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Antibacterial and molecular docking studies of newly synthesized nucleosides and Schiff bases derived from sulfadimidines

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A new series of nucleosides, moieties, and Schiff bases were synthesized from sulfadimidine. Infrared (IR), ¹HNMR, ¹³C NMR, and mass spectrometry techniques and elemental analysis were employed to elucidate the synthesized compounds. The prepared analogues were purified by different chromatographic techniques (preparative TLC and column chromatography). Molecular docking studies of synthesized compounds 3a, 4b, 6a, and 6e demonstrated the binding mode involved in the active site of DNA gyrase. Finally, all synthesized compounds were tested against selected bacterial strains. The most effective synthesized compounds against *S. aureus* were 3a, 4d, 4b, 3b, 3c, 4c, and 6f, which exhibited inhibition zones of inhibition of 24.33 ± 1.528, 24.67 ± 0.577, 23.67 ± 0.577, 22.33 ± 1.528, 18.67 ± 1.528 and 19.33 ± 0.577, respectively. Notably, the smallest zones were observed for 4a, 6d, 6e and 6g (6.33 ± 1.528, 11.33 ± 1.528, 11.67 ± 1.528 and 14.66 ± 1.155, respectively). Finally, 6b and 6c gave negative zone values. *K. pneumoniae* was treated with the same compounds and the following results were obtained. The most effective compounds were 4d, 4c, 4b and 3c, which showed inhibition zones of 29.67 ± 1.528, 24.67 ± 0.577, 23.67 ± 1.155 and 19.33 ± 1.528, respectively, followed by 4a and 3d (15.33 ± 1.528 for both), while moderate results (13.67 ± 1.155 and 11.33 ± 1.528) were obtained for 6f and 6g, respectively. Finally, 6a, 6b, 6c, 3a, and 3b did not show any inhibition. The most effective compounds observed for the treatment of *E. coli* were 4d, 4b, 4c, 3d, 6e and 6f (inhibition zones of 26.33 ± 0.577, 21.67 ± 1.528, 21.67 ± 1.528, 19.67 ± 1.528, 17.67 ± 1.155 and 16.67 ± 1.155, respectively). Compounds 3b, 3c, 6a, 6c, and 6g gave moderate results (13.67 ± 1.528, 12.67 ± 1.528, 11.33 ± 0.577, 15.33 ± 1.528 and 12.67 ± 1.528, respectively), while 6b showed no effect. The MIC values against *S. aureus* ranged from 50 to 3.125 mg, while those against *E. coli* and *K. pneumoniae* ranged from 50 to 1562 mg. In vitro, the antibacterial effects were promising. Further research is required to study the in vivo antibacterial effects of these compounds and determine therapeutic doses.

Abbreviations

AcOH	Acetic acid
Ac ₂ O	Acetic anhydride
ATCC 25923	<i>Staphylococcus aureus</i> Subsp. <i>aureus</i> Rosenbach
ATCC 6633	<i>Bacillus subtilis</i> Subsp. <i>spizizenii</i> Nakamura et al.
ATCC 14028	<i>Salmonella enterica</i> Subsp. <i>enterica</i> (ex Kauffmann and Edwards) Le Minor and Popoff serovar Typhimurium
ATCC 12022	<i>Shigella flexneri</i> Castellani and Chalmers
EtOH	Ethanol
POA 286	<i>P. aeruginosa</i>

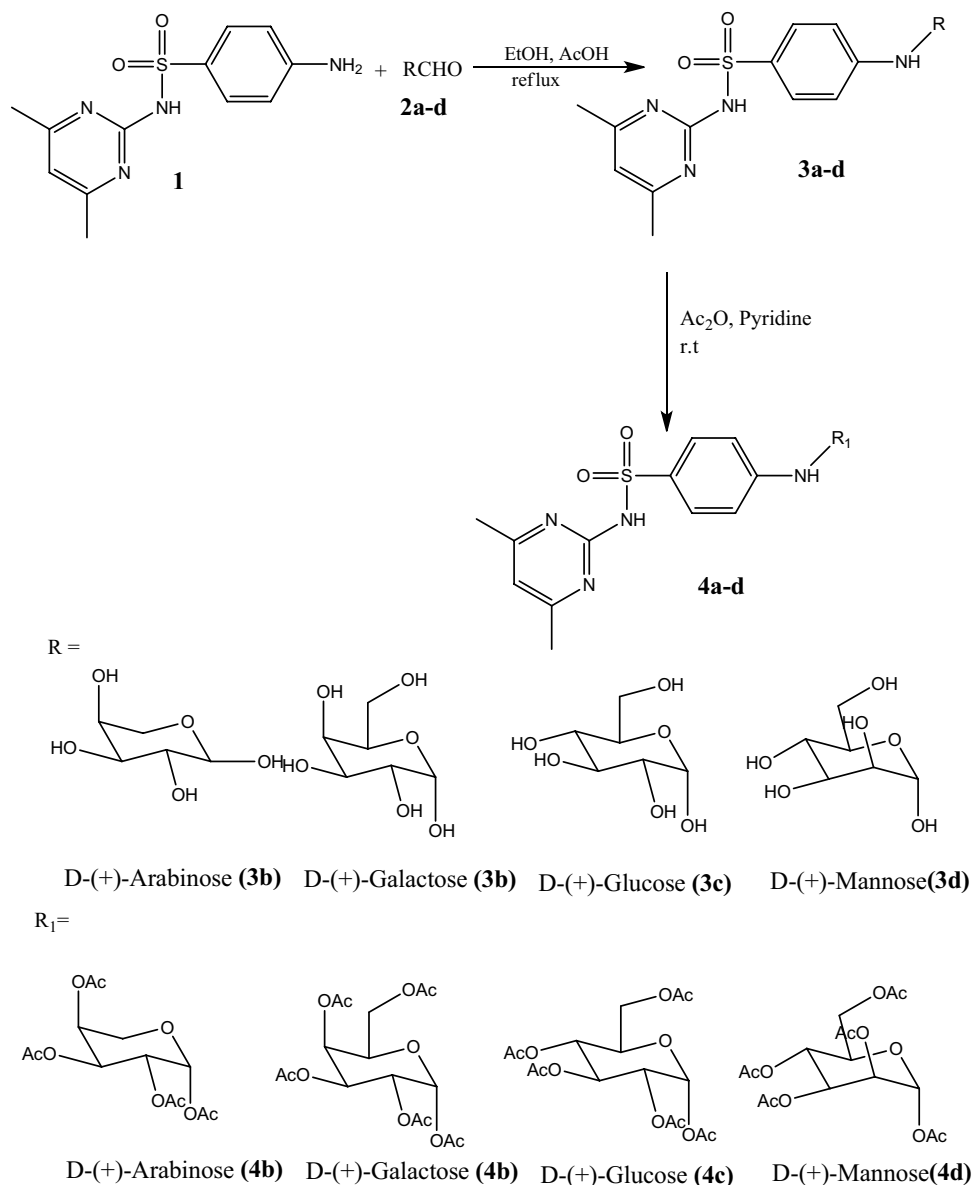
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MIC	Minimum inhibitory concentration
VERO	VERO cell
RMSD	Root mean square deviation
PBM	Pharmacy Benefits Management
CEM	Construction Engineering and Management
MOE	Molecular Operating Environment
MD	Molecular dynamics
NAMD	National Association of Medicaid Directors
EHT	Extended Hückel Theory
PLIFs	Protein–ligand interaction fingerprints
CDCl ₃	Deuterated chloroform
TMS	Trimethyl silane
TLC	Thin layer chromatography
DFT	Discrete Fourier transform
LDF-9025U	Deep freezer LDF 9025U
OD600 0.2	Orion (Knight Electronics, Inc.)
DMSO	Dimethyl sulfoxide

A large number of new sugar hydrazones composed of monosaccharides and heterocyclic rings have reportedly been synthesized from quinoxaline derivatives. By testing the synthesized sugar hydrazones against bacteria, it was found that they have very high activity compared to standard drugs¹. The newly synthesized heterocyclic compounds containing a sulfonamide group showed high activity against microbes in general², and bacteria in particular. Nitrogen-containing heterocyclic compounds associated with sulfonamide groups have received great interest in drug development due to their well-known activity in the fields of pharmaceutical and medicinal chemistry. Heterocyclic sulphonamides have been used as carbonic anhydrase inhibitors^{3–5}. Pyrimidines constitute a large group of heterocyclic compounds that have contributed greatly to the discovery of new drugs due to their known biological activity as antibacterial⁶, anti-inflammatory⁷, analgesic⁸ and anticancer drugs⁹. Schiff bases are one of the most extensively used compound types in the fields of organic and medicinal chemistry due to their strong effects as new drugs. Schiff bases derived from heterocyclic aldehydes have shown good anti-fungal and bacterial activities¹⁰. Nucleosides have been of great importance in medicinal and therapeutic chemistry for a long time due to their anti-viral, anti-parasitic¹¹, antibacterial¹² and anticancer properties¹³. Derivatives of sulfamethazine that are predicted to have high biological activity were designed, synthesized, and tested for their antibacterial activity relative to the antibiotic ofloxacin as the standard antibiotic against *P. aeruginosa* (POA 286), *S. aureus* (ATCC 25923), *B. subtilis* (ATCC 6633), *Salmonella typhi* (ATCC 14028), and *E. coli* (ATCC 12022). The results illustrated that most of the new compounds showed moderate activity against the above-mentioned bacterial strains¹⁴. Five series of sulfonamide derivatives derived from coumarin were also synthesized, and their biological activities were assessed as clinically important gram-positive and gram-negative antibacterial drugs and the derivatives were also evaluated for their in vitro antibacterial activity against selected clinically important strains, such as *Salmonella typhi*, *S. aureus*, *P. aeruginosa* and *B. cereus*. In addition, they were assessed as antifungal compounds against pathogenic diseases such as those caused by *Curvularia lunata*, *Aspergillus niger*, *Trichoderma viride* and *Aspergillus flavus*. The results showed that the compounds containing sulfonamide groups displayed good activity against the mentioned bacterial and fungal strains¹⁵. A new series of heterocyclic compounds containing sulfonamides were synthesized and tested for their antibacterial activity. These compounds were found to be highly active (MIC values of 50–3.1 µg/mL) against *P. aeruginosa*, *S. aureus*, *E. coli*, and *B. cereus*; the highest activity was shown against *E. coli*. By reducing their MIC value to 1/4 or 1/32 of their original value, some of the compounds showed 4–32-fold stronger activity relative to the MIC of the reference antibiotic. Additionally, these compounds showed a low degree of cytotoxicity against the VERO, PBM and CEM cell lines¹⁶. A group of new nucleosides, oxadiazole and arylidene derivatives were synthesized, purified by different chromatographic methods and chemically elucidated by infrared and ¹H and ¹³C nuclear magnetic resonance spectroscopic techniques. The synthesized compounds were tested against three different strains of bacteria. The results showed high to moderate efficacy against the tested bacterial strains. These studies were compared with theoretical reports, such as those that included molecular docking of the compounds, revealing that of one of these compounds bound to the active site of DNA gyrase¹⁷. Phthalimide derivatives and their copper complexes have shown antibacterial activity against three bacterial strains, namely, *S. enterica*, *E. faecalis*, and *S. pneumoniae*¹⁸. Among the most active compounds that reportedly exhibited antibacterial activity were the Schiff bases that were prepared by condensation of sulfonamides such as sulfamethoxy pyridazine, which is a compound in the sulfadimidine family with an aldehyde moiety derived from naphthalene, and notably, their complexes were found to be effective antibacterial agents¹⁹. In this study¹⁴, Schiff bases were synthesized from 6-(phenyl)-N'-((E)-(phenyl) methylidene)-2-methylpyridine-3-carbohydrate and tested as anti-gram-positive (*S. aureus* and *E. faecalis*) and anti-gram-negative (*E. coli* and *P. aeruginosa*) compounds. The synthesized compounds showed high antibacterial activity at a concentration lower than the standard value²⁰. In this work, new nucleosides and Schiff bases were synthesized from sulfadimidine and screened against gram-positive and gram-negative bacteria (Supplementary Information).

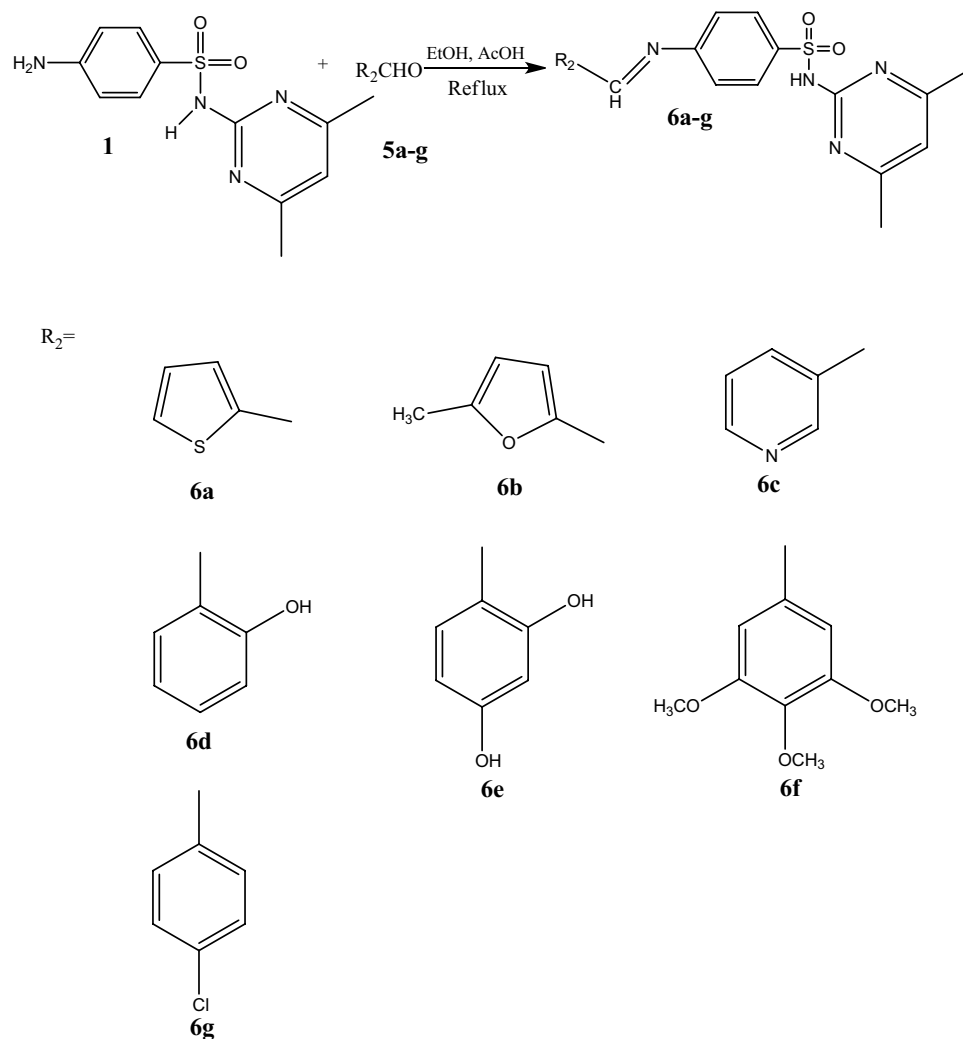
Results and discussion

Chemistry. Sulfadimidine (1) was reacted with different sugar moieties (2a–d) in the presence of a small amount of catalyst (AcOH) using absolute ethanol as the solvent and heated in a condenser to give the corresponding nucleoside derivatives 3a–d in 82–88% yields (Scheme 1). Elucidation of compounds 3a–d by IR



Scheme 1. Synthesis of some Nucleoside derivatives.

spectroscopy showed the disappearance of the (NH₂) group of sulfadimidine (1) and the appearance of the NH group at 3400 cm⁻¹ and the hydroxyl groups of the nucleoside derivatives at 3360 cm⁻¹. The ¹H NMR spectra indicated multiplet peaks at approximately 3.57–4.95 for the CH groups, 3.50 and 3.80 for the OH groups and 8.36 for the NH group in the sugar moieties. The ¹³C NMR spectra showed a peak at 24.24 for the methyl groups of the pyrimidine ring, peaks at approximately 65.00 and 94.00 for the CH group of the sugar moieties, and at 61.93 for the hydroxyl groups of the sugar moieties in compounds 3b, 3c and 3d. The CH aromatic peaks appeared between approximately 110.11 to 168.60. The mass spectra showed molecular ion peaks at 410 [M]⁺, 441 [M+H]⁺, 442 [M+2H]⁺ and 440 [M]⁺ for 3a, 3b, 3c and 3d, respectively. Acetylated nucleosides 4a–d were obtained in 85–93% yields by the acetylation of nucleoside derivatives 3a–d upon reaction with acetic anhydride in dry pyridine at room temperature (Scheme 1). The elucidation of compounds 4a–d by IR spectroscopy showed the disappearance of a peak the 3360 cm⁻¹ attributed to the OH group of the nucleosides in 3a–d and the appearance of a peak at 1740 cm⁻¹ attributed to the acetyl group (COCH₃) of the acetylated nucleoside derivatives 4a–d. The ¹H NMR spectra displayed the disappearance of the peaks from the hydroxyl groups of the sugar moieties between 3.50 and 3.80 and the appearance of multiplet peaks at 2.02 and 2.23 attributed to the acetyl groups (COCH₃), multiplet peaks at approximately 3.78–5.36 for the CH groups of the sugar moieties and at approximately 8.45 for the NH group. The ¹³C NMR spectra showed a peak at 21.00 for the methyl group of the acetyl moiety (COCH₃) and at approximately 24.24 for the methyl groups of the pyrimidine ring. The peaks observed between approximately 61.31 to 90.47 were attributed to the CH group in the sugar moieties, and those at 62.41 were caused by the methylene groups in the sugar moieties of compounds 4b, 4c, and 4d. The CH-aromatic peaks appeared at approximately 110.18 to 168.63 and that of the acetyl carbonyl group was



Scheme 2. Synthesis of some Schiff bases.

at approximately 170.47. The mass spectra showed molecular ion peaks at 539 $[\text{M}]^+$, 608 $[\text{M}]^+$, 609 $[\text{M}+\text{H}]^+$ and 608 $[\text{M}]^+$ for compounds 4a, 4b, 4c and 4d, respectively. Among the nucleoside derivatives, two isomers, E and Z, were present.

Sulfadimidine (1) was reacted with different aromatic aldehydes 5a–g using absolute ethanol as the solvent in the presence of small amounts of catalyst (AcOH) and heated using a condenser to give the corresponding Schiff base derivatives 6a–g in 92–96% yields (Scheme 2). Elucidation of compounds 6a–g by IR spectroscopy showed the disappearance of the NH_2 group of sulfadimidine (1) and the appearance of the peak attributed to the NH group at 3370 cm^{-1} and hydroxyl groups of certain Schiff bases at 3450 cm^{-1} . The ^1H NMR spectra displayed a singlet peak at 2.35 attributed to the methyl groups on the pyrimidine ring, a peak between approximately 5.43–5.76 for the OH groups of the Schiff bases, and multiplet peaks between approximately 6.56–8.50 for Ar-CH and at 9.53 for the NH group. The ^{13}C NMR spectra showed a peak at 22.86 for the methyl groups on the pyrimidine ring and at approximately 54.43 and 61.21 for the methoxy groups (OCH_3). The peaks between approximately 104.03–168.58 were attributed to the CH aromatics, and those at 163.22 were attributed to CH=N groups. The mass spectra showed molecular ion peaks at 372 $[\text{M}]^+$, 370 $[\text{M}]^+$, 367 $[\text{M}]^+$, 382 $[\text{M}]^+$, 299 $[\text{M}+\text{H}]^+$, 458 $[\text{M}+2\text{H}]^+$, and 400 $[\text{M}]^+$ for compounds 6a, 6b, 6c, 6d, 6e, 6f and 6g, respectively.

Docking studies. The docking study used DNA gyrase B PDB ID: 4uro 14 to examine the mode of action of the small molecule compounds as antimicrobial agents. The ligand–protein interaction behaviours were estimated based on the docking score function, as implemented in the Molecular Operating Environment (MOE) 2015.10 package. All calculations of the docking experiment against the crystal structure (PDB: 4uro) are presented in Table 4. Bacterial DNA gyrase plays a vital role during the investigation of antibacterial agents, as it breaks double-stranded DNA by catalysing negative supercoiling, which is essential for DNA replication, transcription, and recombination¹⁵. Analysis of the co-crystallized DNA-gyrase cleavage complex with novobiocin, which is an effective antibacterial agent that cleaves DNA by restricting the ATPase binding site including the vital naked peptidoglycan of the cell wall (Ser55, Ala64, Asn65, Asp89, Thr164, Thr173, and Val79), was per-

Cpd	E.dG	rmsd_	E_place	E_conf	Eele	E.Int	Eele _{MD}	E.Int _{MD}
3a	-5.99	1.02	-136.520	-86.518	-11.286	-20.913	-13.27	-22.00
3b	-5.77	1.66	-298.362	-106.58	-9.846	-10.369	-11.17	-12.74
6a	-4.93	1.52	-158.321	-52.361	-10.685	-21.604	-13.685	-26.02
6e	-4.65	1.23	-358.365	-192.754	-13.971	-29.471	-13.971	-27.02

Table 1. Simulated molecular docking energies (kcal/mol) of the tested ligands. *Ed.G.* final free binding energy, *E_conf* binding energy for the ligand-conformer, *E_place* binding energy of the ligand-receptor, *E.Int.* ligand-receptor affinity interaction energy, *E.ele* ligand-receptor electrostatic interaction, *RMSD* root mean square deviation of the docking pose compared to the co-crystal ligand position, *E.Int_{MD}* interaction energy obtained from molecular dynamics, *E.ele* electrostatic interaction obtained from molecular dynamics.

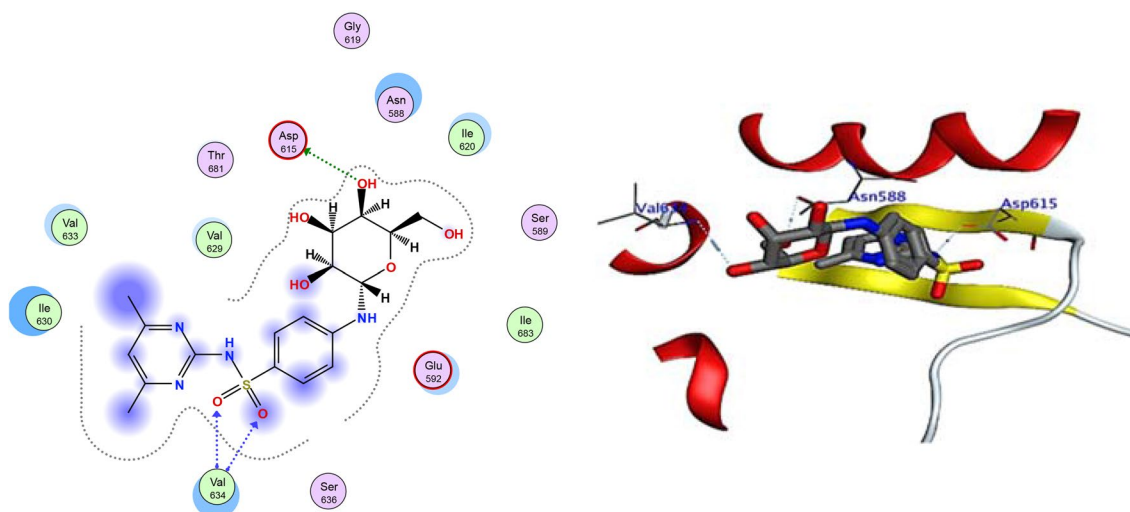


Figure 1. The 2D and 3D binding modes of 3a in the active site of DNA gyrase. H-bonds are represented by blue dashed lines.

formed. The inhibitory effects may be attributed to the distinct differences in the structures of the cell walls between gram-negative and gram-positive bacteria. The cell walls of gram-negative bacteria are composed of a thin peptidoglycan layer (7–8 nm) with an additional outer membrane. Gram-positive bacteria have a thick peptidoglycan layer (20–80 nm) outside the cell wall without an outer membrane. Peptidoglycan is a mesh-like polymer consisting of sugars and amino acids. The peptidoglycan layer protects the microorganism against antibacterial agents such as antibiotics, toxins, chemicals, and degradative enzymes^{16,17}. All of the compounds were redocked into the active site without a reference inhibitor and the ligands were successfully complexed with the active sites of the enzyme. The extracted docked poses of the ligands were energy-minimized using the molecular mechanics force field Amber12:EHT until a gradient convergence of 0.05 kcal/mol was achieved. The poses with a docking score of $E_d.G < -7$ kcal/mol were initially selected. These poses were filtered on the basis of the lowest MOE docking score, which was attributed to those with the smallest root mean square deviation (RMSD) against the reference drugs. The highest MOE scoring function was applied for the tested compounds to evaluate their binding affinities (Table 1). Compound 3a exhibited the highest binding affinity and RMSD to DNA gyrase compared with the other compounds (-5.99 kcal/mol.), as summarized in Table 1. The highest antimicrobial activity observed for compound 3a may be explained by the strong interaction with the binding site through the formation of H-bonds with important amino acids, namely, two bonds with Asp615 and two with Val634 (Fig. 1). The considerable RMSD value of < 2 was attributed to the high stability of these compounds in the binding site (Table 1). The binding energy for 4e was found to be -5.77 kcal/mol, and it interacted with the binding pocket by forming 3 H-bonds with Asp615, Asn588, and Val634 (Fig. 2). Compounds 6a and 6e showed the lowest binding energies (-4.93 and -4.65 kcal/mol). Additionally, compound 6a formed two H-bonds with Thr171 and Ala64 and compound 6e exhibited a π -H bond with Glu592 (Table 2; Figs. 3 and 4). Based on the different interaction modes of the ligands with the hydrophilic amino acid backbone in the gyrase binding site (Fig. 1), it was postulated that the hydrophilic fragments in the synthesized compounds (Asp and Asn) play important roles on the binding of molecules to DNA gyrase. Furthermore, the inhibitory efficiency of the synthesized compounds against bacterial growth may be attributed to the attack of the naked peptidoglycan in the cell wall. Therefore, these compounds were inferred to have good biological functions.

Antibacterial activity. The results of the sensitivity test are shown in Table 3 and Figs. 5 and 6, which include testing of the synthesized compounds against the three bacterial strains by measuring the sizes of the inhibition zones. The most effective synthesized compounds against *S. aureus* were 3a, 4d, 4b, 3b, 3c, 4c, and 6f

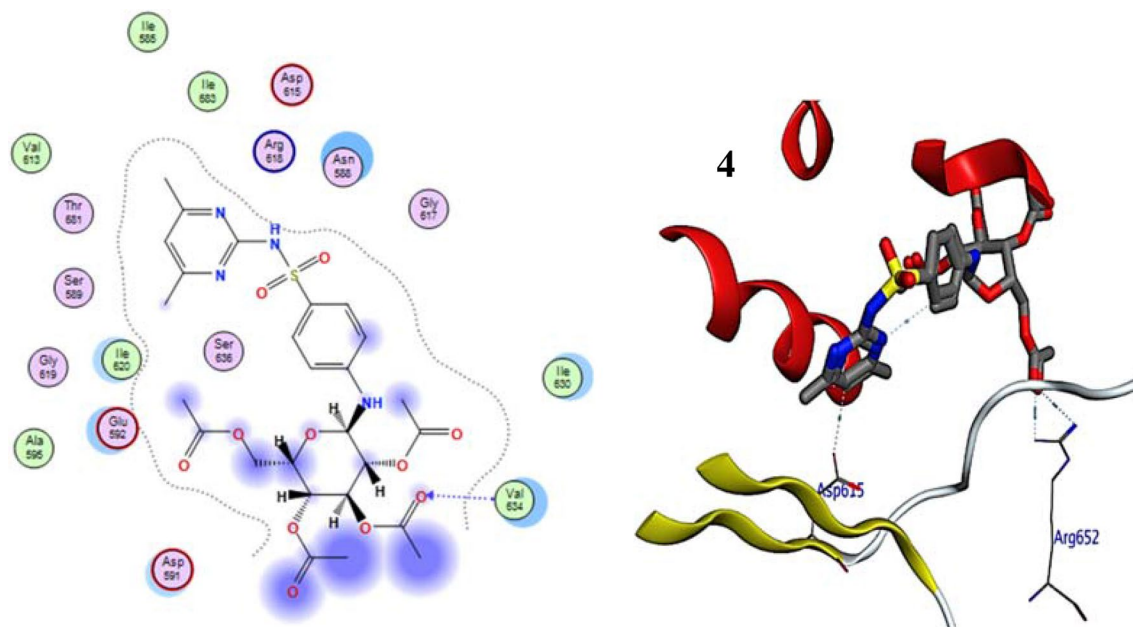


Figure 2. The 2D and 3D binding modes of 4b in the active site of DNA gyrase. H-bonds are represented by blue dashed lines.

Ligand	Receptor	Interaction	Distance	E (kcal/mol)	
Compound 3a					
O38	Asp	615	H-donor	3.01	-1.2
O38	Asp	615	H-donor	2.99	-2.6
O14	Val	634	H-donor	2.89	-9.5
O15	Val	634	H-donor	3.0	-8.0
Compound 4e					
N16	ASP	615	H-donor	2.47	-1.1
O52	ASN	588	H-donor	2.90	-0.7
S38	Val	634	H-acceptor	2.24	-4.1
Compound 6a					
6-ring	THR	171	H-donor	2.87	-1.4
O10	ALA	64	H-acceptor	2.87	-0.9
Compound 6e					
6-ring	GLU	592	π-H	2.35	-4.00

Table 2. Key interactions of the newly designed ligands with the active site of DNA gyrase B (PDB ID: 4uro).

Moreover, the smallest zones were observed with 4a, 6d, 6e, and 6g. Finally, 6b and 6c gave negative zone values. *K. pneumoniae* was tested with the same compounds and displayed the following results. The most effective compounds were 4d, 4c, 4b, and 3c, followed by 4a and 3d, and moderate results were obtained for 6f and 6g. Finally, compounds 6a, 6b, 6c, 3a, and 3b showed no inhibition. The most effective compounds observed against *E. coli* were 4d, 4b, 4c, 3d, 6e, and 6f, followed by moderate results for 3b, 3c, 6a, 6c, and 6g. Finally, 6b showed no effect at all. The inhibition zones with ciprofloxacin were all positive. The zones against *S. aureus* showed a value of 21.22 ± 1.55 mm, while the zones against *K. pneumoniae* were 19.22 ± 1.23 mm and those with *E. coli* were 18.52 ± 1.44 mm.

An MIC test was included in this study to determine the lowest concentration of compound that give the lowest inhibition zone using the agar well diffusion test. The results showed that the highest dilutions for the treatment of *S. aureus* were obtained for 3c, 4b, 4c, and 4d, followed by 3d and 6g. Compounds 4a, 6e, and 6f gave the lowest values, and finally, 3a, 6b, 6c showed no inhibition zones at all. Compounds 4b, 3c, 3d, 3a, and 6e showed the best activity against *E. coli*, followed by 6a, 6c, 6f, and 6g, while 6d and 3a gave the lowest dilution values, and 3b, 4c, 4d, and 6b gave negative results. Finally, the reported results against *K. pneumoniae* showed that compounds 4b, 3d, and 3c gave the best results, followed by 6f and 6g. Notably, 4a gave the smallest result, and 3a, 3b, 4c, 4d, 6a, 6b, 6c, 6d, and 6e gave negative results (Table 4; Fig. 5). The MIC values of the sulfonamide derivatives against the *S. aureus* isolate were also evaluated. The MIC values ranged from 50 to 3.125 mg.

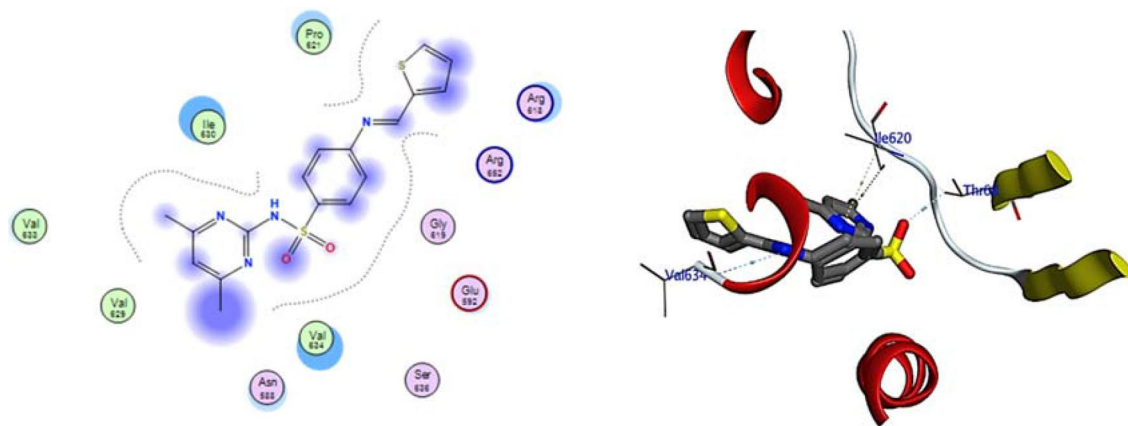


Figure 3. The 2D and 3D binding modes of 6a in the active site of DNA gyrase. H-bonds are represented by blue dashed lines.

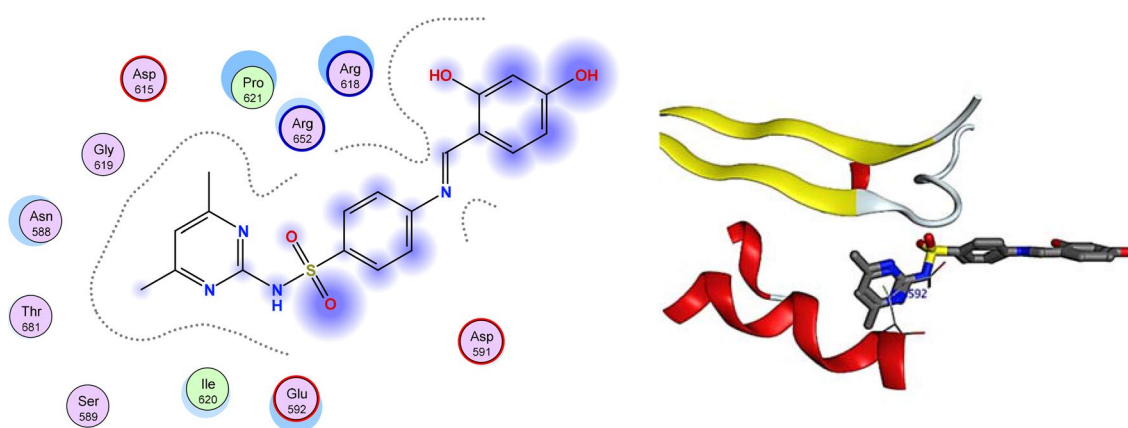


Figure 4. The 2D and 3D binding modes of 6e in the active site of DNA gyrase. H-bonds are represented by blue dashed lines.

However, the values ranged from 50 to 1.562 mg for *E. coli* and *K. pneumoniae*^{21,22}. In this study, the synthesized sulfadimidine derivatives were tested for their antibacterial activity to develop lead compounds for the production of new antibacterial agents. The outcome of our study showed that the sulfadimidine derivatives were effective enough in comparison to previously reported agents²². It was previously reported that sulfadimidine showed the highest inhibitory effects against gram-positive bacteria, i.e., *S. aureus*, and gram-negative bacteria, i.e., *E. coli* and *K. pneumoniae*. The antibacterial activity against these three bacterial isolates was attributed to hydrogen bonding between the functional groups, such as hydroxyl, acetyl, and heterocyclic rings, and the bases in the bacterial DNA gyrase as previously discussed by the docking results.

Material and methods

Chemistry. The melting points were determined using a Kofler block apparatus without correction. The ¹H NMR spectra (results in δ) were acquired on a 400 MHz spectrometer at King Abdel-Aziz University, Jeddah, Saudi Arabia, using CDCl₃ as the solvent and TMS as the internal standard. IR spectra were recorded using KBr disks on an IC50 model FT-IR (Thermo) instrument at the University College of Turbah, Taif University, Turbah, Taif, Saudi Arabia. Mass spectra were obtained using a GC-MS instrument at Taif University, Taif, Saudi Arabia. TLC was performed using aluminium silica gel plates (60 F245) to monitor the progress of the reactions. The antibacterial activities of the synthesized compounds were evaluated at the Biology Laboratory, Department of Biology, Turabah University College, Taif University, Turabah, Saudi Arabia.

General procedure for the preparation of 4-amino-*N*-(4,6-dimethylpyrimidin-2-yl) benzene-sulfonamide nucleosides (3a–d). To a solution of sulfadimidine derivative 1 (1.39 g, 10 mmol) in absolute ethanol (50 mL), the appropriate sugar derivative (*L*-(+)-arabinose, *D*-(+)-galactose, *D*-(+)-glucose or *D*-(+)-mannose; 0.75 g, 0.9 g, 0.9 g or 0.9 g (10 mmol each), respectively) and a few drops of acetic acid were added. The mixture was refluxed for 10–14 h. The formed precipitate was filtered off, washed with absolute ethanol, dried, and recrystallized using an ethanol:methanol mixture (1:2) to give nucleosides 3a–d in 82–88% yields.

Inhibition zone by mm			
Substance code	<i>S. aureus</i>	<i>K. pneumoniae</i>	<i>E. coli</i>
3a	24.33 ± 1.528	nz	11.33 ± 1.528
3b	22.67 ± 1.155	nz	13.67 ± 1.528
3c	22.33 ± 1.528	19.33 ± 1.528	12.67 ± 1.528
3d	18.44 ± 1.432	15.33 ± 1.528	19.67 ± 1.528
4a	6.33 ± 1.528	15.33 ± 1.528	16.33 ± 0.577
4b	23.67 ± 0.577	23.67 ± 1.155	21.67 ± 1.528
4c	18.67 ± 1.528	24.67 ± 0.577	21.67 ± 1.528
4d	24.67 ± 0.577	29.67 ± 1.528	26.33 ± 0.577
6a	15.33 ± 0.577	nz	11.33 ± 0.577
6b	nz	nz	nz
6c	6.66 ± 2.082	nz	15.33 ± 1.528
6d	11.33 ± 1.528	nz	12.67 ± 1.528
6e	11.67 ± 1.528	8.44 ± 1.144	17.67 ± 1.155
6f	19.33 ± 0.577	13.67 ± 1.155	16.67 ± 1.155
6g	14.66 ± 1.155	11.33 ± 1.528	12.67 ± 1.528
Positive control (Ciprofloxacin)	21.22 ± 1.55	19.22 ± 1.23	18.52 ± 1.44
Negative control (DMSO)	NZ	NZ	NZ

Table 3. Antibacterial activity of the tested chemical substances. Inhibition zones are expressed as the mean ± SD (mm). *S. aureus*: *Staphylococcus aureus*, *K. pneumoniae*: *Klebsiella pneumoniae* and *E. coli*: *Escherichia coli*. Nz: no zone.

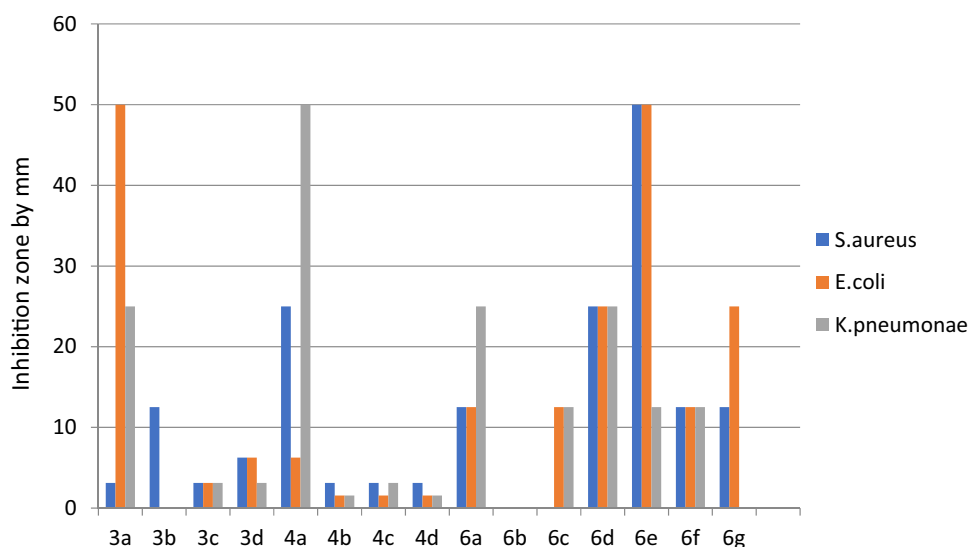


Figure 5. Sensitivity test of the synthesized compounds against three clinical isolates. Inhibition zones are expressed in mm.

N-Arabinofuranosylamino sulphadimidine (3a). Brown powder, 85% yield, m.p.=217–219 °C, Rf=0.50 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3400 (NH), 3360 (OH), 3055 (Ar-H), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.35 (6H, s, 2CH₃), 2.85 (1H, d, J=2.7 Hz, H-1), 3.57 (2H, d, 2xOH), 3.80 (1H, d, OH), 3.57–4.95 (4H, m, H-2, H-3, H-4, H-5), 6.67 (1H, s, CH-Pyrimidine), 7.17 (2H, dd, J=7.00 Hz, CH-aromatic), 7.83 (2H, dd, J=7.3 Hz, CH-aromatic), 8.36 (2H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 24.25 (CH₃), 65.23, 70.25, 73.76, 76.32, 94.00 (CH-Sugar), 110.17, 111.21, 128.22, 130.07, 150.92, 166.48, 168.61 (Ar-CH); MS: m/z (%) 410 (M)⁺. Anal. Calcd for C₁₇H₂₂N₄O₆S: C, 49.75; H, 5.40; N, 13.65. Found C, 49.88; H, 5.62; N, 13.46.

N-Glactopyranosylamino sulphadimidine (3b). White powder, 86% yield, m.p.=244–246 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3390 (NH), 3320 (OH), 3050 (Ar-H), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.32 (6H, s, 2CH₃), 3.59 (2H, dd, J=5.4 Hz, 2xOH), 3.59 (2H, dd, J=5.5 Hz,

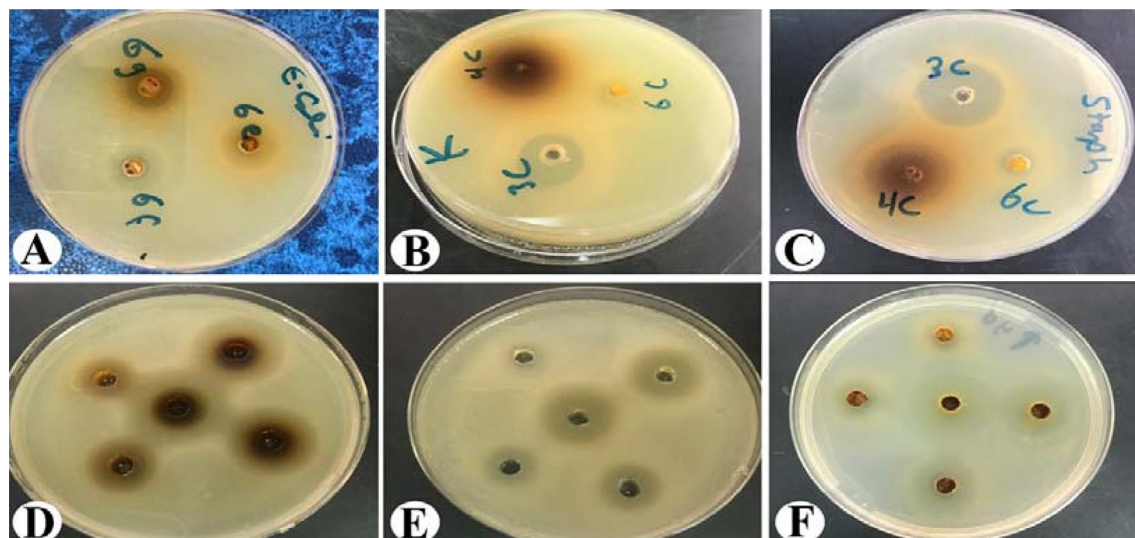


Figure 6. Sensitivity and minimum inhibitory concentration test (MIC) tests of the synthesized compounds against three clinical isolates [(A–C) sensitivity test; (D–F) MIC determinations by the agar dilution method].

Substance code	MIC minimum inhibitory concentration MIC by mg/mL		
	<i>S. aureus</i>	<i>E. coli</i>	<i>K. pneumoniae</i>
3a	3.125	50	25
3b	12.5	N	N
3c	3.125	3.125	3.125
3d	6.25	6.25	3.125
4a	25	6.25	50
4b	3.125	1.562	1.562
4c	3.125	1.562	3.125
4d	3.125	1.562	1.562
6a	12.5	12.5	25
6b	N	N	N
6c	N	12.5	12.5
6d	25	25	25
6e	50	50	12.5
6f	12.5	12.5	12.5
6g	12.5	25	12.5

Table 4. Minimum inhibitory concentration (MIC) test of the synthesized compounds against the three tested clinical isolates. N, no zone of inhibition was present.

2xOH), 4.58 (1H, d, $J=3.00$ Hz, H-1), 3.57–3.97 (6H, m, H-2, H-3, H-4, H-5, H-6), 6.65 (1H, s, CH-Pyrimidine), 7.15 (2H, dd, $J=7.00$ Hz, CH-aromatic), 7.82 (2H, dd, $J=7.3$ Hz, CH-aromatic), 8.22 (2H, brs, 2NH)); ^{13}C NMR (100 MHz, CDCl_3): δ : 24.25 (CH_3), 61.93 (CH_2OH of sugar), 71.24, 74.12, 77.45, 82.72, 91.60 (CH-Sugar), 110.17, 111.11, 128.17, 130.07, 150.92, 166.48, 168.61 (Ar-CH); MS: m/z (%) 441 (M+H)⁺. Anal. Calcd for $\text{C}_{18}\text{H}_{24}\text{N}_4\text{O}_7\text{S}$: C, 49.08; H, 5.49; N, 12.72. Found C, 48.87; H, 5.15; N, 12.40.

N-Glucopyranosylamino sulphadimidine (3c). Pale brown powder, 82% yield, m.p. = 256–258 °C, $R_f=0.40$ (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3390 (NH), 3320 (OH), 3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 2.30 (6H, s, 2 CH_3), 3.58 (2H, dd, $J=5.4$ Hz, 2xOH), 3.60 (2H, dd, $J=5.5$ Hz, 2xOH), 4.55 (1H, d, $J=3.00$ Hz, H-1), 3.57–3.95 (6H, m, H-2, H-3, H-4, H-5, H-6), 6.62 (1H, s, CH-Pyrimidine), 7.13 (2H, dd, $J=7.00$ Hz, CH-aromatic), 7.80 (2H, dd, $J=7.3$ Hz, CH-aromatic), 8.18 (2H, brs, 2NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 24.25 (CH_3), 61.93 (CH_2OH of sugar), 71.24, 74.12, 77.45, 82.72, 91.60 (CH-Sugar), 110.17, 111.11, 128.17, 130.07, 150.92, 166.48, 168.61 (Ar-CH); MS: m/z (%) 442 (M+2H)⁺. Anal. Calcd for $\text{C}_{18}\text{H}_{24}\text{N}_4\text{O}_7\text{S}$: C, 49.08; H, 5.49; N, 12.72. Found C, 49.32; H, 5.64; N, 12.87.

N-Mannopyranosylamino sulphadimidine (3d). White powder, 88% yield, m.p.=270–272 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3390 (NH), 3320 (OH), 3050 (Ar-H), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.30 (6H, s, 2CH₃), 3.53 (2H, dd, J=5.4 Hz, 2xOH), 3.58 (2H, dd, J=5.5 Hz, 2xOH), 4.59 (1H, d, J=3.00 Hz, H-1), 3.59–3.92 (6H, m, H-2, H-3, H-4, H-5, H-6), 6.66 (1H, s, CH-Pyrimidine), 7.20 (2H, dd, J=7.00 Hz, CH-aromatic), 7.87 (2H, dd, J=7.3 Hz, CH-aromatic), 8.25 (2H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 24.25 (CH₃), 61.93 (CH₂OH of sugar), 71.24, 74.12, 77.45, 82.72, 91.60 (CH-Sugar), 110.17, 111.11, 128.17, 130.07, 150.92, 166.48, 168.61 (Ar-CH); MS: m/z (%) 440 (M)⁺. Anal. Calcd for C₁₈H₂₄N₄O₇S: C, 49.08; H, 5.49; N, 12.72. Found C, 49.55; H, 5.56; N, 12.63.

General procedure for the synthesis of acetylated nucleosides 4a–d. A solution of nucleoside derivatives 3a–d (0.01 mol) in 15 mL of anhydrous pyridine was treated with acetic anhydride (0.01 mol). The reaction mixture was stirred for 14 h at room temperature and monitored by TLC. Each mixture was poured into 100 g of ice to give a pale yellow precipitate. The resulting precipitates were filtered, washed with water, and dried to produce acetylated nucleosides 4a–d in 85–93% yields.

4-(2, 3, 5-Tri-O-acetyl)-N-arabinofuranosylamino sulphadimidine (4a). Pale yellow powder, 93% yield, m.p.=265–267 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3350 (NH), 3050 (Ar-H), 1740 (COCH₃), 1625 (C=N), 1375 (CH₃), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.02, 2.11, 2.19 (9H, m, 3x COCH₃), 2.35 (1H, s, CH₃), 3.78–5.20 (4H, m, H-2, H-3, H-4, H-5), 5.36 (1H, d, J=5.5 Hz, H-1), 6.75–7.61 (5H, m, Ar-H), 8.45 (1H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 21.00 (3CH₃ of 3COCH₃ of sugar), 24.25 (2CH₃ of pyrimidine ring), 61.31, 68.23, 70.91, 72.79, 90.47 (CH-Sugar), 110.18, 111.21, 128.19, 130.09, 150.96, 166.50, 168.63 (Ar-CH), 170.47 (3C=O of 3COCH₃ of sugar); MS: m/z (%) 536 (M)⁺. Anal. Calcd for C₂₃H₂₈N₄O₉S: C, 51.49; H, 5.26; N, 10.44. Found C, 51.67; H, 5.47; N, 10.58.

4-(2, 3, 5, 6-Tetra-O-acetyl)-N-galactopyranosylamino sulphadimidine (4b). Pale yellow powder, 90% yield, m.p.=230–232 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3400 (NH), 3050 (Ar-H), 1735 (COCH₃), 1375 (CH₃), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.10, 2.13, 2.18, 2.23 (12H, m, 4xCOCH₃), 2.34 (CH₃), 4.22–5.35 (5H, m, H-2, H-3, H-4, H-5, H-6), 5.50 (1H, d, J=5.5 Hz, H-1), 6.76–7.60 (5H, m, Ar-H), 8.75 (1H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 20.87 (CH₃ of COCH₃ of sugar), 21.11 (3CH₃ of 3COCH₃ of sugar), 23.80 (2CH₃ of pyrimidine ring), 62.41 (CH₂), 68.98, 69.43, 73.08, 77.41, 87.86 (CH-Sugar), 110.18, 111.21, 128.19, 130.09, 150.96, 166.50, 168.63 (Ar-CH), 170.24 (4C=O of 4COCH₃ of sugar); MS: m/z (%) 608 (M)⁺. Anal. Calcd for C₂₆H₃₂N₄O₁₁S: C, 54.21; H, 4.71; N, 8.72. Found C, 54.41; H, 4.87; N, 8.45.

4-(2, 3, 5, 6-Tetra-O-acetyl)-N-glucopyranosylamino sulphadimidine (4c). Pale yellow powder, 85% yield, m.p.=243–245 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3400 (NH), 3050 (Ar-H), 1735 (COCH₃), 1375 (CH₃), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.08, 2.11, 2.17, 2.20 (12H, m, 4xCOCH₃), 2.35 (CH₃), 4.24–5.30 (5H, m, H-2, H-3, H-4, H-5, H-6), 5.45 (1H, d, J=5.5 Hz, H-1), 6.74–7.64 (5H, m, Ar-H), 8.75 (1H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 20.87 (CH₃ of COCH₃ of sugar), 21.11 (3CH₃ of 3COCH₃ of sugar), 23.80 (2CH₃ of pyrimidine ring), 62.41 (CH₂), 68.98, 69.43, 73.08, 77.41, 87.86 (CH-Sugar), 110.18, 111.21, 128.19, 130.09, 150.96, 166.50, 168.63 (Ar-CH), 170.24 (4C=O of 4COCH₃ of sugar); MS: m/z (%) 609 (M+H)⁺. Anal. Calcd for C₂₆H₃₂N₄O₁₁S: C, 54.21; H, 4.71; N, 8.72. Found C, 54.48; H, 4.89; N, 8.50.

4-(2, 3, 5, 6-Tetra-O-acetyl)-N-glucopyranosylamino sulphadimidine (4d). Pale yellow powder, 88% yield, m.p.=270–272 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3400 (NH), 3050 (Ar-H), 1735 (COCH₃), 1375 (CH₃), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.05, 2.10, 2.15, 2.18 (12H, m, 4xCOCH₃), 2.30 (CH₃), 4.25–5.32 (5H, m, H-2, H-3, H-4, H-5, H-6), 5.42 (1H, d, J=5.5 Hz, H-1), 6.74–7.64 (5H, m, Ar-H), 8.85 (1H, brs, 2NH); ¹³C NMR (100 MHz, CDCl₃): δ : 20.87 (CH₃ of COCH₃ of sugar), 21.11 (3CH₃ of 3COCH₃ of sugar), 23.80 (2CH₃ of pyrimidine ring), 62.41 (CH₂), 68.98, 69.43, 73.08, 77.41, 87.86 (CH-Sugar), 110.18, 111.21, 128.19, 130.09, 150.96, 166.50, 168.63 (Ar-CH), 170.24 (4C=O of 4COCH₃ of sugar); MS: m/z (%) 608 (M)⁺. Anal. Calcd for C₂₆H₃₂N₄O₁₁S: C, 54.21; H, 4.71; N, 8.72. Found C, 54.50; H, 4.90; N, 8.47.

General procedure for the synthesis of arylidene derivatives 6a–g. A solution of 1 (0.01 mol) and the appropriate aromatic aldehyde [thiophene-2-aldehyde, 5-methylfurfural, pyridine-3-carboxaldehyde, salicylaldehyde, 2,4-dihydroxybenzaldehyde, 3,4,5-trimethoxybenzaldehyde or p-chlorobenzaldehyde (0.01 mol)] were dissolved in 50 mL of ethanol. A small amount of acetic acid (1 mL) was added to the reaction mixture as the catalyst, which was refluxed for 10 h. The resulting product was filtered off to afford 6a–g in 92–96% yields.

N-(4,6-dimethylpyrimidin-2-yl)-4-((thiophen-2-ylmethylene)amino)benzenesulfonamide (6a). Pale yellow powder, 94% yield, m.p.=155–157 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3370 (NH), 3050 (Ar-H), 1150, 1305 (O=S=O); ¹H NMR (400 MHz, DMSO-d₆) δ : 2.35 (6H, s, 2CH₃), 6.75–7.89 (8H, m, Ar-H), 8.25 (1H, brs, CH), 9.78 (1H, brs, NH); ¹³C NMR (100 MHz, CDCl₃): δ : 24.23 (CH₃), 110.13, 122.62, 127.21, 124.13, 128.63, 130.27, 138.23, 132.43, 151.25, 166.69, 168.57 (Ar-CH), 152.81 (CH=N); MS: m/z (%) 372 (M)⁺. Anal. Calcd for C₁₇H₁₆N₄O₂S₂: C, 54.82; H, 4.33; N, 15.04. Found C, 54.98; H, 4.56; N, 14.97.

N-(4,6-dimethylpyrimidin-2-yl)-4-(((5-methylfuran-2-yl)methylene)amino)benzenesulfonamide (6b). Brown powder, 93% yield, m.p.=181–183 °C, Rf=0.45 (7% methanol in CHCl₃). IR spectra (KBr) (ν , cm⁻¹), 3370 (NH),

3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 2.12 (3H, s, CH_3), 2.30 (6H, s, 2CH_3), 6.65–8.50 (7H, m, Ar-H), 8.88 (1H, brs, CH), 9.86 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 16.50 (CH_3), 24.25 (2CH_3), 106.73, 110.11, 122.64, 128.62, 138.26, 147.33, 151.60, 155.65, 166.56, 168.67 (Ar-CH), 144.95 (CH=N); MS: m/z (%) 370 (M) $^+$. Anal. Calcd for $\text{C}_{18}\text{H}_{18}\text{N}_4\text{O}_3\text{S}$: C, 58.36; H, 4.90; N, 15.12. Found C, 58.47; H, 4.78; N, 15.43.

N-(4,6-dimethylpyrimidin-2-yl)-4-((pyridin-3-ylmethylene)amino)benzenesulfonamide (6c). White powder, 92% yield, m.p. = 167–169 °C, Rf = 0.45 (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3370 (NH), 3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 2.32 (6H, s, 2CH_3), 6.72–7.80 (9H, m, Ar-H), 7.95 (1H, brs, CH), 9.53 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 24.25 (2CH_3), 110.13, 122.61, 123.90, 128.60, 130.44, 133.75, 138.27, 149.04, 151.80, 151.92, 166.66, 168.54 (Ar-CH), 159.78 (CH=N); MS: m/z (%) 367 (M) $^+$. Anal. Calcd for $\text{C}_{18}\text{H}_{17}\text{N}_5\text{O}_2\text{S}$: C, 58.84; H, 4.66; N, 19.06. Found C, 58.70; H, 4.81; N, 18.97.

N-(4,6-dimethylpyrimidin-2-yl)-4-((2-hydroxybenzylidene)amino)benzenesulfonamide (6d). Yellow powder, 96% yield, m.p. = 210–212 °C, Rf = 0.45 (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3370 (NH), 3450 (OH), 3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 2.32 (6H, s, 2CH_3), 5.43 (1H, brs, OH), 6.75–7.75 (9H, m, Ar-H), 8.78 (1H, brs, CH), 9.75 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 24.25 (2CH_3), 110.15, 117.85, 120.44, 121.52, 122.58, 128.65, 132.21, 132.48, 138.37, 155.31, 161.13, 166.54, 168.62 (Ar-CH), 158.98 (CH=N); MS: m/z (%) 382 (M) $^+$. Anal. Calcd for $\text{C}_{19}\text{H}_{18}\text{N}_4\text{O}_3\text{S}$: C, 59.67; H, 4.74; N, 14.65. Found C, 59.87; H, 4.55; N, 14.76.

4-((2,4-dihydroxybenzylidene)amino)-*N*-(4,6-dimethylpyrimidin-2-yl)benzenesulfonamide (6e). Orange powder, 94% yield, m.p. = 190–192 °C, Rf = 0.45 (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3370 (NH), 3450 (OH), 3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 2.22 (6H, s, 2CH_3), 5.67 (2H, brs, 2OH), 6.45–7.80 (9H, m, Ar-H), 8.85 (1H, brs, CH), 9.92 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 23.35 (2CH_3), 103.43, 108.73, 110.25, 113.23, 122.67, 128.68, 133.89, 138.29, 155.24, 162.31, 162.54, 166.47, 168.56 (Ar-CH), 160.14 (CH=N); MS: m/z (%) 399 (M+H) $^+$. Anal. Calcd for $\text{C}_{19}\text{H}_{18}\text{N}_4\text{O}_4\text{S}$: C, 57.27; H, 4.55; N, 14.06. Found C, 57.45; H, 4.71; N, 14.25.

N-(4,6-dimethylpyrimidin-2-yl)-4-((3,4,5-trimethoxybenzylidene)amino)benzenesulfonamide (6f). White powder, 93% yield, m.p. = 190–192 °C, Rf = 0.45 (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3370 (NH), 3050 (Ar-H), 1375 (CH_3), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 1.98 (6H, s, 2CH_3), 3.65 (9H, s, 3OCH_3), 6.80–7.87 (7H, m, Ar-H), 8.76 (1H, brs, CH), 10.00 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 22.86 (2CH_3), 56.43 (2OCH_3), 61.21 (OCH_3), 104.03, 110.13, 113.23, 122.57, 128.46, 133.32, 138.43, 141.54, 153.23, 155.34, 166.61, 168.58 (Ar-CH), 163.22 (CH=N); MS: m/z (%) 458 (M+2H) $^+$. Anal. Calcd for $\text{C}_{22}\text{H}_{24}\text{N}_4\text{O}_5\text{S}$: C, 57.88; H, 5.33; N, 12.27. Found C, 57.77; H, 5.56; N, 12.48.

4-((4-chlorobenzylidene)amino)-*N*-(4,6-dimethylpyrimidin-2-yl)benzenesulfonamide (6g). Yellow powder, 92% yield, m.p. = 212–214 °C, Rf = 0.45 (7% methanol in CHCl_3). IR spectra (KBr) (ν , cm^{-1}), 3370 (NH), 3050 (Ar-H), 1150, 1305 (O=S=O); ^1H NMR (400 MHz, DMSO- d_6) δ : 1.88 (6H, s, 2CH_3), 6.73–7.85 (7H, m, Ar-H), 8.56 (1H, brs, CH), 9.56 (1H, brs, NH); ^{13}C NMR (100 MHz, CDCl_3): δ : 24.16 (2CH_3), 110.15, 122.60, 128.55, 128.98, 130.45, 134.76, 136.67, 138.54, 155.34, 166.71, 168.63 (Ar-CH), 160.00 (CH=N); MS: m/z (%) 400 (M) $^+$. Anal. Calcd for $\text{C}_{19}\text{H}_{17}\text{ClN}_4\text{O}_2\text{S}$: C, 56.93; H, 4.27; N, 13.98. Found C, 57.08; H, 4.47; N, 14.11.

Ethical approval. The ethical committee of Taif University, KSA, approved all procedures used in the current study.

Computational study

Preparation of the ligands and protein. From the RCSB protein data bank, the 3D structure of the protein was downloaded (ID: 4uro²³). To observe the data from the proteins, the BLAST P tool was employed to analyse the FASTA sequence. Then, the CLUSTALW18 package was utilized for multiple amino acid alignments, as reported earlier^{19,20,24–30}.

Procedure for the docking experiment. The docking study was completed with the MOE 2015 package 28 [Molecular Operating Environment (MOE)]. The tested compounds were constructed, and their energy was minimized on the basis of PM3. DFT studies of the compounds based on B3LYP/6–311G were performed. Error correction for the structure of the catalytic site in DNA gyrase B (ID: 4uro) was performed by the supplemented hydrogen and partial charges using Amber12:EHT and then minimized by using the same force field with an RMSD value of 0.100. The catalytic site was identified and analysed by the Site Finder program. This method was based on alpha spheres as well as the energy model29. The catalytic zone was predicted by the MOE-Site finder30. The water and inhibitor molecules were eliminated, and the H atoms were supplemented to obtain the crystal structure. The charges were regulated using the MMFF94x force field. The alpha-site spheres were added on the basis of the Site Finder module. The final energies, generated poses, and scores were measured using the procedures mentioned above³¹.

Antibacterial activity

Bacterial isolates. *S. aureus*, *E. coli*, and *K. pneumoniae* were obtained from the Microbiology Unit of King Faisal Hospital after ethical approval from the Taif Directorate of Health Affairs, Taif, Saudi Arabia. The *S. aureus* and *E. coli* strains were originally isolated from urine samples. The *K. pneumoniae* strain was isolated from wounds. The strains were identified phenotypically, biochemically, serologically, and genotypically and saved in the LDF-9025U at $-80\text{ }^{\circ}\text{C}$. Each isolate was cultivated separately in tryptic soy broth (Difco) at $37\text{ }^{\circ}\text{C}$ for 24 h. The cells were harvested by centrifugation at $5000g$ for 10 min. The cells were then washed twice with normal PBS and finally resuspended to obtain a final cell density of 1.8×10^8 cfu/mL (OD600 0.2) in sterile saline solution (0.85% NaCl).

Antimicrobial assay. The antibacterial activity of the synthesized compounds was assessed using the method of Duraipandiyan³² with slight modifications. Specifically, the bacterial counts of the *S. aureus*, *E. coli* and *K. pneumoniae* isolates were adjusted to 0.5 McFarland turbidity standards using sterile saline solution. The bacterial cultures were poured onto sterile Petri dishes containing Mueller–Hinton agar (Difco). The freshly cultured bacterial isolates were uniformly spread using sterile cotton swabs. A total volume of 100 μL of each stock solution of the synthesized chemicals was dissolved in DMSO and loaded in 6 mm wells. Later, the plates were incubated at $37\text{ }^{\circ}\text{C}$ for 24 h. The results are represented as the size of the inhibition zone around the wells. The negative control wells contained 100 μL of DMSO.

Minimum inhibitory concentrations. The synthesized sulfadimidine derivatives exhibited antibacterial activity at a concentration of 100 mg in DMSO (w/v). Therefore, this concentration was selected to determine their minimum inhibitory concentration (MIC) using the agar well diffusion method³³. The substrates were diluted twice (tenfold dilutions). Approximately 1 mL of each freshly prepared bacterial inoculum was pipetted into sterile Petri dishes, followed by the addition of molten Muller–Hinton agar. Later, five wells were made on each plate, and 100 μL of each diluted substance was transferred to the respective wells in a clockwise manner. The plates were kept in the refrigerator for 20 min and then incubated overnight at $37\text{ }^{\circ}\text{C}$. The MIC was considered the lowest concentration at which the growth of the respective bacteria was inhibited. DMSO was used as the negative control.

Conclusions

This study focused on the synthesis of new nucleoside and Schiff base derivatives and their activities against bacterial strains. Our study concluded that the arabinose nucleoside 3a and acetylated mannose nucleoside 4d derived from sulfadimidine were the most active against *S. aureus* while 4d and the acetylated glucose nucleoside 4c were the most active against *K. pneumoniae*. Finally, 4d, 4c and the acetylated galactose nucleoside 4b were the most active against *E. coli*. The MIC values against *S. aureus* ranged from 50 mg to 3.125 mg, while those against *E. coli* and *K. pneumoniae* ranged from 50 mg to 1562 mg. In vitro, the antibacterial effects were promising. Further research is required to study the in vivo antibacterial effects of these compounds and determine therapeutic doses.

Data availability

Data are available upon request.

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References

1. Ayoup, M. S. *et al.* Synthesis, docking, and evaluation of antimicrobial activity of a new series of acyclo C-nucleosides of 1, 2, 4-triazolo [4, 3-a] quinoxaline derivatives. *J. Heterocyc. Chem.* **53**, 153–163 (2016).
2. Azab, M. E., Youssef, M. M. & El-Bordany, E. A. J. M. Synthesis and antibacterial evaluation of novel heterocyclic compounds containing a sulfonamido moiety. *Molecules* **18**, 832–844 (2013).
3. Di Fiore, A. *et al.* Carbonic anhydrase inhibitors: Crystallographic and solution binding studies for the interaction of a boron-containing aromatic sulfamide with mammalian isoforms I–XV. *Bioorg. Med. Chem. Lett.* **20**, 3601–3605 (2010).
4. Smaine, F.-Z. *et al.* Carbonic anhydrase inhibitors: 2-Substituted-1, 3, 4-thiadiazole-5-sulfamides act as powerful and selective inhibitors of the mitochondrial isozymes VA and VB over the cytosolic and membrane-associated carbonic anhydrases I, II and IV. *Bioorg. Med. Chem. Lett.* **18**, 6332–6335 (2008).
5. Supuran, C. T., Scozzafava, A., Jurca, B. C. & Ilies, M. A. Carbonic anhydrase inhibitors–Part 49: Synthesis of substituted ureido and thioureido derivatives of aromatic/heterocyclic sulfonamides with increased affinities for isozyme I. *Eur. J. Med. Chem.* **33**, 83–93 (1998).
6. Liu, H.-B., Gao, W.-W., Tanganchu, V. K. R., Zhou, C.-H. & Geng, R.-X. Novel aminopyrimidinyl benzimidazoles as potentially antimicrobial agents: Design, synthesis and biological evaluation. *Eur. J. Med. Chem.* **143**, 66–84 (2018).
7. Kalčić, F., Kolman, V., Ajani, H., Zidek, Z. & Janeba, Z. J. C. Polysubstituted pyrimidines as mPGE2 inhibitors: Discovery of potent inhibitors of PGE2 production with a strong anti-inflammatory effects in carrageenan-induced rat paw edema. *ChemMedChem* **15**, 1398–1407 (2020).
8. Yejella, R. P. & Atla, S. R. A study of anti-inflammatory and analgesic activity of new 2, 4, 6-trisubstituted pyrimidines. *Chem. Pharm. Bull.* **59**, 1079–1082 (2011).
9. Dwivedi, A. R. *et al.* Anti-proliferative potential of triphenyl substituted pyrimidines against MDA-MB-231, HCT-116 and HT-29 cancer cell lines. *Bioorg. Med. Chem. Lett.* **30**, 127468 (2020).
10. Mesbah, M. *et al.* Synthesis, characterization, spectroscopic studies and antimicrobial activity of three new Schiff bases derived from Heterocyclic moiety. *J. Mol. Struct.* **1151**, 41–48 (2018).
11. Cawrse, B., Robinson, N., Lee, N., Wilson, G. & Seley-Radtke, K. Structural and biological investigations for a series of N-5 substituted Pyrrolo[3,2-d]pyrimidines as potential anti-cancer therapeutics. *Molecules* **24**, 2656–2668 (2019).

12. Amer, H. H., Ali, O. M. & El-Kafaween, I. K. Synthesis of some new nucleosides derived from 2-mercapto benzimidazole with expected biological activity. *Orient. J. Chem.* **33**, 2303–2310 (2017).
13. Amer, H. H. *et al.* Synthesis of some new 1, 3, 4-oxadiazole derivatives bearing sugars and α -aminophosphonate derived from 4-nitrophenol as anticancer agents. *Natl. J. Physiol. Pharm. Pharmacol.* **8**, 1275–1286 (2018).
14. Siddiqui, H., Haniffa, H. M., Jabeen, A., Rahman, A.-U. & Choudhary, M. I. Sulphamethazine derivatives as immunomodulating agents: New therapeutic strategies for inflammatory diseases. *PLoS One* **13**, e0208933 (2018).
15. Lal, J., Gupta, S. K., Thavaselvam, D. & Agarwal, D. D. Biological activity, design, synthesis and structure activity relationship of some novel derivatives of curcumin containing sulfonamides. *Eur. J. Med. Chem.* **64**, 579–588 (2013).
16. Naaz, F. *et al.* Molecular modeling, synthesis, antibacterial and cytotoxicity evaluation of sulfonamide derivatives of benzimidazole, indazole, benzothiazole and thiazole. *Bioorg. Med. Chem.* **26**, 3414–3428 (2018).
17. Alotaibi, S. H. & Amer, H. H. Synthesis, spectroscopic and molecular docking studies on new Schiff bases, nucleosides and α -aminophosphonate derivatives as antibacterial agents. *Saudi J. Biol. Sci.* **27**, 3481–3488 (2020).
18. Arif, R. *et al.* CU(II) complex: In vitro antibacterial, hemolytic and antioxidant assessment. *J. Mol. Struct.* **1160**, 142–153 (2018).
19. Khan, M. N., Khan, A. M., Ullah, H., Hussain, S. & Khattak, A. K. Synthesis and antibacterial activity of the sulfonamide based schiff base and its transition metal (II) complexes. *Pak. J. Pharm. Sci.* **31**, 103–111 (2028).
20. Nayak, S. G. & Poojary, B. Synthesis of novel Schiff bases containing arylpyrimidines as promising antibacterial agents. *Heliyon* **5**, e02318 (2019).
21. Dale, G. E., Kostrewa, D., Gsell, B., Steiger, M. & D'Arcy, A. Crystal engineering: Deletion mutagenesis of the 24 kDa fragment of the DNA gyrase B subunit from *Staphylococcus aureus*. *Acta Cryst.* **55**, 1626–1629 (1999).
22. Rusakov, A. A., Frisch, M. J. & Scuseria, G. E. Space group symmetry applied to SCF calculations with periodic boundary conditions and Gaussian orbitals. *J. Chem. Phys.* **139**, 114110 (2013).
23. Gupta, M., Sharma, R., Kumar, A. J. O. P. & Medicine, E. Comparative potential of simvastatin, rosuvastatin and fluvastatin against bacterial infection: An in silico and in vitro study. *Orient. Pharm. Exp. Med.* **19**, 259–275 (2019).
24. Ames, B. N., McCann, J. & Yamasaki, E. Methods for mutagenicity test. *Mutat. Res.* **31**, 347–363 (1975).
25. Aldred, K. J., McPherson, S. A., Turnbough, C. L. Jr., Kerns, R. J. & Osheroff, N. Topoisomerase IV-quinolone interactions are mediated through a water-metal ion bridge: mechanistic basis of quinolone resistance. *Nucleic Acids Res.* <https://doi.org/10.1093/nar/gkt124> (2013).
26. Jung, W. K. *et al.* Antibacterial activity and mechanism of action of the silver ion in *Staphylococcus aureus* and *Escherichia coli*. *J. Biomed. Mater. Res.* **74**, 2171–2178 (2008).
27. Feng, Q. L. *et al.* A mechanistic study of the antibacterial effect of silver ions on *Escherichia coli* and *Staphylococcus aureus*. *J. Biomed. Mater. Res.* **52**, 662–668 (2000).
28. Gellibert, F. *et al.* Identification of 1, 5-naphthyridine derivatives as a novel series of potent and selective TGF- β type I receptor inhibitors. *J. Med. Chem.* **47**, 4494–4506 (2004).
29. Podust, L. M., Poulos, T. L. & Waterman, M. R. Crystal structure of cytochrome P450 14 α -sterol demethylase (CYP51) from *Mycobacterium tuberculosis* in complex with azole inhibitors. *Proc. Natl. Acad. Sci.* **98**, 3068–3073 (2001).
30. Goujon, M. *et al.* A new bioinformatics analysis tools framework at EMBL–EBI. *Nucleic Acids Res.* **38**, W695–W699 (2010).
31. Kant, K., Lal, U. R., Kumar, A. & Ghosh, M. A merged molecular docking, ADME-T and dynamics approaches towards the genus of Arisaema as herpes simplex virus type 1 and type 2 inhibitors. *Comput. Biol. Chem.* **78**, 217–226 (2019).
32. Navyashree, V., Kant, K. & Kumar, A. Natural chemical entities from Arisaema Genus might be a promising break-through against Japanese encephalitis virus infection: A molecular docking and dynamics approach. *J. Biomol. Struct. Dyn.* **39**, 1404–1416 (2021).

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Author contributions

All authors contributed equally to the completion of this finished work. H.H.A. as responsible for the design, synthesis, purification of the synthesized nucleosides and Schiff bases derived from sulfadimidine and the docking analysis. E.H.E, S.M.A. and M.Y.H. were responsible for the conception and design of the sensitivity tests and MIC tests. S.H.A. was responsible for the conception of the spectroscopic analyses. Y.S.A. was responsible for the writing of the manuscript and data analysis.

Competing interests

The authors declare no competing interests.

Additional information

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