



## Article

# Synthesis, Identification and Antibacterial Activities of Amino Acid Schiff Base Cu(II) Complexes with Chlorinated Aromatic Moieties

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**Abstract:** Amino acid Schiff base Cu(II) complexes were synthesized under microwave irradiation using methanol as a solvent, to maximize the best conditions to obtain the attained compounds, containing aromatics possessing no, one or two chlorine atoms. The compounds' antibacterial activities were tested against Gram-positive and Gram-negative bacteria, and the most active were tested for their antioxidant activities, and as *E. coli*, in particular, was found to be sensitive to these compounds, their interaction with this bacterium was investigated. It was found that, depending on the amino acid used for the formation of the Schiff base ligand, its LogP<sub>o/w</sub> mono-chlorinated or bis-chlorinated compounds are the most efficient against the tested bacteria.



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**Keywords:** amino acid Schiff base Cu(II) complexes; chlorinated salicylaldehyde; antibacterial activity; interaction with *E. coli*

## 1. Introduction

Schiff bases were discovered by Hugo Schiff, and their metal complex derivatives were largely synthesized, for example, incorporating amino acids [1]. Their fields of application are varied and concern, for example, environmental sensors [2], catalysis [3,4], and anticancer [5,6] or antioxidant agents, but they also serve as antimicrobial agents, particularly against bacterial and fungal pathogens, showing that, compared to Co(II), Ni(II) or Zn(II), Cu(II) complexes show lower MIC values, revealing that these complexes present a better growth inhibitory activity [7]. Other Schiff bases' copper complexes were described from salicylaldehyde derivatives and tested as antimicrobials [8,9], but in all the previous literature on amino acid copper-complex Schiff base syntheses [10], it is noticeable that the reaction time is somehow long, lasting from several hours to days.

Since the use of microwave heating to accelerate organic chemical transformations was first reported by the group of Gedye and Giguere/Majetich in 1986 [11,12], many studies using microwaves have been published [13] about the wavelengths, frequencies and applications of various radio waves used in our daily lives. Controlled microwave heating under closed vessel conditions greatly accelerates reactions compared to conventional synthesis methods using external heating, resulting in higher yields of products, fewer unwanted side reactions and the ability to synthesize high-purity compounds with fewer raw materials [13,14]. Microwave-assisted synthesis typically utilizes "microwave dielectric heating" phenomena, such as dipolar polarization and ionic conduction mechanisms, and relies on the ability of the reacting mixture to efficiently absorb microwave energy [15]. The ability of a specific solvent to convert microwave energy into heat at a given frequency and temperature is determined by the so-called loss tangent ( $\tan \delta$ ), expressed as the quotient,  $\tan \delta = \epsilon'' / \epsilon'$ , where  $\epsilon''$  is the dielectric loss, indicative of the efficiency with

which electromagnetic radiation is converted into heat, and  $\epsilon'$  is the dielectric constant, describing the ability of molecules to be polarized by the electric field. A reaction medium with a high  $\tan \delta$  at the standard operating frequency of a microwave synthesis reactor (2.45 GHz) is required for good absorption and, consequently, efficient heating [15]. In general, solvents used for microwave synthesis are classified as having high ( $\tan \delta > 0.5$ ), medium ( $\tan \delta 0.1\text{--}0.5$ ) or low ( $\tan \delta < 0.1$ ) microwave absorption. Microwave synthesis in low-absorbing or microwave-permeable solvents are often not feasible. Among the solvents possessing a high  $\tan \delta$ , methanol is one of the most suitable solvents for microwave synthesis because of its high microwave absorption with a loss  $\tan \delta$  of 0.659 [16]. That is why this method was previously employed for the synthesis of copper Schiff base complexes by microwave from a 2-step procedure, lowering the global reaction time to 10 min [17], leading, for example, to di-chlorinated compounds and, thanks to the synthetic optimized method, new mono-chlorinated Schiff base copper complexes were synthesized in this work, and all these compounds were tested against model bacteria to investigate how they inhibit the growth of these bacteria.

## 2. Materials and Methods

### 2.1. Compound Syntheses

Known compounds C1.1–C8.1 were prepared in the previous paper and used as described in [17]. New compounds entries C1.2–C12.2 were prepared and characterized as follows.

$\text{Cu}(\text{CH}_3\text{COO})_2 \cdot \text{H}_2\text{O}$ , *L*-alanine, *L*-leucine, *L*-serine and *L*-threonine used in this study were purchased from Wako Fujifilm (Osaka, Japan), and salicylaldehyde, 3-chlorosalicylaldehyde, 4-chlorosalicylaldehyde and 5-chlorosalicylaldehyde were purchased from TCI (Tokyo, Japan). The synthesis of the compounds was conducted at 358 K using an Initiator+ microwave apparatus (Biotage, Tokyo, Japan).

#### 2.1.1. General Procedures

To a methanol solution (20 mL) of *L*-amino acid (0.2 mmol) was added the proper chlorosalicylaldehyde (0.2 mmol), and the mixture was stirred under microwave irradiation at 358 K for 10 min. Then, a methanol solution (20 mL) of  $\text{Cu}(\text{CH}_3\text{COO})_2 \cdot \text{H}_2\text{O}$  (0.0399 g, 0.2 mmol) was added and the mixture was stirred under microwave irradiation for 10 min at 358 K to produce the final product. For additional spectra, see Supplementary Materials.

#### 2.1.2. Physical Measurements

Elemental analyses were conducted with a Perkin-Elmer 2400II CHNS/O analyzer, (Waltham, MA, USA) at the Tokyo University of Science. Infrared (IR) spectra were recorded on a JASCO FT-IR 4200 spectrophotometer (JASCO, Tokyo, Japan) in the range of  $4000\text{--}400\text{ cm}^{-1}$  at 298 K. Absorption electronic (UV-Vis) spectra were measured on a JASCO V-570 spectrophotometer in the range of  $800\text{--}250\text{ nm}$  at 298 K.

#### 2.1.3. Preparation of C1.2

Using the general procedure with *L*-leucine (0.02623 g, 0.2 mmol) and 3-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.06985 g, 54.73%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether and ethanol, and dried in a desiccator for several days. This product was then filtered. Anal. Calcd. for  $\text{C}_{13}\text{H}_{16}\text{ClCuNO}_4$ : C, 44.71; H, 4.62; N, 4.01%; found: C, 45.69; H, 3.5; N, 3.84%.

#### 2.1.4. Preparation of C2.2

Using the general procedure with *L*-leucine (0.02623 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.06985 g, 32.85%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $\text{C}_{13}\text{H}_{16}\text{ClCuNO}_4$ : C, 44.71; H, 4.62; N, 4.01%; found: C, 43.39; H, 2.51; N, 1.20%.

#### 2.1.5. Preparation of C3.2

Using the general procedure with *L*-leucine (0.02623 g, 0.2 mmol) and 5-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.03918 g, 56.09%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{13}H_{16}ClCuNO_4$ : C, 44.71; H, 4.62; N, 4.01%; found: C, 45.1; H, 3.71; N, 3.82%.

#### 2.1.6. Preparation of C4.2

Using the general procedure with *L*-alanine (0.01782 g, 0.2 mmol) and 3-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.40000 g, 65.11%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_4$ : C, 39.10; H, 3.28; N, 4.56%; found: C, 40.78; H, 2.45; N, 4.43%.

#### 2.1.7. Preparation of C5.2

Using the general procedure with *L*-alanine (0.01782 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.04146 g, 67.48%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_4$ : C, 39.10; H, 3.28; N, 4.56%; found: C, 41.39; H, 2.32; N, 4.54%.

#### 2.1.8. Preparation of C6.2

To a methanol solution (20 mL) of alanine (0.01782 g, 0.2 mmol) was added 4-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) and stirred at 358 K for 10 min to obtain a pale yellow color. Then, a methanol solution (20 mL) of  $Cu(CH_3COO)_2 \cdot H_2O$  (0.03990 g, 0.2 mmol) was added, and the mixture was stirred for 10 min at 358 K to produce a green compound (yield 0.01060 g, 17.25%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_4$ : C, 39.10; H, 3.28; N, 4.56%; found: C, 36.6; H, 3.13; N, 4.08%.

#### 2.1.9. Preparation of C7.2

To a methanol solution (20 mL) of serine (0.02102 g, 0.2 mmol) was added 3-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) and stirred at 358 K for 10 min to obtain a pale yellow color. Then, a methanol solution (20 mL) of  $Cu(CH_3COO)_2 \cdot H_2O$  (0.0399 g, 0.2 mmol) was added, and the mixture was stirred for 10 min at 358 K to produce a green compound (yield 0.03737 g, 58.04%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_5$ : C, 37.16; H, 3.12; N, 4.33%; found: C, 33.17; H, 3.27; N, 7.17%.

#### 2.1.10. Preparation of C8.2

Using the general procedure with *L*-serine (0.02102 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.04404 g, 68.40%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_5$ : C, 37.16; H, 3.12; N, 4.33%; found: C, 35.31; H, 2.88; N, 4.95%.

#### 2.1.11. Preparation of C9.2

Using the general procedure with *L*-serine (0.02102 g, 0.2 mmol) and 5-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.04678 g, 72.65%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for  $C_{10}H_{10}ClCuNO_5$ : C, 37.16; H, 3.12; N, 4.33%; found: C, 34.34; H, 2.42; N, 3.17%.

#### 2.1.12. Preparation of C10.2

Using the general procedure with *L*-threonine (0.02380 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03131 g, 0.2 mmol) produced a green compound (yield 0.03446 g, 51.09%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for C<sub>11</sub>H<sub>12</sub>ClCuNO<sub>5</sub>: C, 39.18; H, 3.59; N, 4.15%; found: C, 34.34; H, 3.28; N, 6.37%.

#### 2.1.13. Preparation of C11.2

Using the general procedure with *L*-threonine (0.02382 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03135 g, 0.2 mmol) produced a green compound (yield 0.02844 g, 42.17%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for C<sub>11</sub>H<sub>12</sub>ClCuNO<sub>5</sub>: C, 39.18; H, 3.59; N, 4.15%; found: C, 44.48; H, 1.62; N, 6.71%.

#### 2.1.14. Preparation of C12.2

Using the general procedure, *L*-threonine (0.02382 g, 0.2 mmol) and 4-chlorosalicylaldehyde (0.03135 g, 0.2 mmol) produced a green compound (yield 0.13912 g, 20.63%). The resulting crude compound was filtered and the precipitate was washed with diethyl ether, acetone and ethanol, then dried in a desiccator for several days. This product was filtered. Anal. Calcd. for C<sub>11</sub>H<sub>12</sub>ClCuNO<sub>5</sub>: C, 39.18; H, 3.59; N, 4.15%; found: C, 44.62; H, 1.53; N, 14.03%.

### 2.2. Antimicrobial Assays

The bacterial strains used were *E. coli* ATCC 25922TM (American Type Culture Collection, Manassas, VA, USA), *M. luteus* CRBIP 107660 (Institut Pasteur, Lille, France), *S. saprophyticus* (isolated by TIMR Laboratory) and *B. subtilis* ATCC 6051 (American Type Culture Collection, Manassas, VA, USA). They were grown on trypto-casein soy solid medium (TSA, Conda, Madrid, Spain) and incubated at 30 °C in the dark (SANYO, incubator, MIR-253) for 24 h. The solid cultures were then stored in a cold room at 4 °C or used to inoculate 5 mL of liquid mineral medium 24 h before the microplate test (glucose 10 g/L, KCl 0.250 g/L, NaH<sub>2</sub>PO<sub>4</sub> 6.464 g/L, Na<sub>2</sub>HPO<sub>4</sub>·2H<sub>2</sub>O 10.408 g/L, MgSO<sub>4</sub> 0.244g/L, NO<sub>3</sub>NH<sub>4</sub> 1 g/L, MgCl<sub>2</sub> 0.05 g/L; pH = 7). A suspension of bacterial inoculum with an optical density (OD) between 0.1 and 0.2 (Ultrospec10 Amersham Bioscience) was prepared from a pre-culture for 24 h. The tests were performed on 96-well microplates (Thermo Scientific, Nunc™ Edge) in triplicate. The tested products were dissolved in DMSO as 100-times-concentrated stock solutions. Blanks were composed of non-inoculated mineral medium with or without the tested compounds to withdraw their eventual variation of absorbance along with time; positive controls were conducted with inoculated mineral medium without the compound and tests were inoculated media with the tested compounds. Growth monitoring was performed with a spectrophotometer Thermo Fisher Scientific Multiskan GO (type 1510). It was set to perform OD measurements at 600 nm of each well every 15 min for 24 h to follow the growth kinetics.

The percentage of inhibition was calculated as follows:

$$\%Inhibition = \left( 1 - \frac{ODvar\ growth}{ODvar\ control} \right) * 100 \quad (1)$$

*ODvar growth* was the difference in OD between the highest and lowest points on the growth curve in the tests. *ODvar control* was the difference in OD between the highest and lowest points on the positive control curve.

The MIC<sub>95</sub> value was considered to be the lowest tested concentration value at which at least 95% inhibition was achieved.

### 2.3. Antioxidant Assays

The tests were performed on 96-well microplates (Thermo Scientific, Nunc™ Edge) in triplicate. The tested products were dissolved in DMSO. Oxidation monitoring was performed with a spectrophotometer Thermo Fisher Scientific Multiskan GO (Type 1510). It was set to perform OD measurements at 450 nm of each well every 1 min for 6 h to follow the oxidation kinetics (2  $\mu$ L of sodium persulfate at 5 mg/mL; 2  $\mu$ L of the tested compound of 50  $\mu$ L ABTS at 0.08%) in a citrate-phosphate buffer (0.1 M) for a total volume of 200  $\mu$ L. The maximum oxidation curve for the positive control reached the absorbance of 0.7. The percentage of inhibition was calculated as follows:

$$\%Inhibition = (1 - (ODvar\ test)/(ODvar\ control)) * 100 \quad (2)$$

*ODvar test* was the difference in OD between the highest and the lowest points on the test curve. *ODvar control* was the difference in OD between the highest and lowest points on the control curve.

### 2.4. Interaction with *E. coli*

The tested compounds in DMSO (200  $\mu$ g/mL as the final concentration) were mixed in culture media with *E. coli*. Their UV-Vis spectra were recorded after filtration on a 0.2  $\mu$ m PTFE syringe filter at different times of incubation at 30 °C for up to 24 h. The values were recorded depending on the complex  $\lambda_{max}$  values (384 nm for C4.1, 372 nm for C6.1, 369 nm for C2.2, 357 nm for C8.2, 366 nm for C9.2 and 378 nm for C11.2) and compared to the solutions without culture media.

### 2.5. Log $P_{o/w}$ Calculation

The theoretical Log  $P_{o/w}$  were calculated thanks to SwissADME from the Swiss Institute of Bioinformatics under the CC-BY 4.0 Creative Commons 4.0 International License.

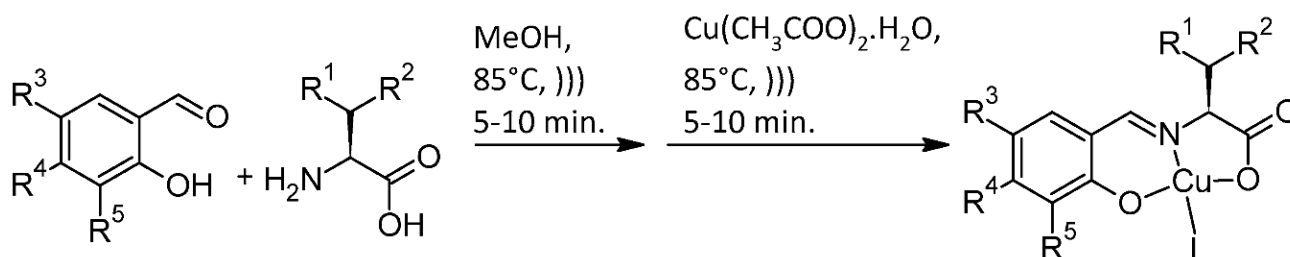
### 2.6. Statistical Analysis

Data are expressed as means  $\pm$  standard deviation, and the statistical significance ( $p < 0.05$ ) was determined by one-way ANOVA with Tukey's post hoc analysis.

## 3. Results and Discussions

### 3.1. Chemistry

For the preparation of *L*-amino acid derivative Schiff base Cu(II) complexes, two-step reactions were employed, namely, (1) imine condensation of primary amine (*L*-amino acid) and aldehyde, and (2) coordination of the Cu(II) ion from an acetate source (Scheme 1). In previous work, microwave syntheses, conventional heating and mechanochemistry for the corresponding *L*-amino acid derivative Schiff base Cu (II) complexes (series 1) [17] were compared, and this compound library was enriched with various mono-chloride compounds (series 2) to better understand the structure–activity link between the amino acid functional group/number and place of the chlorine moiety, as well as the antibacterial activity.



**Scheme 1.** Typical reaction scheme in a solution for the *L*-amino acid derivative Schiff base Cu(II) complexes.

From the previous work (series 1), it is noteworthy that a two-step 5 + 5 (min) microwave synthesis at 85 °C is really efficient for the reaction between valine or threonine and salicylaldehyde (entries 1 and 7, 86% yield), while dichloro salicylaldehyde with threonine or alanine produced lower results (entries 6 and 8, 75% and 78% yield). The new series was then synthesized thanks to the screening antimicrobial results (see 3.2 Antibacterial Effect), and *L*-leucine (non-polar amino acid, entries 9–11) showed that the chlorine position in R<sup>3</sup> or R<sup>5</sup> produced a yield of 55% and 56%; whereas, with R<sup>4</sup> = Cl, only a 33% yield was obtained. With *L*-alanine (non-polar amino acid, entries 12–14), the best result was obtained for R<sup>4</sup> = Cl (67% yield); whereas, for *L*-threonine (non-charged polar amino acid, entries 18–20), the best result was obtained with R<sup>5</sup> = Cl (51% yield, respectively) lowered to R<sup>3</sup> = Cl or both amino acids (17% and 21% yields, respectively). As for the *L*-serine derivatives (non-charged polar amino acid entries 15–17), the best result was obtained with R<sup>3</sup> = Cl (73% yield) and the lower yield was obtained for R<sup>5</sup> = Cl (58%). All these results show that there is clearly an impact from the chlorine position and the amino acid functional group on the yield obtained, not really correlated to their polarity or their charge.

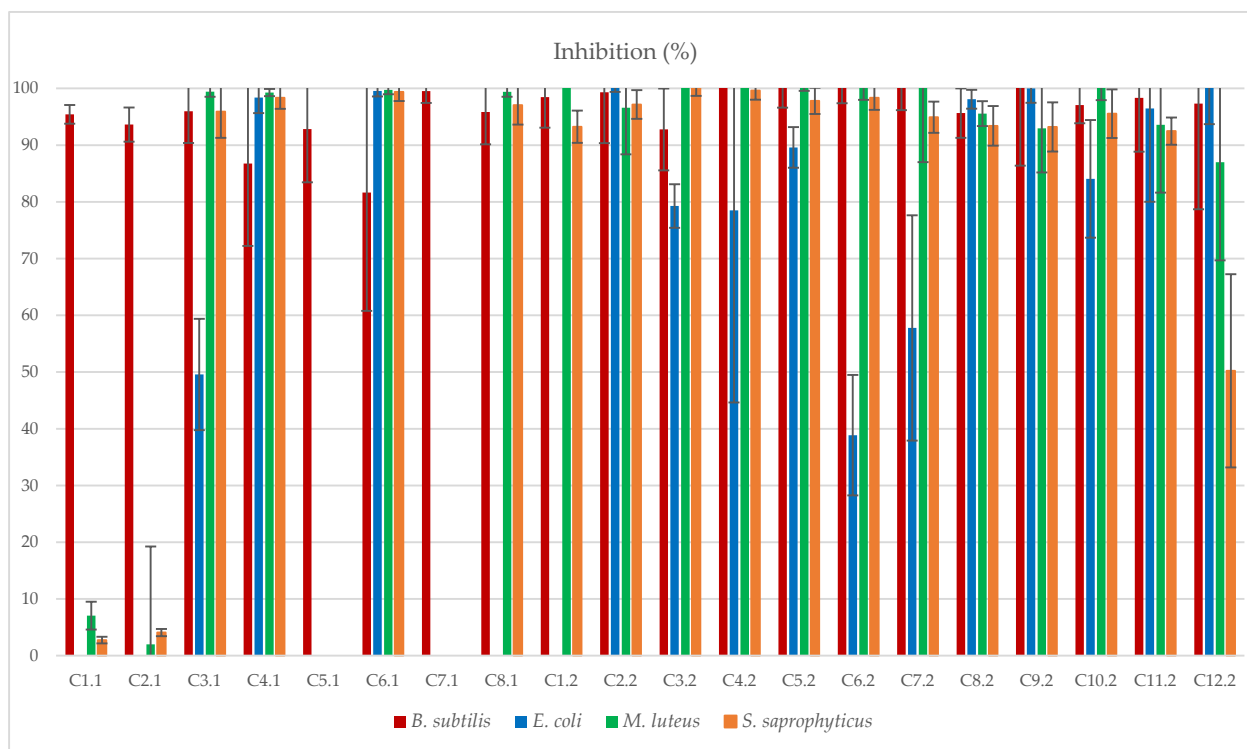
### 3.2. Antibacterial Effect

The synthesized 20 compounds were tested at a concentration of 50 µg/mL against four bacteria *Bacillus subtilis* (Gram (+), rod-shaped), *Staphylococcus saprophyticus* (Gram (+), coccus), *Micrococcus luteus* (Gram (+), coccus) and *Escherichia coli* (Gram (–), rod-shaped). These bacterial strains were chosen in order to have a good representativity of bacterial morphologies, cell grouping and cell wall structures, which are important factors influencing resistance to antibacterial molecules. Moreover, the species studied are described as human pathogens regarding *E. coli* [18] and *S. saprophyticus* [19], or in food poisoning with *B. subtilis* [20] and *M. luteus* [21]. The first group (Table 1, Cn.1, entries 1–8) was chosen for its various functions led by the amino acids (R<sup>1</sup> and R<sup>2</sup>), as well as the presence of 2 or no chlorides (R<sup>3</sup> and R<sup>5</sup>). Then, the second group (Table 1, Cn.2, entries 9–20) was chosen according to the best antibacterial results obtained with group 1, and the mono-chlorinated positions varied (R<sup>3</sup>, R<sup>4</sup> and R<sup>5</sup>). It was clear from the antibacterial tests of the first group that compounds C3.1, C4.1 and C6.1 (entries 3, 4 and 6) had the best antibacterial effect (Figure 1). Indeed, C3.1 was active against all the Gram (+) bacteria, while C4.1 and C6.1 were active against both Gram (+) and Gram (–) bacteria. The last compound, C8.1, was found to be active against all the Gram (+) bacteria, but totally inactive against *E. coli*. The common factor of these three compounds compared to the others (C1.1, C2.1, C5.1 and C7.1) was the presence of the bis-chlorine atoms. Moreover, focusing on the Gram (–) bacteria, C4.1 and C6.1, which were active against this strain, were exempted from oxygen moiety afforded by the amino acid part, whereas both C3.1 and C8.1 possessed a hydroxyl group. Lastly, between the two hydroxylated compounds, the methyl moiety seemed to have a deleterious effect on the activity compared to the H atom. Based on these first excellent results by C3.1, C4.1, C6.1 and C8.1, the second group (Cn.2) was synthesized, but with moderating the place of the mono-chlorine atom (R<sup>3</sup>, R<sup>4</sup> and R<sup>5</sup>) and then tested against the four same bacteria. First, it was really interesting to observe that C6.1 mono-chlorinated isosteres (C4.2, C5.2, C6.2) were less interesting than the bis-chlorinated compound. Then, comparing C1.2, C2.2 and C3.2 led to the observation that the most active compounds against all the strains possessed chlorine in the R<sup>4</sup> position. In a similar manner, comparing C7.2, C8.2 and C9.2 found that C8.2 and C9.2 were the most active compounds and, most importantly, even better than their mono-chlorinated isostere C3.1. As expected, C10.2, C11.2 and C12.2 compounds were less active as their isostere C8.1, even though the mono-chlorinated compound C11.2 was the best of them. According to this antibacterial screening result, MIC (minimum inhibition concentration) values were tested and produced the following results.

**Table 1.** Summary of the results for the obtained 2-step compounds.

Entry	Series	Compound	R <sup>1</sup>	R <sup>2</sup>	R <sup>3</sup>	R <sup>4</sup>	R <sup>5</sup>	Yield <sup>a</sup>
1	1	C1.1	CH <sub>3</sub>	CH <sub>3</sub>	H	H	H	86% <sup>b</sup>
2		C2.1	OH	H	H	H	H	65% <sup>b</sup>
3		C3.1	OH	H	Cl	H	Cl	49% <sup>b</sup>
4		C4.1	<sup>i</sup> Pr	H	Cl	H	Cl	31% <sup>b</sup>
5		C5.1	H	H	H	H	H	67% <sup>b</sup>
6		C6.1	H	H	Cl	H	Cl	75% <sup>b</sup>
7		C7.1	OH	CH <sub>3</sub>	H	H	H	86% <sup>b</sup>
8		C8.1	OH	CH <sub>3</sub>	Cl	H	Cl	78% <sup>b</sup>
9	2	C1.2	<sup>i</sup> Pr	H	H	H	Cl	55%
10		C2.2	<sup>i</sup> Pr	H	H	Cl	H	33%
11		C3.2	<sup>i</sup> Pr	H	Cl	H	H	56%
12		C4.2	H	H	H	H	Cl	65%
13		C5.2	H	H	H	Cl	H	67%
14		C6.2	H	H	Cl	H	H	17%
15		C7.2	OH	H	H	H	Cl	58%
16		C8.2	OH	H	H	Cl	H	68%
17		C9.2	OH	H	Cl	H	H	73%
18		C10.2	OH	CH <sub>3</sub>	H	H	Cl	51%
19		C11.2	OH	CH <sub>3</sub>	H	Cl	H	42%
20		C12.2	OH	CH <sub>3</sub>	Cl	H	H	21%

<sup>a</sup> Isolated yields; <sup>b</sup> from previous work [17].



**Figure 1.** Inhibition rate of Cn.1 and Cn.2 against *B. subtilis*, *E. coli*, *M. luteus* and *S. saprophyticus*.



### 3.3. MIC<sub>95</sub> and MIC<sub>50</sub> Values

MIC<sub>95</sub> values are the minimal concentration where the inhibition reaches at least 95% and MIC<sub>50</sub> values are minimal concentration where the inhibition reaches at least 50%. The results for the tested concentrations are 50, 25, 12.5, 6.25, 3.12, 1.56 and 0.78 µg/mL. From Table 2, it can be observed that, for the bacteria *S. saprophyticus*, the bis-chlorinated molecules exhibited the lowest MIC value. This was totally inverted as for one of the other Gram (+) strains, *B. subtilis*, since the mono-chlorinated compounds possessed the lowest MIC values, maybe due to the difference of cell morphology between the two bacteria. Surprisingly, for *E. coli*, which is a Gram (–) bacteria, the MIC values were the lowest, whatever the molecules, with the best values for C6.1, C2.2 and C11.2. Indeed Gram (–) bacteria are generally described as more resistant against antibacterial molecules due to their more selective cell wall and other more specific mechanisms of resistance [22]. By comparison, between all the tested molecules, on *S. saprophyticus*, *E. coli* and *M. luteus*, the compounds C8.2 and C9.2 exhibited the lowest activity. Moreover, C2.2 and C11.2 were more active against the rod-shaped molecules than the coccus.

**Table 2.** MIC<sub>95</sub>/MIC<sub>50</sub> values (µg/mL).

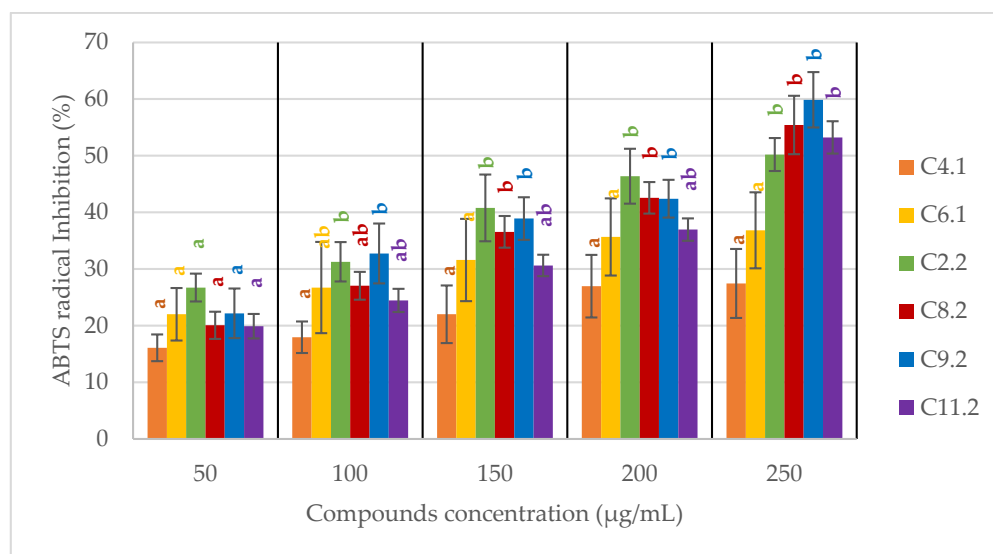
Compounds	<i>S. saprophyticus</i>	<i>E. coli</i>	<i>M. luteus</i>	<i>B. subtilis</i>
C4.1	25/25	25/25	50/12.5	>50/50
C6.1	12.5/12.5	12.5/12.5	25/6.25	>50/50
C2.2	50/25	6.25/6.25	50/12.5	6.25/3.12
C8.2	50/50	50/50	>50/12.5	25/12.5
C9.2	50/50	50/50	>50/50	25/12.5
C11.2	50/50	12.5/12.5	>50/50	12.5/12.5
Streptomycin	25/6.25	-	-	-
Daptomycin	-	-	7.75/2	30/15
Polymyxin B	-	3.75/1	-	-

### 3.4. Antioxidant Activity

ROS (reactive oxygen species) were suspected to be the cause of the bacterial inhibition, so oxidation tests were performed with ABTS (2,2'-azinobis-3-ethylbenzothiazoline-6-sulfonic acid) and copper compounds (C4.1, C6.1, C2.2, C8.2, C9.2 and C11.2), and compared to sodium persulfate, in the dark or under a white lamp (60 watt). The result produced absolutely no oxidized ABTS species, And, as mentioned in the literature, copper Schiff-based compounds can exhibit antioxidant activity [23–25]. The antioxidant activity of the best antibacterial compounds C4.1, C6.1, C2.2, C8.2, C9.2 and C11.2 was kinetically tested against the oxidation of ABTS to the ABTS<sup>+</sup> radical [26] by sodium persulfate (Figure 2).

Two groups of antiradical compounds were identified, as it was clear that the first group (C4.1 and C6.1) did not reach a 50% inhibition with a plateau of antioxidant capability from 200–250 µg/mL and over. However, the second group of compounds (C2.2, C8.2, C9.2 and C11.2) was able to reach 50% inhibition at a 250 µg/mL concentration and in the same statistical group. The conclusion may be that the mono-chlorinated compounds of these series of Schiff bases were more active against the radical oxidation of ABTS than the bis-chlorinated ones. IC<sub>50</sub> was thus calculated from this second group of compounds and compared to the standard *L*-ascorbic acid (AA) (Table 3).





**Figure 2.** ABTS radical inhibition by C4.1, C6.1, C8.2, C9.2 and C11.2 compounds. Bars with different letters indicate statistically significant differences among the groups.

**Table 3.** IC<sub>50</sub> (mM) compared to standard L-ascorbic acid.

	C2.2	C8.2	C9.2	C11.2	AA
IC <sub>50</sub> (mM)	0.72	0.77	0.77	0.74	0.14

### 3.5. Interaction of Copper Complexes with *E. coli*

Copper ions and complexes are known to be biologically active due to the redox processes involved [27]. However, if *E. coli* is equipped with multiple systems to ensure safe copper handling under varying environmental conditions [28,29], in our case, this bacteria was the most sensitive to the copper Schiff base compounds. Therefore, the method of Joseph et al. [30] was adapted to evaluate the capability of the most active compounds to interact with *E. coli*. The absorption band of the complex in the culture media at their respective  $\lambda_{\max}$  wavelengths lowered after interacting with *E. coli* (Table 4).

**Table 4.** Absorbance reduction (%) of copper Schiff base compounds in solution after *E. coli* incubation.

	C4.1	C6.1	C2.2	C8.2	C9.2	C11.2
Absorbance reduction (%)	60.3	34.3	34.4	11.9	2.3	82.3

This absorbance reduction after 24 h of incubation at 30 °C can be correlated to the compound activity. Indeed, the lowest absorbance reductions (2% and 12%) were observed for C9.2 and C8.2, respectively, and for these two compounds, the MIC<sub>95</sub> values of 50 µg/mL were the highest. As for the other compounds, good absorbance reduction greater than 34% led to lower MIC<sub>95</sub> from 6.25 to 25 µg/mL. In this group of active compounds, it is important to notice that, even if C2.2 was less absorbed than C11.2 or C4.1 by *E. coli*, its better activity might be attributed to its functional groups and chlorine placement. Additionally, the theoretical LogP<sub>o/w</sub> is interesting while comparing the uncomplexed ligands. Indeed, while this Log P<sub>o/w</sub> for L4.1 led to C4.1 when complexed copper was high (3.31), the values for L8.2 and L9.2 were low (1.07 and 1.06, respectively). This could also explain why if C4.1 is highly absorbed by *E. coli*, its activity is not as good as the activities of C2.2, C6.1 and C11.2 (Log P<sub>o/w</sub> for L2.2 = 2.69, for L6.1 = 2.37 and for L11.2 = 1.39).

#### 4. Conclusions

Amino acid Schiff base Cu(II) complexes were synthesized under microwave irradiation using methanol as a solvent, to maximize the best conditions to obtain the obtained compounds. First, it was found that some compounds possessing bis-chlorinated moieties had better antibacterial activity compared to non-chlorinated aromatic. The place and number of chlorine atoms on the salicylaldehyde was then investigated and, depending on the amino acid used for the formation of the Schiff base ligand, its  $\text{LogP}_{o/w}$  mono-chlorinated or bis-chlorinated compounds were the most efficient against the tested bacteria. Finally, the light antioxidant effect and the sensitivity of *E. coli* towards the most active compounds led us to test their interaction capability to be absorbed by *E. coli*. Subsequently, we plan to verify the toxicity of the molecules to guide the application, depending on the results obtained for skin, kidney, liver cells.

**Supplementary Materials:** The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/applmicrobiol2020032/s1>: Figures S1–S6: IR spectra, Figures S7–S12: UV-Vis spectra, Figure S13: Tentative crystal structures; Table S1: Compounds.

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