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Antimicrobial Activity and Molecular Modeling Study of Schiff Base Derived from Sulfamerazine

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Abstract

Condensation of 4-amino-*N*-(4-methylpyrimidin-2-yl)benzene sulfonamide (Sulphamerazine Drug) with 1,3-Benzodioxole-5-carbaldehyde (Piperonal) yielded Schiff base derivative in good yield. Spectroscopic characterization such as, IR and NMR of synthesized compound have been obtained by using nuclear magnetic resonance 600 MHz. The synthesized compound was screened for antibacterial activity against *Staphylococcus aureus*, *Escherichia coli*, *Bacillus cereus* and *klebsella pneumonia* and fungicidal activity against *Candida albicans*, *Candida trobicalis*, *Candida krusi*, *Aspergillus fumigatus* and *Aspergillus niger*. A new compound exhibited potent antibacterial and antifungal activity. Molecular modeling studies were performed, showing the hydrogen bindings and hydrophobic interactions.

Keywords: Sulphamerazine, Piperonal, Schiff-base, Microbial activity, Molecular modeling.

Introduction

New sources of antimicrobial agents is need to be discovered due to the presence and constant development of resistant micro-organisms, the emergence of new infections diseases and the toxicity concerns of some of the currently used antimicrobial treatments¹.

The random use of traditional antibiotics and synthetic antimicrobial drugs has resulted in the emergence of resistance microbes such as Methicillin Resistance *Staphylococcus aureus* (MRSA). Vancomycin resistance enterococci and multidrug resistance strain of *Klebsiella pneumonia* and *Pseudomonas aeruginosa* which have come to be a cause of concern for scientists to find a different approach for treating such infection. In a high percentage of asian countries 90-95% of *S. aureus* are penicillin resistance and 75% are methicillin resistance^{2,3}. Sulfa drugs, developed in the 1930s, were the first medications effective against bacterial disease and infection. They seemed as the first “miracle drugs” at a time when death from bacterial infections such as pneumonia and blood poisoning were unrestricted⁴. Moreover, sulfonamide derivatives have received considerable interest from researchers, and many sulfonamide derivatives have been prepared for

their importance in the pharmacological and biological fields. Several Schiff-base compounds derived from sulfonamide have been prepared for use in many biological applications^{5,6}.

Schiff bases are versatile C=N (Imine) containing compounds possessing broad spectrum of biological activity. Schiff base are the compound containing azomethine group (-HC=N-). They are condensation products of ketones (or) aldehydes (aldehyde and ketones) with primary amines and were first reported by Hugo Schiff in 1864⁷. R. C. Maurya *et al* were prepared some Schiff base derived from salicylaldehyde and the sulfa drug sulfamerazine, [N-(salicylidene)-sulfamerazine]⁸.

Schiff base derived from o-vanillin and Sulfamerazine, β – lactam and Zn(II), Cu(I) complexes have been Synthesized and showed highly antimicrobial activity toward mentioned some bacteria⁹.

The aim of present work is to synthesis of new Schiff base derived from sulfamerazine drug and piperonal (Scheme1) and study of their antimicrobial activity and molecular modeling.

Fig. 1: ^1H NMR of synthesized compound

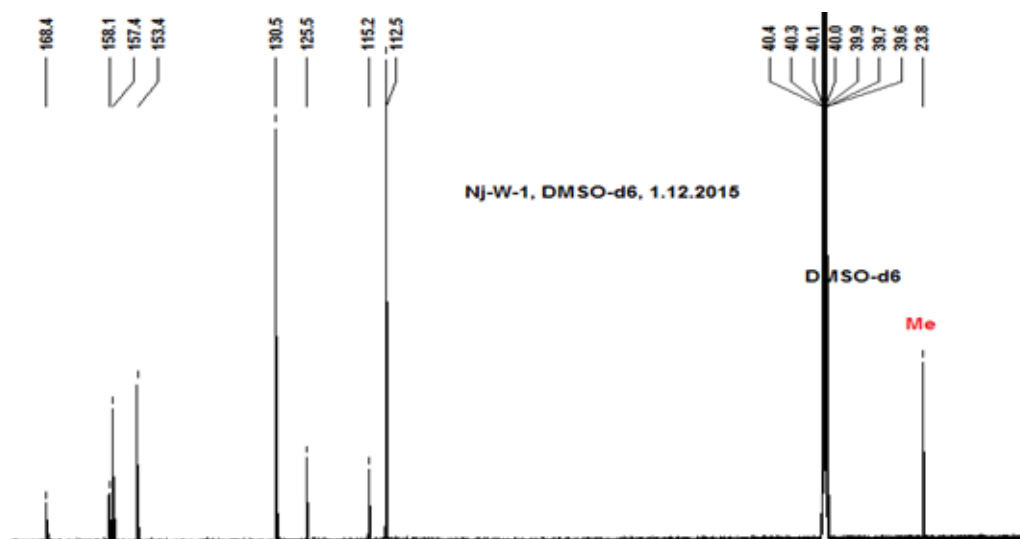
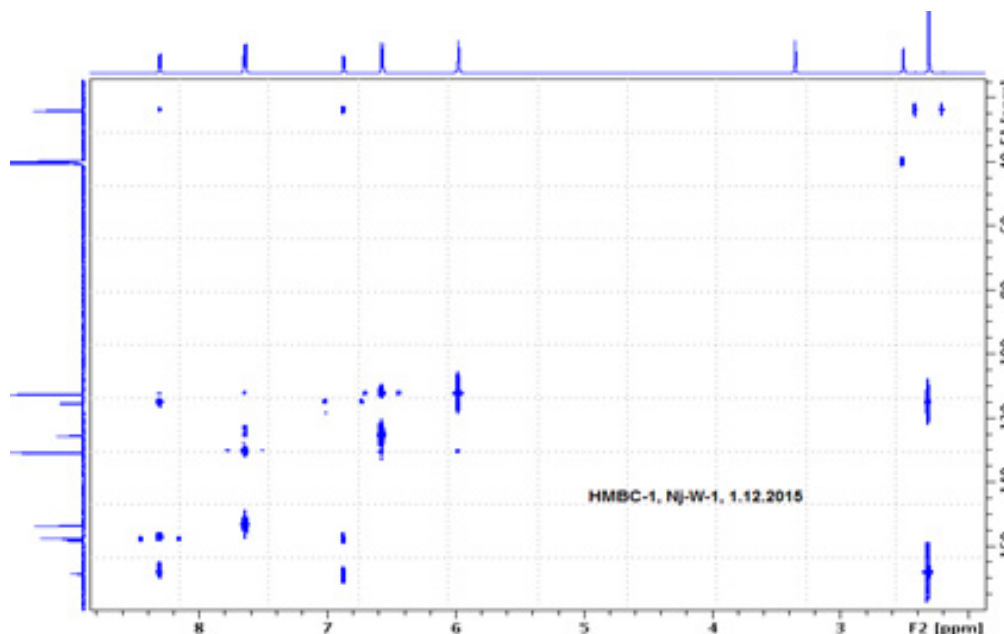
Fig. 2: ^{13}C NMR of synthesized compound

Fig. 3: HMBC- NMR of synthesized compound

Antimicrobial activity

Even though pharmacological industries have produced a number of new antibiotics in the last three decades, resistance to these drugs by microorganisms has increased. In general, bacteria have the genetic ability to transmit and acquire resistance to drugs, which are utilized as therapeutic agents¹¹. In the present work the studied compound are evaluated *in vitro* for their

antibacterial and antifungal activities by using paper disc-agar diffusion technique¹⁰ with the microorganisms as seen in Table 1. Antibiotic drug ampicillin and Nystatin were used as control for bacteria and fungi, respectively.

Table 1: The antibacterial and antifungal activity of the studied compounds(inhibition zone mm)

Bacteria and fungi types	Zone inhibition of antibiotic disc (control)(mm)		Zone inhibition of antimicrobial sensitivity test of compounds (mm)					
		Compound 1(S) µg/ml	Compound 2 µg/ml					
	Ampicillin 25µg/ml	Nystatin 30µg/ml	200	100	50	200	100	50
Staphylococcus aureus	22	-	18	15	12	16	12	10
Escherichia coli	20	-	15	12	12	11	10	10
Klebsiella pneumonia	25	-	14	12	10	10	8	6
Bacillus cereus	30	-	0	0	0	0	0	0
Candida krusei	-	11	30	25	20	21	16	15
Candida tropicalis	-	-	15	12	10	18	16	12
Candida albicans	-	-	18	15	12	18	15	12
Aspergillus niger	-	11	15	14	9	20	17	14
Aspergillus fumigatus	-	10	12	10	6	15	15	12

Molecular modeling analysis

The molecular docking was performed using SYBYL-X 1.1 and the docking results were shown by PyMOL¹². Our molecular docking analysis of the new analogue based on the modeling study which was performed to understand the binding mode of these analogues with UDP-N-acetylglucosamine 1-carboxyvinyl transferase of *E. coli*¹³ binding pocket (PDB code: 1ahg¹⁴). Compound **2** has been selected for the docking modelling study, since its binding energy score

-9.8, with indicating a selectivity of 4-((benzo[d][1,3]dioxo-5-ylmethylene)amino)-N-(5-methylpyrimidin-2-yl)benzenesulfonamide in binding to the enzyme pocket (Figure 4) via its and its azo and benzo[d][1,3]dioxo groups. As shown in Figure 4, the aromatic ring of new compound was fitted into an aromatic rich sub-pocket surrounded by the aromatic side chains of Phe328 showing an hydrophobic interaction via the pi stacking, in addition to two hydrogen bonding were observed. The azo-sulfamethazine backbone was located in the middle of the binding pocket, anchoring the nitrogen atom of the azo group in a favourable position for hydrogen bonding with the NH₂ of the amino acid of *E. coli* Asn23, whereas the other hydrogen bond was assigned between the oxygen atom of benzo[d][1,3]dioxo group

and the NH₂ group of Asn305. Overall, the combination of hydrophobic interaction and pi stacking appears to govern the binding of new compound with amino acids of transferase of *E. coli*. Furthermore, the amino acids of the enzyme: Tyr399 and Ala374 have surrounded the sulfamerazine backbone.

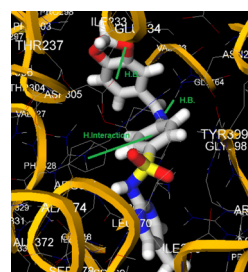


Figure 4: Docked conformation of synthesized compound showing two hydrogen bonds: NH₂ group of Asn305 with oxygen atom of benzo[d][1,3]dioxo moiety, and Asn23 with nitrogen atom of the azo group.

It also exhibits also hydrophobic interaction between phenyl ring of sulfamethazine residue and Phe328 of UDP-GlcNAc enolpyruvyl transferase enzyme residue UDP-N-acetylglucosamine 1-carboxyvinyl transferase of *E. coli* (PDB: 1A2NMurA (UDP-GlcNAc enolpyruvyl transferase), the first enzyme in bacterial peptidoglycan biosynthesis, catalyzes the enolpyruvyl transfer from phosphoenolpyruvate (PEP) to the 3'-OH of UDP-GlcNAc by an addition-elimination mechanism that proceeds through a tetrahedral ketal

intermediate. The crystal structure of the Cys115-to-Ala is shown in Fig. 5.



Fig. 5: UDP-N-acetylglucosamine 1-carboxyvinyltransferase of *E. coli*

Discussion

The search of novel antimicrobial agents still continues as the available antibiotics are not active against multi drug resistant pathogenic strains. In current study the Schiff base derived from sulfamerazine is synthesized and their chemical structure was confirmed by IR and NMR spectral analysis. The IR spectra confirm the presence of the azomethine group ($-\text{CH}=\text{N}$) stretching with a sharp region around 1540 cm^{-1} . ^1H NMR spectra of synthesized compound shows signal due to azomethine proton ($\text{CH}=\text{N}$) at 8.31 ppm . ^1H NMR spectra of synthesized compound show a singlet at 11.13 ppm due to NH . The region at $7.65\text{--}5.98\text{ ppm}$ due to aromatic protons. ^1H NMR spectra of synthesized compound shows singlet's at 3.35 and 2.31 ppm respectively, due to aliphatic protons, Fig. 1.

The ^{13}C NMR spectrum of synthesized compound was measured in $\text{DMSO-}d_6$. ^{13}C NMR spectra gave further support to the formation of new compound. The spectra revealed the presence of $\text{CH}=\text{N}$ group around 158.1 ppm . The signals around $157.4\text{--}112.5\text{ ppm}$ due to C-aromatic groups. The signal at 168.4 ppm due to C-SO_2 group, Figure 2. The two dimensional ^1H , ^{13}C -HMBC NMR spectra are support the chemical structure of synthesized compound, Figure 3.

The antibacterial activity of the synthesized compounds were tested against two Gram positive bacteria (*Staphylococcus aureus*, *Bacillus cereus*, and

two Gram negative bacteria (*Klebsiella pneumonia*, *Escherichia coli*) at a concentration of 50 , 100 and $200\text{ }\mu\text{g/mL}$ using DMSO as a solvent, which have no effect on the growth of microbes¹⁵. According to the results on Table (1) the antimicrobial activity of the tested compounds were showed a slight to intermediate activity against *Klebsiella pneumonia* and *Staphylococcus aureus* but show no activity against *Escherichia coli* and *Bacillus cereus*. On the other hand, the synthesized compounds were showed a good activity against all the fungi species. However, the compounds had the highest effect against *Candida krusei* and *Aspergillus niger* and intermediate activity against *Candida tropicalis*, *Candida albicans* and *Aspergillus fumigatus*. The difference of susceptibility of the tested microorganisms could be attributed to their major properties of microorganisms that are related to the permeability of their cell surface to the antibiotics and drugs.

Conclusion

In conclusion the studied compound could be a probable source to obtain effective against multi-drug resistant strains of microorganisms. Conversely, it is necessary to calculate the MIC index (MBC/MIC) and to determine the toxicity and the side effects of the studied compounds.

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Disclosure statement

Conflict of Interests: The authors declare that they have no conflict of interest.

Author contributions: All authors contributed equally to this work.

Ethical Clearance: All ethical guidelines have been adhered according to committee on the ethics of dealing with laboratory animals.

Source of Funding: Self

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