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Synthesis, spectroscopic characterization, crystal structure, Hirshfeld surface analysis and antimicrobial activities of two triazole Schiff bases and their silver complexes

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Abstract

The present work describes the synthesis, structural characterization, Hirshfeld analysis, and antibacterial assays of two hydroxy Schiff bases with triazole moiety: *N*-[(*E*)-(3-hydroxy)methylidene]-4*H*-1-2-4 triazol-4-amine (**L1**) and *N*-[(*E*)-(2,4-dihydroxy)methylidene]-4*H*-1-2-4 triazol-4-amine (**L2**). The interaction of these two Schiff bases with AgNO₃ leads to Ag(I) dimer [Ag(**L1**)](NO₃) in the case of the first ligand (**L1**) and polymer [Ag(**L2**)]_n(NO₃)_n(H₂O) for (**L2**). Crystal structure determination and Hirshfeld analysis reveal the presence of two important hydrogen bond interactions in the four compounds, O...H and H...N, involving nitrogen atoms of triazole ring. Contribution of these two types of hydrogen bonds evolves inversely from (**L1**) which contains one –OH, to (**L2**) containing two –OH, then to [Ag(**L1**)](NO₃) dimer and finally to [Ag(**L2**)]_n(NO₃)_n(H₂O) polymer. Indeed, we observed an increase of the percentage to the Hirshfeld surfaces of O...H contribution from (**L1**) to [Ag(**L2**)]_n(NO₃)_n(H₂O) passing by (**L2**) and [Ag(**L1**)](NO₃), while the percentage of N...H decrease in the same way. Antibacterial activity of the free ligands (**L1**), (**L2**) and their silver complexes [Ag(**L1**)](NO₃) and [Ag(**L2**)]_n(NO₃)_n(H₂O) were evaluated against four Gram-negative (*E. coli*, *Salmonella typhimurium*, *Klebsiella*

pneumoniae and *serratia marcescens*) and two Gram-positive (*Staphylococcus aureus*, *streptococcus*) bacteria strains, using the standard antibiotics, Colistin, Fosfomycin, Gentamycin, Erythromycin, Rifampicin, Amikacin, Ofloxacin and Amoxicillin as positive controls. AgNO₃ salt was also used as second control test. The results of this evaluation reveal that, while the two Schiff bases inhibit the growth of the *Serratia Marcescens* bacterium only, the two Ag(I) complexes are active for all bacteria, with activities comparable to the ones of silver nitrate. However, [Ag(L2)]_n(NO₃)_n(H₂O) complex appears significantly more efficient than the AgNO₃ salt against *Klebsiella Pneumoniae*.

Keywords

Schiff bases; silver complexes; hydrogen bonding; Hirshfeld surfaces; antimicrobial activity

1. Introduction

Schiff bases *i.e.* R₂C=NR' (R and R'≠H), are an important class of organic compounds reported for the first time in 1864 by Hugo Schiff [1]. This family of compounds is used as starting materials in the synthesis of important drugs, such as antibiotics and antitumor substances [2-4]. A number of these bases have been reported to possess antioxidant, [5-9] anticonvulsant, [10] analgesic, [11] antitubercotic, [12] and diuretic [13] activities. Schiff bases have often been used as chelating ligands in the field of coordination chemistry for obtaining metal-based supramolecular assemblies, with a large number of potential applications such as catalysis, [14-15] photochemistry, [16] luminescence, [17] sensing, [18] magnetism, [19-20] gas storage, [21-23] gas purification, [24-25] and medicine [26-30].

Our interests have been in the synthesis, the characterization and the biological properties of heterocyclic Schiff bases and their Ag(I) complexes, where heterocycles are chosen as 1-2-4 triazole groups. Triazole moieties are indeed known as important polar pharmacophore able to act either as hydrogen bond donor or acceptor, and thus ready to interact with a variety of partners in biological systems. Triazole heterocyclic rings are thus considered as pharmacologically important scaffolds and are indeed used in many drugs associated to diverse biological activities [31, 32]. On the other hand, ionic silver is known for its antimicrobial properties for many centuries. Even if the precise mechanisms by which it is able to inhibit bacterial growth is still studied nowadays, it is commonly admitted that silver can kill bacteria by entering cells through impairment of enzymes binding Ag(I) ions on the surface of cell wall. Once inside the cell, proposed mechanisms range from the deactivation or

the denaturation of essential enzymes by interaction between silver and sulfhydryl-containing molecules, to DNA denaturation by intercalation between base pairs. In any case, these mechanisms lead to a deep disruption of the cell metabolism and, finally, to cell death [33, 34]. Despite the historical knowledge of the antimicrobial properties of silver ions, the discovery and the development of antibiotics significantly reduced the search of new silver-based bactericidal agents and, overall, the use of silver in medicine. Nowadays the use of silver in medicine is in practice limited to burns treatment with silver sulfadiazine and wound management, for instance using dressings or medicinal devices coated with silver nanoparticles [35]. However, the emergence of antibiotic resistant bacteria strains and of multidrug-resistant organisms quite recently changed the situation and prompted the search of new antimicrobial drugs relying on the well-known properties of silver, such as silver coordination compounds. For instance, the literature is abundant concerning the development of silver complexes with N-heterocycles (including triazoles), or N-heterocyclic carbenes [36], and the evaluation of their antibiotic properties. In addition to the evaluation of biological activities of our triazole substituted Schiff bases and their Ag(I) complexes, this study can be a plus for the construction of metal-organic frameworks (MOFs) with novel topology and for the crystal engineering of molecular architectures organized by coordination bonds and supramolecular contacts (such as hydrogen bonding, π - π interactions). Indeed, Ag(I) is a favourable building block for constructing MOFs because of its flexible coordination numbers and weak metal-metal or metal-ligand interactions [37-38].

2. Experimental

2.1. Reagents and measurements

All reagents were commercially available, purchased from Sigma-Aldrich, and used without further purification. The IR spectra were recorded on a Nicolet 5700 FT-IR Thermo Scientific spectrophotometer using powder in mixed KBr pellets in the range of 400-4000 cm^{-1} . The UV-Vis spectra were recorded at room temperature in the range of 100-900 nm, on a Cary 4000 Varian UV-Visible spectrophotometer by dissolving the compounds in DMSO.

2.2. Procedure of synthesis of the two Schiff bases and their silver complexes

2.2.1. Synthesis of *N*-[(*E*)-(3-hydroxyphenyl)methylidene]-4H-1,2,4-triazol-4-amine. (L1):

3-Hydroxybenzaldehyde (0.244 g, 2 mmol), was added to an ethanolic solution (15 ml) of 4-amino-4H-1,2,4-triazole (0.161 g, 2 mmol) containing few drops of glacial acetic acid. The reaction mixture was stirred for 6h under reflux at 75°. The resulting yellow solution was

cooled in an ice bath. The crystalline powder obtained was filtered off, washed with ethanol and dried under vacuum. Pure colourless crystals for (**L1**) diffraction were obtained by recrystallization in acetonitrile/H₂O. ¹H NMR (250 MHz, [D6] DMSO, 300K): δ =9.04 (2H, s, H₁, H₂), 8.85 (1H, s, H₃), 7.36 (1H, m, H₈), 7.31 (1H, m, H₅), 7.26 (1H, m, H₇), 6.89 (1H, m, H₉) ppm. ¹³C NMR (250 MHz, [D6] DMSO, 300K) δ = 159.76 (C₆), 158.27 (C₃), 140.09 (C₁-C₂), 134.00 (C₈), 131.50 (C₉), 121.69 (C₅), 120.83 (C₇), 115.07 (C₄) ppm. (**supp.1**). C₉H₈N₄O (188.19): calcd. C 57.44, H 4.28, N 29.77, O 8.50; found C 57.54, H 4.35, N 29.90, O 8.55.

2.2.2. Synthesis of *N*-[(*E*)-(2,4-dihydroxyphenyl)methylidene]-4H-1-2-4 triazol-4-amine. (**L2**) :

The pure crystals of second Schiff base (**L2**) was synthesized by the same procedure used for (**L1**). However, in this case, we use 2,4-dihydroxybenzaldehyde (0.276 g, 2 mmol) instead of 3-hydroxybenzaldehyde. ¹H NMR (250 MHz, [D6] DMSO, 300 K): δ = 8.98 (2H, s, H₁, H₂), 8.89 (1H, s, H₃), 7.56 (1H, d, *j*=8.6 Hz, H₉), 6.41 (1H, dd, *j*= 8.6 Hz, 2.0 Hz, H₈), 6.34 (1H, d, *j*=2.0 Hz, H₆) ppm. ¹³C NMR (250 MHz, [D6] DMSO, 300K) δ = 163.80 (C₅), 161.12 (C₇), 158.43 (C₃), 139.90 (C₁-C₂), 132.37 (C₉), 110.57 (C₆), 109.75 (C₈), 103.57 (C₄) ppm. (**supp.2**). C₉H₈N₄O₂ (188.19): calcd. C 52.89, H 3.92, N 27.43, O 15.67; found C 53.00, H 3.99, N 27.44, O 15.80.

2.2.3. Synthesis of [Ag(**L1**)](NO₃) and [Ag(**L2**)]_n(NO₃)_n(H₂O) :

Complexes [Ag(**L1**)](NO₃) and [Ag(**L2**)](NO₃)(H₂O) were prepared by the reaction of (**L1**) and (**L2**) respectively with AgNO₃ (0.3:0.3 mmol) in 14 ml of H₂O/CH₃CN (1:1) solution. After stirring for about 30 min at 50 °C, the resulting solutions were filtered. Colourless single crystals suitable for X-ray diffraction were obtained by slow evaporation of the two solutions after few days. For [Ag(**L1**)](NO₃): AgC₁₈H₁₆N₉O₅ (546.27): Ag 19.75 C 39.54, H 2.93, N 23.06, O 14.64; found Ag 19.90 C 39.54, H 3.05, N 23.09, O 14.70. For [Ag(**L2**)](NO₃)(H₂O): AgC₉H₁₀N₅O₆ (392.09): Ag 27.51 C 27.54, H 2.55, N 17.85, O 24.48; found Ag 27.60 C 27.56, H 2.80, N 17.95, O 24.67.

2.3. X-ray crystallography

A suitable crystal of each compound was glued to a glass fiber mounted on dual source supernova with the large area Atlas CCD detector. Diffraction data were collected using the Mo-K α radiation at 100 K. Correction for Lorentz-polarization effect, peak integration and

reduction were carried out with CrysAlisPro package [39]. The structures were solved by direct methods and refined by full-matrix least squares method using SHELX programs [40]. All non-hydrogen atoms are placed appropriately and refined anisotropically. All H atoms were located in difference electron-density maps and were treated as riding on their parent atoms, with C-H = 0.93 Å, N-H = 0.85 Å, O-H = 0.84 Å and $U_{\text{iso}}(\text{H}) = -1.2U_{\text{eq}}(\text{X})$ (X = C and N) and $-1.5 U_{\text{eq}}(\text{O})$.

The crystallographic data for the structures reported in this paper have been deposited with the Cambridge Crystallographic Data Centre as supplementary publications no. CCDC 1582965 - 1582968. Copies of available material can be obtained, free of charge, on application to the Director, CCDC, 12 Union Road, Cambridge CB2 1EZ, UK, fax: +44 (0)1223 336033 or Email: deposit@ccdc.cam.ac.uk. Relevant crystal data are listed in **table 1** together with refinement detail.

Table 1. Crystal data, data collection and structure refinement for the four title compounds

Crystal data	(L1)	(L2)	[Ag(L1)](NO ₃)	[Ag(L2)](NO ₃)(H ₂ O)
Chemical formula	C ₉ H ₈ N ₄ O	C ₉ H ₈ N ₄ O ₂	C ₁₈ H ₁₆ AgN ₈ O ₂ . NO ₃	C ₉ H ₈ AgN ₄ O ₂ . NO ₃ . H ₂ O
M_r	188.19	204.19	546.27	392.09
Cell setting, space group	Orthorhombic, $Pna2_1$	Monoclinic, $P2_1/n$	Triclinic, $P\bar{1}$	Monoclinic, $P2_1/n$
Temperature (K)	100 (2)	100 (2)	100 (2)	100 (2)
Unit cell parameters	$a = 22.7284(5)$ Å $b = 9.8594(2)$ Å $c = 3.7808(1)$ Å $\alpha = 90^\circ$ $\beta = 90^\circ$ $\gamma = 90^\circ$	$a = 5.8440(5)$ Å $b = 21.118(2)$ Å $c = 7.5764(8)$ Å $\alpha = 90^\circ$ $\beta = 107.094(5)^\circ$ $\gamma = 90^\circ$	$a = 8.751(5)$ Å $b = 9.625(5)$ Å $c = 12.313(5)$ Å $\alpha = 90.081(5)^\circ$ $\beta = 97.473(5)^\circ$ $\gamma = 99.838(5)^\circ$	$a = 12.5270(4)$ Å $b = 8.2643(3)$ Å $c = 12.5996(3)$ Å $\alpha = 90^\circ$ $\beta = 103.151(3)^\circ$ $\gamma = 90^\circ$
V (Å ³)	847.23(3)	893.74(15)	1012.9(9)	1270.16(7)
Z	4	4	2	4
D_x (Mg m ⁻³)	1.475	1.518	1.791	2.050
Radiation type	Cu $K\alpha$	Mo $K\alpha$	Mo $K\alpha$	Mo $K\alpha$
μ (mm ⁻¹)	0.86	0.11	1.05	1.63
Crystal form, colour	Prism, colourless	Prism, colourless	Prism, colourless	Prism, colourless
Crystal size (mm)	0.15 × 0.10 × 0.09	0.20 × 0.10 × 0.10	0.10 × 0.10 × 0.08	0.20 × 0.10 × 0.10

Diffractometer	Agilent SuperNova CCD diffractometer	Agilent SuperNova CCD diffractometer	Agilent SuperNova CCD diffractometer	Agilent SuperNova CCD diffractometer
Data collection method	ω -scans	ω -scans	ω -scans	ω -scans
Absorption correction	Integration	Integration	Integration	Integration
$T_{\min}$, $T_{\max}$	0.908, 0.938	0.889, 0.940	0.708, 0.750	0.896, 0.946
No. of measured, independent and observed reflections ($I > 2\sigma(I)$)	15602, 1787, 1574	8605, 2267, 1440	21173, 6041, 4082	32791, 6508, 4856
$R_{\text{int}}^{\text{a}}$	0.059	0.049	0.037	0.036
$\theta_{\max}$ ($^{\circ}$)	76.5	28.6	30.5	37.7
Refinement on	F^2	F^2	F^2	F^2
$R[F^2 > 2\sigma(F^2)]^{\text{b}}$, $wR(F^2)^{\text{c}}$, S^{d}	0.053, 0.135, 1.04	0.057, 0.200, 1.08	0.041, 0.164, 1.09	0.029, 0.070, 1.05
No. of reflections	1663 reflections	2267 reflections	6041 reflections	2815 reflections
No. of parameters	127	136	298	167
$(\Delta/\sigma)_{\max}$	< 0.01	< 0.01	0.001	0.001
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ ($\text{e } \text{\AA}^{-3}$)	0.60, -0.32	0.42, -0.37	0.63, -0.93	0.75, -0.83

$$^{\text{a}} R_{\text{int}} = \Sigma(F_o^2 - \langle F_o^2 \rangle) / \Sigma(F_o^2)$$

$$^{\text{b}} R(F^2) = \Sigma||F_o| - |F_c|| / \Sigma|F_o|, \quad ^{\text{c}} wR(F^2) = [\Sigma w(F_o^2 - F_c^2)^2] / [\Sigma w(F_o^2)^2]^{1/2}.$$

$$^{\text{d}} \text{Goodness-of-fit } S = [\Sigma w(F_o^2 - F_c^2)^2 / (n - p)]^{1/2}, \text{ where } n \text{ is the number of reflections and } p \text{ the number of parameters.}$$

2.4. Hirshfeld surface analysis

In order to have a better understanding of the intermolecular interactions within the crystal structure, Hirshfeld surface analysis [41] has been undertaken. It consists in partitioning the crystal electron density into molecular fragments. Thus, the Hirshfeld surface can be defined as the border between regions where promolecule electron density surpasses that from all neighbouring molecules over the rest of the crystal. The Hirshfeld surface, which envelops the molecule, is defined as an iso-surface of $w(r) = 0.5$, where $w(r)$ is a molecular weight function derived from the spherically averaged atomic electron densities $\rho_i(r)$ [42]:

$$w(r) = \frac{\sum_{i \in \text{molecule}} \rho_i(r)}{\sum_{i \in \text{crystal}} \rho_i(r)}$$

The normalized contact distance d_{norm} is the sum of the distances of any surface point to the nearest interior (d_i) and exterior (d_e) atom, normalized by the van der Waals radii of the atoms, Spackman et al. [43]. The Hirshfeld surfaces were computed to explore the contributions of non-covalent interactions in the crystal structures using Crystal Explorer 3.1 software [44].

3. Results and discussion

FT-IR spectroscopy:

The spectrum of the triazole Schiff bases, (**L1**) and (**L2**), exhibit remarkable bands at 1611 and 1605 cm^{-1} respectively due to the formation of the imine group ($\text{C}=\text{N}$) [45-46], **supp.3** (supplementary materials). Furthermore, the $\nu(\text{O-H})$ bands of the weak intensity were observed in the range 3100-3150 cm^{-1} . These bands do not shift to lower wave number in the spectra of the complexes indicating the non-participation of the imine nitrogen and the hydroxyl group in coordination to the metal ion [47]. In addition, the bands detected in the range of 1300-1500 cm^{-1} is due to the $\nu(\text{N-N})$ stretching vibration of the triazole moiety. This band is shifted to higher wave number in the complex spectra suggesting the participation of the two nitrogen atoms N1 and N2 of triazole in coordination to the metal ion [48].

UV-visible spectroscopy:

The UV-visible absorption of the four compounds display two bands observed in DMSO in the range of 260-400 nm. The first band observed around 320 nm is owing to the low energy $\pi - \pi^*$ transition of the imine group. The second band detected around 280 nm was assigned as a moderate $\pi - \pi^*$ transition of the aromatic ring **table S1** and **supp.4** (supplementary materials). These bands undergo a hypsochromic shift due to the coordination of the free ligand with the metal ion.

X-ray structure analysis.

Crystal structure of (L1) and (L2)

The crystal structure of ligand (**L2**) has been reported by François Safin and co-workers [49], however, the data collection was done at room temperature on a twinned crystal, with

insufficient data to have a good refinement of the structure (*Ratio NO/NV*: 6.8). These constraints affected the accuracy of the values of distances and angles.

The molecular structures of the two free ligands (**L1**) and (**L2**) are depicted in **Fig.1**. The two compounds crystallize in the orthorhombic group space $Pna2_1$ for (**L1**) and in the monoclinic group space $P2_1/n$ for (**L2**) with four molecules in the unit cell for each compound. The two Schiff bases (**L1**) and (**L2**) differ only by the positions and the nature of the substitutions on the six-membered aromatic ring: (**L1**) bears a single hydroxyl group at the C(6A) meta position while (**L2**) presents two hydroxyl substituent at the ortho and para positions. Both (**L1**) and (**L2**) are in their enolimine form, confirmed by the lengths of C-O bonds (1.368(3), 1.366(3) and 1.348(8) Å), and of C-C single bonds (1.469(3) and 1.451(3) Å) and C=N imine bond (1.278(3) and 1.279(3) Å) (**Fig.1**). The two molecules have an almost planar *E* configuration with respect to the imine C=N double bond, as indicated by the value of $-177.1(2)^\circ$ for (**L1**) and $177.21(17)^\circ$ for (**L2**) of the N(3A)-N(4A)-C(3A)-C(4A) torsion angle. The dihedral angles between aromatic rings in both ligands are comparable: 17.16° for (**L1**) and 19.51° (**L2**). The bond angles and distances in (**L1**) and (**L2**) are in agreement with similar compounds [50-52].

(a)

(b)

Fig.1: ORTEP of the two ligands (**L1**) and (**L2**) with thermal displacement ellipsoids drawn at the 50% probability level. (a) (**L1**) and (b) (**L2**).

In the crystal structures formed by (**L1**) and (**L2**), the nitrogen atoms of triazole moieties interact with hydroxyl group through intermolecular O-H...N hydrogen bonds. In the case of (**L1**), the single hydroxyl group is hydrogen bonded to N(1A) allowing to form infinite chains along *a* axis with a C(10) graph set motif [53-54]. The presence of the second OH in the case of (**L2**) has drastic consequences for the crystal packing. It leads indeed to the formation of another O-H...N hydrogen bond involving the second triazole nitrogen atom, transforming the infinite chain observed in the case of (**L1**) into $R_4^4(29)$ rings, generating two-dimensional layers developing parallel to the (101) plane (**Fig.2** and (**table S2** in supplementary materials)). These hydrogen bonds replace the intramolecular ones between *o*-OH and imine nitrogen atom which are broken by the rotation of the phenolic ring [55].

(a)

Fig.2: O-H...N hydrogen bonds. *(a)* Infinite chains in **(L1)** and 2D layers *(b)* in **(L2)**

In **(L1)**, the azomethine nitrogen (N4A), which is not involved in O-H...N interaction, acts nevertheless as acceptor in a C4A-H4A...N4A hydrogen bond along *b* axis, giving rise to 2D layers built on the basis of alternating $R_6^6(33)$ and $R_3^3(18)$ rings, (**Fig.3a**). Moreover, along the third direction (*c* axis), C1A-H1...O1A bonds between triazole and hydroxyl, and C2A-H2A...N1A bond between two triazole moieties ensure the supramolecular cohesion of the 2D layers, (**Fig.3b**). The same C-H...O hydrogen bond links the 2D layers in **(L2)** along the *b* axis and leads to a three dimensional structure. In both crystal structures, $\pi\cdots\pi$ interactions coexist with hydrogen bonding and participate in the final stability of the molecular assemblies. As shown in **Fig.4**, in the case of **(L1)**, the crystal packing presents clear triazole ... triazole and phenyl...phenyl $\pi\cdots\pi$ interactions. In the case of **(L2)**, a single weaker $\pi\cdots\pi$ stacking appears to form partially between the phenyl ring of a molecule and both the triazole moiety and the imine double bond of the symmetry equivalent molecule obtained by translation along the *a* axis.

Fig 3: C-H...N and C-H...O hydrogen bonds in (**L1**). (a) 2D layers generated by interactions involving azomethine nitrogen atoms. (b) Supramolecular structure of (**L1**).

Fig.4: $\pi \dots \pi$ interactions.

(a) phenyl...phenyl and triazole...triazole in (**L1**). (b) phenyl...triazole in (**L2**)

Crystal structure of the silver complexes [Ag(L1)](NO₃) and [Ag(L2)]_n(NO₃)_n(H₂O)

Reaction of AgNO₃ with (**L1**) and (**L2**) yield a Ag(I) dimer for the first ligand and a polymer for the second one. As observed in the crystal structures of the isolated ligands, the presence of an extra hydroxyl group in (**L2**) compared to (**L1**) led to very different crystal structures of their corresponding silver complexes. The structure of the dimer issued by (**L1**) contains a dinuclear cation formed by two-symmetry related Ag(I) coordinated to four (**L1**) ligands, [Ag₂(**L1**)₄], and uncoordinated NO₃⁻ anions, **Fig.5**. Each Ag(I) ion adopts a trigonal environment and the sum of angles around it, equal to 360.0°, indicates planarity of its geometry [56-57]. Ag(I) and Ag(I)(i) [*i*] 2-*x*, 1-*y*, 1-*z*] are bridged by two ligands, hereafter noted A and B, to form a six-membered ring {Ag₂N₄}. The Ag(I)...Ag(I) separation is 3.917 Å, which is significantly longer than the sum of the van der Waals radii of two Ag(I) cations (3.44 Å) [58]. The Ag-N separations (2.1812(12) Å and 2.1919(12) Å) are very similar in structures with such coordination bonds [59]. The ligand A is bitopic with the two nitrogen atoms N(1A) and N(2A), however, ligand B interacts with Ag(I) by only one nitrogen atom (N1B). The dihedral angles formed between the aromatic and triazole rings are 4.63° for A and 7.65° for B, much less than the 17.16° observed for the free ligand (**L1**), which indicate that they are nearly coplanar in the silver complex. In [Ag(**L1**)](NO₃), the O-H...N interaction, between two ligands, organizes the dimers into polymeric chains, built on the succession of R₂²(32) rings and developing in the [110] direction, **Fig.6**. O-H...O and C-H...O hydrogen bonds between ligands and nitrate anion organize these chains into 3D supramolecular network. In addition, $\pi \dots \pi$ and Ag... π interactions in conjunction with all hydrogen bonds contribute to the stability of the structure.

In the crystal structure of the second complex, $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$, the coordination geometry of Ag(I) ion is square pyramidal with a basal plane constituted by N(1A) and N(2A) triazole nitrogen atoms in two (**L2**) molecules and by oxygen atoms of two NO_3^- anions. The apical direction of the square pyramid is defined by the O(1A) atom of the phenyl hydroxyl group located at ortho position. Ag(I) ions, being coordinated by the nitrogen atoms of triazole moiety, is found in distorted linear geometry (N(1A)–Ag(1)–N(2A) of $179.31(5)^\circ$). This slight deviation from perfect linear geometry, is essentially due to the weak Ag...O interactions between Ag(I) and nitrate anion on one side and hydroxyl group on the other side ($2.6892(12)$ Å and $2.7181(14)$ Å respectively). In the Cambridge Structural Database (CSD-2017, update Feb 2017), only two structures are reported considering these two weak interactions as Ag–O bond [60]. Ag(I) ions are connected via bridging ligand molecules to form polymeric chains along *b* axis **Fig.7**. The dihedral angle between triazole and phenyl rings in the bitopic ligand (**L2**) in this polymer is the same as found for the free one (19.52°).

Fig.5: ORTEP representation of the dimer of $[\text{Ag}(\text{L1})](\text{NO}_3)$ with thermal displacement ellipsoids drawn at the 50% probability level. (i) $2-x, 1-y, 1-z$

Fig.6: Interligand O-H...N hydrogen bond connecting dimers

Fig.7: Polymeric chains of $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ developing along **b** axis

Interestingly, unlike in the free (**L2**) ligand, in the polymer $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$, one S(6) intra-ligand O-H...N hydrogen bond is observed between the *o*-OH group of the phenyl ring and the azomethine nitrogen atom, **Fig.8**. O-H...O hydroxyl-water hydrogen bonds connect polymeric chains in 2D sheets parallel to (bc) plane C-H...O ligand-nitrate anion and $\pi\cdots\pi$ interactions reinforce the crystal structure and contribute to the construction of 3D supramolecular network. In the structure of $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$, the presence of a structured water molecule is

noteworthy. This water molecule fills a void formed by five (**L2**) molecules and a NO_3^- ion and is involved in three hydrogen bonds: two as donor to oxygen atoms of a NO_3^- ion and of a hydroxyl group of the ligand, and one as acceptor of hydrogen atom of the same hydroxyl group located in a symmetry related molecule (**Fig.8**). This water molecule has probably a significant impact on the packing stability as the hydrogen bonds in which it is involved are among the shortest in the crystal structure, with oxygen-oxygen distances of 2.617(2) and 2.812(2) Å (see Table S3 in supplementary materials).

Fig.8: 2D Sheets built on the basis of O-H...O and O-H...N hydrogen bonds in $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$

Hirshfeld surfaces analyses of (L1), (L2):

The intermolecular interactions within the crystal structures of the ligands (**L1**) and (**L2**), and of their silver complexes were analysed by combining 3D Hirshfeld surface and 2D fingerprint plots. A better understanding of the molecular structure requires quantitative insights into the relative contributions of intermolecular interactions within the crystal [61]. For that purpose, Hirshfeld surfaces of the title compounds have been generated, and mapped over d_{norm} value (**Fig. 9**). The red hot spots on the surface indicate the presence of atoms in very close proximity to the outside of the Hirshfeld surface. The white surface means that the

atoms are in medium proximity, while the blue surface shows little proximity of outside atoms.

Fig. 9: Hirshfeld surfaces of the title compounds mapped with

$$d_{norm} = \frac{d_i - r_i^{vdW}}{r_i^{vdW}} + \frac{d_e - r_e^{vdW}}{r_e^{vdW}}$$

Recently, these type of triazole compounds have been studied by Safin *et al.* [49]. In this paper we have used the Hirshfeld surfaces and 2D fingerprint plots analyses to perform a comparative study of the intermolecular interactions in two different families, the free ligands and their silver complexes. From the combination of the 3D Hirshfeld surface and 2D fingerprint plots of the **(L1)** and **(L2)** compounds, we notice that the intermolecular H...H contacts have the main contribution, corresponding to 31.3% and 29.5% of the total Hirshfeld surfaces areas (**supp.5**). For **(L1)** we observe the presence of short intermolecular H...H contact where $d_e = d_i \approx 1.13$ Å. Two red spots are depicted on the d_{norm} surfaces, **Fig.10**. On the other hand, the intermolecular H...H contacts for **(L2)** are bit longer in the vicinity of 2.65 and 2.86 Å, these contacts are manifested as white spots on d_{norm} which are considered as weak interaction.

Fig. 10: Hirshfeld surfaces mapped with d_{norm} for (L1) and (L2) showing H...H contacts

As shown previously, the stabilization of the crystal structure is provided for the two structures by hydrogen bonds. In the fingerprint plots of both (L1) and (L2), N...H contacts are depicted in **supp.5** (supplementary materials), as a pair of sharp spikes at $d_e + d_i \approx 1.8$ Å. These contacts in the two structures (L1) and (L2) have considerable contribution on Hirshfeld surface of 27.3% and 23.3%, respectively (**Fig. 12**).

The red spots on the d_{norm} surfaces in **Fig.11** are due to the $H_C...N$ and $H_O...N$ contacts corresponding to C-H...N and O-H...N hydrogen bonds, respectively. Thus, in the d_{norm} surface of (L1) and (L2); the red spots of $H_O...N$ appear larger than $H_C...N$, which is related to the hydrogen bonds interaction distances, and hence to their strength.

Fig.11: Hirshfeld surfaces mapped with d_{norm} for (L1) and (L2) showing O-H...N and C-H...N hydrogen bonds

The ligand (L2) contains an extra OH function in the phenolic part of the molecule compared to (L1). As expected, this function increases the proportion of O...H contacts, and decreases the C...H ones, on Hirshfeld surface in (L2). The O...H contacts show up as a sharp spike in the fingerprint plots of the molecules at $d_e + d_i \approx 2.4$ Å. The packing cohesion of the two structures is also provided by C...H interactions, representing 16.4% and 11.3% of the total surface area for (L1) and (L2), respectively, (**supp.5** and **supp.6**). As stated in the previous section, $\pi... \pi$ stacking interactions also contribute to the cohesion of the crystal packing. Such interactions appear on Hirshfeld surfaces as C...C and C...N intermolecular contacts. However, these contacts in the two structures represent a small proportion of the total

Hirshfeld surface areas. The fingerprint plots show the C...N interactions as a region of blue/green colour on the diagonal at $d_e \approx d_i \approx 1.80 \text{ \AA}$ (supp.5 and supp.6).

In the study by Safin *et al.* [49] the same molecule (**L2**) had different fingerprint plots (Fig .12), reflecting differences in the structures due to the way the X-ray data were collected. The fingerprint plot is highly sensitive to the environment surrounding the molecule, and unique for each molecule in a particular polymorphic form [42]. The distinction can be observed in the contribution of each intermolecular contacts and in the sum of d_e , d_i . Furthermore, the area and the volume closed by Hirshfeld surfaces ($223.56 \text{ \AA}^2$, $218.04 \text{ \AA}^3$) for (**L2**) and ($222.81 \text{ \AA}^2$, $216.60 \text{ \AA}^3$) for the same one previously reported by Safin and co-workers, named in this work (**L2bis**), show this distinction [49]. The packing energy (PE) is one of the most important property which can be calculated to characterize the crystal packing cohesion. Using the dispersive-repulsive forces only, the packing energy can be approximated by the empirical formula developed by Gavezzotti and Filippini studying the geometry of intermolecular X-H...Y (X, Y = N,O) hydrogen bond and the calibration of its empirical potentials [62]:

$$PE = \frac{1}{2} \sum \sum E(R_{ij}) \text{ where } E(R_{ij}) = A \exp(-BR_{ij}) - CR_{ij}^{-6}$$

Using this potential as implement in the software « Mercury CSD 2.0 - New Features for the Visualization and Investigation of Crystal Structures » [63], the packing energies of (**L1**) and (**L2**) crystal structures are -153.4 kJ/mol and -191.4 kJ/mol, respectively. The fact that the crystal structure of ligand (**L2**) appears more stable than the one of (**L1**) can be explained by the extra OH function which gives more stability to the crystal packing, and by the presence of the water molecule mediating intermolecular interactions by strong O-H...O hydrogen bonds [64].

Fig.12: 2D Fingerprint plot of (**L2**, this work) and (**L2bis**, Safin *et al.* [49])

Hirshfeld surfaces analyses of [Ag (L1)] (NO₃) and [Ag (L2)]_n(NO₃)_n(H₂O):

The fingerprint plots of the silver complexes are drastically different from the ones obtained from the crystal structures of the free ligands.

The main share of total Hirshfeld surface for the complexes [Ag(L1)](NO₃), [Ag(L2)]_n(NO₃)_n(H₂O) is attributed to O...H interactions with 21.4% and 25.8% of the total surfaces, respectively. O...H interactions appear as two sharp spikes in the fingerprint plots of the two complexes (**supp.5**). The upper spike in bottom left (donor) region of the fingerprint plot where ($d_e = 1.19$ Å, $d_i = 0.83$ Å) for [Ag(L1)](NO₃) and ($d_e = 1$ Å, $d_i = 0.65$ Å) for [Ag(L2)]_n(NO₃)_n(H₂O) involves 17.5% and 19.9% of the total surface, respectively. The lower spike which correspond to the hydrogen-bond acceptor in the region ($d_e = 1.1$ Å, $d_i = 1.4$ Å) for [Ag(L1)](NO₃) and ($d_e = 0.7$ Å, $d_i = 1.1$ Å) for [Ag(L2)]_n(NO₃)_n(H₂O) corresponds to 4% and 5.9% of the total surface, respectively. With a comparable proportion than for O...H interactions, H...H contacts account for the quarter of the total Hirshfeld surfaces area of each complex. These contacts are showed in the middle of scattered points in the fingerprint plots (**supp.5**). Furthermore, in the [Ag(L1)](NO₃) structure N...H contacts play the main role in the stabilization of crystal packing. The N...H contact are observable as a very sharp spike ($d_e = 0.9$ Å, $d_i = 1.19$ Å) (**Supp.5**). Interestingly, these contacts represent 16.9% of the total surface, while only 7.2% for [Ag(L2)]_n(NO₃)_n(H₂O). The π ... π stacking (C...C) contacts is manifested in the 2D fingerprint plot, (**Supp.5**), as a region of blue/green colour ($d_e = d_i \approx 1.65$ Å) corresponding to an interlayer distance of 3.3 Å for [Ag(L1)](NO₃). For [Ag(L2)]_n(NO₃)_n(H₂O) the π ... π stacking are visible in the fingerprint plot where ($d_e = d_i \approx 1.8$ Å) corresponding to an interlayer distance of 3.6 Å. The pattern of red and blue triangles on the Hirshfeld surface coloured by the shape index value is characteristic of π ... π stacking (**Fig.13**).

Fig.13 : Hirshfeld surfaces mapped with shape index (front and back views) for $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$.

The C...H interactions appear as wings in fingerprint plot for both structures $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$. These contacts occupy 12% and 9.2% of the total Hirshfeld surface area, respectively.

The counter-ion NO_3^- contribute in the cohesion of the crystal packing. Its oxygen atoms are involved in C-H...O and O-H...O hydrogen bonds in both structures, (**Fig.14**). The Hirshfeld surface around the counter-ion in $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ shows the interactions between oxygen and Ag(I) cations which represent 15.7% of the total surface (**Fig. 15(c-d)**).

Fig.14: Hirshfeld surfaces mapped with d_{norm} for NO_3^- in the $[\text{Ag}(\text{L1})](\text{NO}_3)$ (a,b) and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ (c,d) crystal structures. The surfaces are shown in two different orientations

In order to acquire a better understanding of the silver contribution in the crystal lattice cohesion, the atomic Hirshfeld surface analysis has been undertaken. The Hirshfeld surface

around Ag(I) mapped over curvedness (**Fig.15**), is introduced to understand the coordination and the chemical interactions in the two complexes $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$.

Fig.15 : Curvedness plotted on the atomic Hirshfeld surface is mapped from -1,4 (flat; red) to -0,3 (sphere-like, blue). (a): $[\text{Ag}(\text{L1})](\text{NO}_3)$ and (b): $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$

The Hirshfeld surface was divided into a set of patches, where each patch represents a different interaction[8]. The shape of the surface indicates the regions of interactions, which varies from high curvature (blue) to low curvature (red), (**Fig.15**). Curvedness plotted on the Hirshfeld surface reflects closer contacts between adjacent atoms [62] and allows also to estimate the coordination number of the metal [65-66]. The flat regions on the Hirshfeld surface around Ag(I) ion in both structures $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$, reveal the number and the nature of the interatomic interaction. The large flat regions are related to the strong interaction [60]. These regions on the surface around Ag(I) cation in the $[\text{Ag}(\text{L1})](\text{NO}_3)$ dimer is related to the covalent bonds Ag-N, Where the metal is tricoordinated by three nitrogens of triazole rings. The remaining small flat regions on the surface refer to the medium and weak interactions Ag... π and Ag...O, respectively. In the $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ polymer structure, the Hirshfeld surface around the Ag(I) cation shows two large flat regions related to the covalent bonds Ag-N, which refer to coordination number 2 (linear structure). Besides, the small flat regions denote medium and weak interactions Ag...O and Ag... π , respectively. The atomic Hirshfeld volumes have been calculated for the Ag(I) cation in both structure to be $27.03 \text{ \AA}^3$ for the dimer and $22.95 \text{ \AA}^3$ for the polymer. Due to the coordination number of the silver in the two structures, the Hirshfeld volume of the Ag(I) cation in the dimer is a bit larger than that one of the same atom in the polymer structure.

Antibacterial studies:

Antibacterial activity of the free ligands (**L1**), (**L2**) and of their silver complexes $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ has been evaluated using the well-diffusion method [67] at 20 mg.ml^{-1} concentration in DMSO against four Gram-negative (*E. coli*, *Salmonella typhimurium*, *Klebsiella pneumoniae* and *Serratia marcescens*) and two Gram-positive (*Staphylococcus aureus*, *Streptococcus*) bacteria strains. The standard antibiotics, Colistin, Fosfomycin, Gentamycin, Erythromycin, Rifampicin, Amikacin, Ofloxacin and Amoxicillin have been used as positive controls against some of the tested bacteria. Solvent control test was also carried out to study the DMSO effect on the bacteria growth. Similarly, in an attempt to dissociate the inhibitory and antimicrobial effects of ionic silver from the one of the silver complexes described in this study, AgNO_3 salt was also used as second control test. The antibacterial activity results of these compounds are listed **table S4** in supplementary materials, and represented in **Fig.16**, as averages of four measurements.

It was observed that the DMSO alone did not inhibit the growth of these bacteria while, as expected, the silver nitrate exhibits significant antibacterial activity. Indeed, even if the precise antimicrobial mechanism of ionic silver is not totally elucidated, several plausible mechanisms have been described in the literature. It is thought that silver ions can kill bacteria by entering the cell through interactions with enzymes present on cell surface, and once inside the cell can bind sulphur-containing groups such as thiol in proteins, leading to the deactivation or the denaturation of enzymes essential for the cell metabolism [68]. Moreover, TEM imaging of bacteria exposed to silver nitrate showed phenomenon of DNA condensation, leading to the loss of its replication ability, along with deep structural damage of cell membranes [34]. Also, the antibacterial activity of silver may be due to its binding to the surface of the cell membrane disrupting the permeability and the respiration [69, 70].

In our experiments, the antibacterial activity of silver nitrate was especially strong against *Staphylococcus aureus*, with a diameter of inhibition zone of $22.0(6.7) \text{ mm}$, significantly larger than the one of *Rifampicin* ($d = 13 \text{ mm}$), a drug used generally as an adjunctive in the treatment of *Staphylococcus aureus* infections [71].

Experiments performed using the free ligands indicated that they do not display any measurable antibacterial activity, excepted against *Serratia Marcescens* for which inhibition zones of $13.8(1.7) \text{ mm}$ and $9.5(1.7) \text{ mm}$ diameters were observed for (**L1**) and (**L2**) ligands, respectively. These clearance zones diameters compare, for (**L1**), with the one of silver nitrate ($d = 15.0(1.1) \text{ mm}$), but is significantly smaller in the case of (**L2**). Nevertheless, this is

especially remarkable that the standard antibiotic Gentamycin did not lead to any measurable inhibition zone in the case of the *Serratia* strain tested in this study. Even if the precise mechanism of action of these ligands on *Serratia Marcescens* is unknown, the fact that they are active against a Gram-negative bacteria indicate that the lipophilicity of **(L1)** and **(L2)** molecules allows them to cross, or at least to interfere with their outer phospholipidic membrane. The outer membrane permeability barrier has indeed a big impact on the susceptibility of the microorganism to antibiotics [72-73].

Antimicrobial activity measurements were also performed to test the metal complexes described in this study. In the case of Gram-positive bacteria, **Fig. 16** shows that the complexes $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ display clearance zones of respectively 19(1) mm /17.3(1.3) mm diameters for *Staphylococcus aureus* and 16.0(2.5) mm /15.0(1.1) mm for *Streptococcus*. These antimicrobial activities are however comparable to the ones obtained, under the same experimental conditions, with silver nitrate alone ($d = 22(7)$ mm for *Staphylococcus* and 15.3(1.3) mm for *Streptococcus*). Hence, given that the parent ligands alone did not display any measurable activity for these bacteria strains, these results indicate that the antimicrobial effect of the $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ complexes cannot reliably be distinguished from the one of ionic silver.

Experiments performed to test the silver complexes against Gram-negative bacteria led to variable outcomes. At first, it appears for *Salmonella typhimurim* that the clearance zones obtained for both complexes, accounting for standard errors, compare with the one of the silver nitrate alone ($d = 16.3(1.3)$ mm, 15.3(1.3) mm and 16.5(8) mm for $[\text{Ag}(\text{L1})](\text{NO}_3)$, $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ and AgNO_3 , respectively, (**Fig. 16**), leading to inconclusive results concerning a possible antimicrobial activity of the silver complexes in this case. For *Serratia Marcescens*, both **(L1)** and **(L2)** ligands displayed moderate antimicrobial activities, comparable or lower than the one of silver nitrate. When these ligands are being tested in their silver complex form, diameters of inhibition zones, again, are comparable to the ones obtained with silver nitrate (**Fig. 16**). Besides, the most significant results were obtained for antimicrobial activities of the complexes against *E. Coli* and *Klebsiella Pneumoniae*. Indeed, for the first bacterium, *E. Coli*, both Ag(I) complexes exhibit more efficient antibacterial activity than AgNO_3 ($d = 11.0(1.1)$ mm), and the polymer $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ ($d = 17.8(1.7)$ mm) has a better activity than the dimer $[\text{Ag}(\text{L1})](\text{NO}_3)$ ($d = 14(0)$ mm). For the second bacterium, *Klebsiella pneumoniae*, only the polymer based on **(L2)** ligand exhibits a significantly better activity ($d = 26.0(2.5)$ mm) than AgNO_3 ($d = 14.5(1.8)$ mm). This result is especially remarkable as the isolated ligand **(L2)** seemed inactive against *Klebsiella*

Pneumoniae, and the antibacterial activity of the corresponding complex compare with the one obtained for the standard antibiotic Gentamycin tested in the experiments ($d = 29$ mm). Even if the elucidation of a precise molecular mechanism of the bactericidal effect of $[\text{Ag}(\text{L}2)]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ against *Klebsiella pneumoniae* is beyond the scope of this study, plausible explanations can nevertheless be proposed. At first, the lipophilicity of the ligand (**L2**), which has been shown able to be moderately active against the Gram-negative bacteria *Serratia Marcescens*, can be enhanced upon complexation with Ag(I) [74], allowing the $[\text{Ag}(\text{L}2)]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ complex to cross the phospholipidic membrane of *Klebsiella pneumoniae*. Once inside the cell, assuming a favourable kinetic of de-ligation reaction, and given that the ligand (**L2**) possesses a triazole group known to be a strong nitrogen donor toward transition metals, the chelation theory may explain the measured activity. Indeed, chelating agents are able to disturb the metal metabolism of microorganisms by interfering with metal acquisition needed for crucial reactions, or by inhibiting the biological role of metal-dependent proteins [75]. In this scenario, the observed activity of $[\text{Ag}(\text{L}2)]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ is thus interpreted as a synergetic effect of the chelating properties of the ligand (**L2**) and the known antimicrobial activity of ionic silver, in conjunction with an enhanced lipophilicity of the ligand upon formation of the silver complex. This may explain why the isolated (**L2**) ligand does not seem to be active and why the silver complex based on (**L2**) appears significantly more active than the silver nitrate control.

Fig.16: Antimicrobial activity of the four title compounds. Error bars correspond to 95% confidence intervals using a student *t*-distribution based on quadruplicate measurements.

Conclusion:

Two hydroxy Schiff bases have been synthesized by condensation of 4-amino-4H-1,2,4-triazole with 3-hydroxybenzaldehyde for the first ligand (**L1**) and 2,4-dihydroxybenzaldehyde for the second one (**L2**). The preliminary IR spectroscopy study of these two ligands has highlighted the formation of imine bond $\text{C}=\text{N}$ characteristic of Schiff base compounds (band around 1600 cm^{-1}). The two compounds differ by the presence of an additional OH in the second Schiff base (**L2**). This difference influences directly the hydrogen bond network involving the two ligands. Infinite chains formed by $\text{O-H}\dots\text{N}$ hydrogen bonds in (**L1**) are indeed transformed into 2D layers in (**L2**). Three dimensional supramolecular assemblies of (**L1**) and (**L2**) in their respective crystal structures is then ensured by $\text{C-H}\dots\text{N}$ hydrogen bonds, and $\pi\dots\pi$ stacking interactions.

The reaction of these two Schiff bases with AgNO_3 leads to the formation of Ag(I) dimer $[\text{Ag}(\text{L1})](\text{NO}_3)$ with (**L1**) and polymer $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ with (**L2**) which includes a water solvent molecule. Ag adopts a trigonal geometry in the dimer and a square pyramidal one in the polymer. This polymer expands on 1D direction and the $\text{O-H}\dots\text{O}$ hydroxyl-water hydrogen bonds connect the polymeric chains in 2D sheets parallel to (bc) plane. $\text{C-H}\dots\text{N}$, $\text{C-H}\dots\text{O}$ and $\pi\dots\pi$ stacking reinforce the crystal structure of the two compounds and ensure the three dimensionality of their networks. Analyse based on Hirshfeld surfaces confirm the presence of these interactions and reveals that $\text{N}\dots\text{H}$ and $\text{O}\dots\text{H}$ interactions evolve inversely in the way (**L1**), (**L2**), $[\text{Ag}(\text{L1})](\text{NO}_3)$ and $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$. The percentage of $\text{O}\dots\text{H}$ interaction increases in this way while the one of $\text{N}\dots\text{H}$ contacts decreases, and it is probably due to the presence of additional OH in the second ligand (**L2**), and of a water molecule in the polymer $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$. Antimicrobial activity of the four compounds confirms also the importance of Ag(I) in bacteria inhibition, indeed, while the free ligands are only active for one bacterium (*Serratia Marcescens*), the silver (I) dimer and polymer inhibit the growth of all bacteria strains tested in this study. However, activities of the complexes are generally comparable to the ones of the silver nitrate tested alone, except in the case of the polymer $[\text{Ag}(\text{L2})]_n(\text{NO}_3)_n(\text{H}_2\text{O})$ which has been shown to be significantly more active than ionic silver against *Klebsiella pneumoniae*.

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Highlights

- Synthesis of new triazole Schiff bases and their silver complexes.
- X-ray crystal structure and Hirshfield surfaces.
- Spectroscopy properties of prepared compounds.
- Antimicrobial activity •

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