; Figure S41: Observed low antibacterial effect for compounds 1–6 against P. aeruginosa—a, K. pneumoniae—b, and B. subtilis—c, compared to kanamycin reference; Figure S42: Observed antibacterial effect for compounds 1–6 against E. coli—a, and S. aureus—b compared to kanamycin reference; Table S1. The most important parameters from the SCXRD experiment of compounds 1–6, Table S2: Selected bond lengths and angles determined from the SCXRD experiment of compounds 1–6, Table S3: Halogen bonding interactions and short contacts for the crystal structures of 1–6. Table S4: Measured inhibition zone for compounds 3 and 5 against E. coli and S. aureus.

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