Naphthoquinones and Their Derivatives

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In the current era, an ever-emerging threat of multidrug-resistant (MDR) pathogens pose serious health challenges to mankind. Researchers are uninterruptedly putting their efforts to design and develop alternative, innovative strategies to tackle the antibiotic resistance displayed by varied pathogens. Among several naturally derived and chemically synthesized compounds, quinones have achieved a distinct position to defeat microbial pathogens.

efflux pumps MDR ESKAPE pathogens naphthoquinones plasmid curing

reactive oxygen species

topoisomerase

1. Introduction

Antibiotics represent world-class, assured molecules that have captured a gigantic share in the global market to combat ever-rising and prevalent infections. Over decades, different types of antibiotics have come into medical practice. Penicillin occupied the European and U.S. markets since its discovery in 1928 by Sir Alexander Fleming, followed by its commercial production in the 1940s ^[1]. Further, the world was gifted with the discovery of another antibiotic, streptomycin, by Albert Schatz, Bugie, and Waksman in 1943. This antibiotic was able to inhibit bacteria, predominantly the organisms responsible for tuberculosis ^[2]. After the success stories of penicillin and streptomycin, a huge number of antibiotics succeeded commercially, such that the projected rise of the global antibiotic market is up to US \$67.25 billion by 2026 ^[3].

The challenges of resistance acquired by microbial pathogens towards existing antibiotics led to the advent of new antimicrobials. Presently, healthcare sectors are severely affected due to eternally escalating antimicrobial resistance (AMR) shown by pathogenic bacteria, parasites, viruses, and fungi. This serious threat associated with public health needs urgent attention and an immediate action plan from government policymakers, as well as private industries. It is essential to note that the challenges associated with AMR have led to a substantial cost escalation for pharmaceutical and health-care products. Patients suffering from microbial infections are ultimately the victims of long-term illness, and are therefore loaded with an additional monetary burden in the form of expensive tests and drugs ^{[4][5]}. There is also an increased morbidity and mortality rate in patients. These multidrug-resistant (MDR) pathogens are therefore referred to as "Super Bugs" ^[6]. MDR bacterial pathogens also comprise the ESKAPE group ^[7]. The abbreviation 'ESKAPE' has been used to designate a group having *Enterococcus faecium, Staphylococcus aureus, Klebsiella pneumoniae, Acinetobacter baumannii, Pseudomonas aeruginosa,* and *Enterobacters*pp. In the year 2008, Rice ^[8] had coined the terminology 'the ESKAPE bugs' to denote the ability of pathogens to escape from the lethal activity of antibiotics and impose severe menace to

human health. These pathogens exhibit resistance to antimicrobial drugs like carbapenems, fluoroquinolones, glycopeptides, β -lactams, β -lactam- β -lactamase inhibitor combinations, lipopeptides, macrolides, tetracyclines, and polymyxins etc. ^[Z]. In the year 2017, the World Health Organization (WHO) published a list of priority pathogens—Priority 1 (Critical), Priority 2 (High) and Priority 3 (Medium)—exhibiting resistance to antimicrobial agents ^[9]. These pathogens can worsen emergency situations and therefore, need urgent surveillance so that new and more effective compounds can be brought through the pipelines. Since 2015, WHO has introduced World Antibiotic Awareness Week (WAAW) to create awareness in the public community, health workforce and among policy-formulating personnel to restrict further emergence of antibiotic resistance and its spread. Since 2020, the Tripartite Executive Committee declared WAAW dates to be 18–24 November ^[10].

To gain AMR, pathogens acquire plasmids (R-drug resistance) or transposons and also possess multidrug efflux pumps (EPs) to drive out the drug molecules from their system [11]. Besides, other strategies are used, like (1) inactivation, alteration, or degradation of the drug by bacterial enzymes, (2) modification of drug binding sites on the bacterial cell, (3) biofilm formation (restricting the entry of the drug), and (4) reduction in intracellular drug accumulation ^[12]. Challenges associated with AMR encouraged researchers to explore a variety of naturally existing and chemically synthesized compounds over the decades. Since ancient times, medicinal plants have been evidenced as a great support to tackle dreaded illnesses ^[13]. Recent advances in the area of phytochemicals and synthesized derivatives have been looked forward to due to their multifunctional therapeutic approaches for dealing with AMR-associated challenges [14][15][16][17]. The unique structural, biological, and functional properties of naphthoguinones (NQs), along with their derivatives, have gained enormous consideration, particularly from a medicinal chemistry perspective [18]. NOs are widely distributed as natural pigments in plants, fungi, and some animals ^[19]. NQ derivatives bearing hydroxyl, methyl, nitrogen, sulfur, halide, phenylamino-phenylthio, or sulphide possess exceptional biological activity. Derivatives bearing hydroxyl groups are seeking more consideration due to their broad-spectrum pharmacological properties ^[20]. NQs possess widespread antibacterial, antiparasitic ^{[21][22]}, antifungal, ^{[23][24]}, antiviral ^[25], and antimalarial properties ^[26]. In the field of cancer biology, NQs are also noticeably identified for their abilities to produce reactive oxygen species (ROS) in cancer cell lines [27][28][29]. NOs are propitious candidates over the other chemotherapeutic drugs used currently. Until today, an ample number of NQ derivatives have been analyzed for their functional potential against various pathogens. This encouraged us to present a comprehensive review of the structural diversity and multifunctional potentiality of NQs in medicinal chemistry. We also shed a light on the current understanding of the mechanistic roles of NQs in combating microbial infections. We also emphasize molecular docking-a powerful approach used in predicting the interactions of NQ molecules in biological systems.

2. Structural Diversity of Naphthoquinones Entities

Structurally, NQs constitute bicyclic structures with two carbonyl groups placed either in positions 1, 4 or 1, 2. The latter case is less frequent (<u>Figure 1</u>A). The chemistry and biological activities of quinones are primarily dependent on the position and chemical nature of the side groups attached (R). Groups like hydrogen, hydroxyl, methyl, nitrogen, sulfur, halide, etc. are attached to the NQ's ring structure. Generally, the presence of a hydroxyl and/or

methyl group in quinone structure is found in nature. These derivatives have been reported for a broad range of applications in pharmacology ^[19]. Recently, NQ derivatives isolated from plant sources including lawsone, juglone, plumbagin, shikonin and lapachol (<u>Table 1</u>) have fascinated researchers due to their (1) abundance, (2) structural diversity, and (3) broad-spectrum therapeutic potential ^[20]. It is imperative to state that 1,4-NQs derivatized at the 2nd and 3rd position with different chemical groups are recurrently reported for their biological properties (<u>Table 1</u>). NQs having oxygen entities at 1, 4 positions in the aromatic ring exhibit promising antimicrobial properties. Sparse literature is evident on 1,2-NQs. Realizing the significance of 1,4-NQ derivatives for biological applications, the major focus of this review remains on 1,4-NQ derivatives.



Figure 1. Diversity of naphthoquinone molecules. (**A**): structures of chemically synthesized naphthoquinones. (**B**): production, purification, characterization, and biological applications of 1,4-naphthoquinone derivatives from plant material.

Table 1. Chemical structures of plant-originated naphthoquinone derivatives.



3.1. 1,4-Naphthoquinone Derivatives of Natural Origin

Naphthoquinones, phenol-rich compounds, are seen abundantly in a variety of plants ^{[30][32]}, animals ^[41], and fungi ^[42]. The functional potency of any antimicrobial agent is uniquely determined from its minimum inhibitory concentration (MIC). The MIC represents the lowest concentration that hampers the visible growth of microbes. Minimum bactericidal concentrations (MBCs) indicate the lowest concentration of a compound that prevents the growth of an organism by killing it. Therefore, MICs and MBCs are imperative for any antimicrobial agent to be considered for pharmacological applications. MICs of plant-derived NQs against pathogenic bacteria are represented in <u>Table 2</u>.

Table 2. Naphthoquinones extracted from plants along with their minimum inhibitory concentration (MIC).

Type of Naphthoquinone	Active against Bacterial Strains	MIC (µg/mL)	Reference
Shikonin from Arnebia euchroma	Staphylococcus aureus	128	[<u>36</u>]
root extract			

Type of Naphthoquinone	Active against Bacterial Strains MIC (µg/mL)		Reference	
	Streptococcus agalactiae	128		
	Escherichia coli	256		
	Salmonella isolates	256		
	Pseudomonas aeruginosa	512		
	Staphylococcus aureus	0.5		
Plumbagin from <i>Plumbago</i>	Escherichia coli	8	[43]	
zeylanica	Klebsiella pneumoniae	2		
	Pseudomonas aeruginosa	8		
Plumbagin from Plumbago zeylanica	S. aureus (MRSA)	4–8	[<u>31]</u>	
	Staphylococcus aureus	400		
Lawsone from <i>Plumbago</i> <i>zeylanica</i> root extract	Salmonella typhi	200		
	Bacillus cereus	200		
	Bacillus subtilis	200		
	Pseudomonas aeruginosa	800		
	Escherichia coli	800	[<u>34]</u>	
	Shigella dysenteriae	400		
	Serratia marcescens	>1600		
	Proteus mirabilis	>1600		
	Klebsiellapneumoniae	800		
	Enterobacter	800		
	Acinetobacter baumannii	800		
Naphthoquinone pigments from	Bacillus megaterium	9.54–54.28	[44]	
Onosina visianii	Montrichardia arborescens	6.82–54.28		
	Micrococcus luteus	9.54–76.20		
	Staphylococcus epidermidis	9.54–54.28		

Type of Naphthoquinone	Active against Bacterial Strains	MIC (µg/mL)	Reference	
	Enterococcus faecalis	6.82-38.10		
	Citrobacter koseri	6.82-76.20		
	Hafnia alvei	6.82–54.28		
	Pseudomonas proteolytica	4.77–76.20		
	Stenotrophomonas maltophilia	4.77-76.20		
	Yersinia intermedia	6.82-76.20		
	-	(values depict MIC ₉₀)		
Shikonin from Lithospermum	Staphylococcus aureus (MRSA)	7.8–31.2	[37]	
erythrorhizon	Staphylococcus aureus (MSSA)	7.8		
	Proteus vulgaris	Strong activity		
Plumbago auriculata root extracts	Klebsiella. pneumoniae	Strong activity	[<u>32]</u>	
	Escherichia coli	Moderate activity		
	Pseudomonas aeruginosa	Less activity		
	Bacillus megaterium	125		
	Bacillus mesentericus	125		
	Bacillus mycoides	125		
	Bacillus subtilis	125		
Naphthoquinone from <i>Alkanna</i> orientalis extracts	Brevibacterium flavum	250		
	Enterococcus hirae	250	[<u>45</u>]	
	Micrococcus luteus	250		
	Staphylococcus aureus	125		
	Staphylococcus citreus	500		
	Staphylococcus roseus	125		
	Escherichia. coli	500		
	Salmonella typhimurium	750		

Type of Naphthoquinone	Active against Bacterial Strains	MIC (µg/mL)	Reference	
Lawsone from <i>Lawsonia inermis</i>	Bacillus subtilis	Activity at 1000 μg/disc		
	Staphylococcus aureus	No activity	[<u>35</u>]	
	Escherichia coli	No activity		
	Pseudomonas aeruginosa	No activity		
<i>Plumbago indica</i> from root extracts	Cutibacterium acnes (formerly Propionibacterium acnes)	12.5	[<u>33]</u>	
	Staphylococcus aureus	12.5		
	Staphylococcus epidermidis	0.78		
Naphthoquinones from Caesalpinia sappan	Clostridium perfringens	Activity observed		
	Bacillus bifidum	Activity observed		
	Bacillus. breve	No significant activity	[<u>30]</u>	
	Lactobacillus casei	Minute activity		
	Escherichia coli	Minute activity	[<u>30</u>]	

5-hydroxy-1,4-naphthoquinone (jugione) (A in <u>lable 1</u>) from *Caesalpinia sappan* heartwood and examined its activity against intestinal cultures, viz. *Bifidobacterium bifidum, Bifidobacterium breve, Clostridium perfringens, Escherichia coli,* and *Lactobacillus casei.* Authors had purified those bioactive molecules using different solvents, like methanol, EA, butanol, chloroform, hexane, and water. The powerful growth-inhibition activity of NQ was obtained from EA, butanol, and methanol extracts against *B. bifidum* and *C. perfringens.* Chloroform fractions resulted in moderate antimicrobial activity, whereas water and hexane fractions were ineffective against both pathogens. The scenario presented here is quite interesting and justifies the selectivity and specificity of bioactive compounds isolated from *C. sappan* heartwood using different solvents. Authors also screened commercially accessible compounds like 5-hydroxy-1,4-NQ, 5-hydroxy-2-methyl-1,4-NQ, 1,4-NQ, and 1,2-NQ against the abovementioned test organisms at concentrations of 0.25, 0.5, 1, 2, and 5 mg/disk. Their analysis recommended the dose-dependent activity of 1,4-NQ, and 1,2-NQ against *C. perfringens*, rather than other isolates used by them. Additionally, the 1,4-NQ derivative showed superior activity against four other organisms (*B. bifidum, B. breve, E. coli,* and *L. casei*) as compared to other derivatives (5-hydroxy-1,4-NQ and 5-hydroxy-2-methyl-1,4-NQ). Results also confirmed that the antibacterial activity of 1,4-NQ was reduced significantly due to the presence of a –CH₃ group at the 2nd and an –OH group at the 5th position.

Plumbagin (B in <u>Table 1</u>) is another widely explored secondary plant metabolite. Plumbagin is of immense interest due to its abilities to (1) generate ROS, (2) inhibit EPs, (3) accumulate drugs, and (4) cure plasmids from bacteria. Thus, due to such exceptional properties, it has been utilized as an effective antibacterial agent ^[47]. Periasamy et al. ^[31] evaluated the antibacterial activity of plumbagin (from *Plumbago zeylanica*) against 100 methicillin-resistant

S. aureus (MRSA) strains, including MDR phenotypes. *P. zeylanica* root extract was obtained using ACN, DMSO, DMF, IPA, methanol and water. All fractions were used to analyze their antibacterial potential where active constituents of ACN and water showed the highest and lowest antibacterial activity respectively. Thus, Periasamy et al. ^[31] demonstrated that NQ constituents obtained using ACN possess promising and consistent antibacterial activity (at MIC of 4–8 µg/mL) against MRSA strains.

Similarly, Adusei et al. [43] reported a bacteriostatic effect of plumbagin (*P. zeylanica*) extract using EA. The authors verified its antibacterial effect at 0.5, 8, 2, and 8 µg/mL against S. aureus, E. coli, K. pneumoniae and P. aeruginosa respectively. This research group also assessed the resistance-modulating potential of antibiotics using plumbagin at its subinhibitory concentration of 4 µg/mL. This assessment demonstrated a range of MICs for ciprofloxacin (0.25 to 2 µg/mL), amoxicillin (0.25 to 256 µg/mL), ampicillin (0.25 to 256 µg/mL), and ketoconazole (no activity) against test pathogens. Their findings presented the successful resistance-modulating potential of three antibiotics, viz. ciprofloxacin, amoxicillin, and ampicillin, with plumbagin against S. aureus and E. coli. The activity of those three antibiotics was enhanced up to 2-fold when they were used in combination with plumbagin. Surprisingly, the enhancement of the 2-fold activity of ciprofloxacin was found against P. aeruginosa, whereas the activity of amoxicillin and ampicillin was reduced to 2-fold against the same isolate. For K. pneumoniae, the activity of amoxicillin and ampicillin was found to be enhanced up to 6-fold. At the same time, the efficiency of ciprofloxacin was reduced by 2-fold for *K. pneumoniae* [43]. The synergistic approaches are unquestionably valuable to prove the potential of plumbagin as a favorable drug. Adusei et al. [43] further reported the considerable antibiofilm activity of plumbagin (32 to 128 µg/mL) against the above-mentioned five pathogens. The antibiofilm activity of plumbagin was noticeably higher against S. aureus as compared to ciprofloxacin. Both plumbagin and ciprofloxacin displayed antibiofilm activity (>50%) against the other three pathogens (S. aureus, E. coli, and K. pneumoniae) at 128 µg/mL concentration. The combination of plumbagin with other drugs undoubtedly impede the growth and infection of biofilm-forming pathogens.

The literature suggests that plumbago species has always been a choice NQ among the ones of natural sources. Patwardhan et al. ^[32] documented the effectiveness of *P. auriculata* root extract against 23 nosocomial pathogenic strains including *P. aeruginosa, P. vulgaris, E. coli,* and *K. pneumoniae*. Solvent extracts of plumbago roots were evaluated at various concentrations from 250 to 4000 μ g/mL. Plumbago extract obtained using ethanol displayed the highest antimicrobial activity as compared with other solvents. Ethanolic based extracts impeded the growth of *P. vulgaris* and *K. pneumoniae*, followed by *E. coli*. It is important to mention that *P. aeruginosa* was found to be the least affected with the same ethanolic plumbago extract.

A report documented by Kaewbumrung and Panichayupakaranant ^[33] explains the stability and the yield of plumbagin from *P. indica* roots. For the extraction procedure, authors used ethanol, EA, DCM, diethyl ether, and isopropanol. Out of those five solvents, ethanol was found to be the most suitable to result extract (11.5% w/w) having a high amount (5.79 mg/g) of the plumbagin. Further purification and elution of pure plumbagin was achieved using a silica gel column with a mixture of hexane: EA (9.2%:0.8% v/v). This step enhanced the total plumbagin content (13.26% w/w). Impurities soluble in hexane and EA were removed to result in a purified form of a derivative. The promising bactericidal activity of plumbagin extract was observed against *S. aureus*, *S.*

epidermidis, and P. acnes. Like EA, petroleum ether has been the solvent of choice to extract NQs from plant sources. Vukic et al. ^[44] used petroleum ether along with EA to isolate NQs from *Onosma visianii.* Thus, the selection of appropriate solvents is important to obtain biologically active components from plant materials. From the above discussed literature, we underscore the pivotal role of solvents to extract NQs from natural sources. Attention to this concept could be helpful to enhance the overall yield of NQs from the desired biological sources. In conclusion, plumbago species are one of the preferred natural sources to isolate NQs. Among various solvents, ethanol is a virtuous choice for the extraction of bioactive compounds, followed by EA and hexane to demonstrate the assured biological activity of NQs.

Like plumbagin, another NQ—namely lawsone (C in <u>Table 1</u>)—has stimulated researchers to explore its medicinal potential. The foremost report on the extraction of lawsone (2-hydroxy-1,4-NQ) from *P. zeylanica* was accomplished by Patwardhan et al. ^[34]. Authors extracted lawsone via the solvents acetone, benzene, chloroform, cyclohexane, diethyl ether, ethanol, methanol, and petroleum ether. Consequently, Patwardhan and collaborators ^[34] focused on the biological activity, purification, and characterization of lawsone extracted using ethanol. The antibacterial potential of ethanolic root extract (ranging from 200 to 800 µg/mL) was effective against three Grampositive cultures (*S. aureus, B. cereus,* and *B. subtilis*) and six Gram-negative cultures (*E. coli, S. typhi, Enterococcus spp., K. pneumoniae, A. baumannii,* and *S. dysenteriae*). However, the same ethanolic extract was apperative against *S. marcescens* and *Proteus mirabilis* at higher concentrations (>1600 µg/mL). Patwardhan et al.

A liposoluble red-colored pigment, shikonin (D in <u>Table 1</u>), is usually extracted from the roots of various plants like *Alkanna tinctorial*, *Lithospermum erythrorhizon*, or *Arnebia decumbens* L. Shikonin is one of the unique bioactive NQs routinely extracted from the roots of *L. erythrorhizon*, and is popular for several biological activities.

Lee et al. [37] isolated shikonin from the roots of *L. erythrorhizon* and examined its antibacterial potency against seven MRSA strains. This study revealed the MIC of shikonin to be 7.8 to 31.2 µg/mL. MRSA strains were found to be more susceptible to shikonin, as compared to ampicillin and oxacillin (antibiotics of the penicillin class, cell wall attacking). Shikonin was evaluated in combination with Tris and Triton X-100 (membrane-permeabilizing agents), sodium azide and N,N'-dicyclohexylcarbodiimide (ATPase inhibitors), and peptidoglycan (derived from S. aureus). The work was supported through experiments like (1) the broth microdilution method (to analyze the susceptibility of bacteria to antibacterial agents), (2) the time-kill test (to determine the bactericidal/bacteriostatic activity of antibacterial agents over time), and (3) transmission electron microscopy (predicting the underlying mechanism of antibacterial agents affecting cell morphology). Studies have discovered the enhanced antibacterial activity of shikonin in the presence of Tris and Triton X-100 and ATPase inhibitors. Lee et al. [37] suggested that shikonin can be proposed as a natural antibiotic and even to realize the underlying mechanism responsible for antimicrobial action. Al-Mussawi [48] had also isolated shikonin from Arnebia decumbens L. and purified it using column chromatography. Researchers proved the antibacterial activity of shikonin against P. aeruginosa, E. coli, S. aureus, and K. pneumonia. Moreover, shikonin has been widely explored for anti-inflammatory, wound-healing, antithrombotic, and anticancer properties ^[49]. Large-scale production of any bioactive molecule is mandatory to utilize them for pharmacological purposes. When NQs are extracted from natural plant sources, the yield and purity cannot be neglected. Recently, Huang et al. [36] developed an ultrasound-assisted extraction (UAE) technique to

isolate shikonin from *A. euchroma*. The response surface methodology (RSM) was used to design an appropriate extraction set-up for the production of shikonin. This experiment set-up can encourage researchers to extract shikonin at ease using an energy-saving approach. Authors magnificently reported around a 1.26% yield of shikonin under the ideal extraction protocol using ultrasound power at 93 W (in 87 min at 39 °C) with a liquid–solid ratio of 11:1. The same studies supplemented the antimicrobial activity of shikonin with clinical isolates (three) along with standard ones (five) at MICs of 128–1024 μ g/mL. Extracts obtained from the medicinal plants find applications in the manufacturing of ointment to treat patients with burn infections. Aljanaby ^[50] isolated an aqueous extract having alkannin esters and shikonin from *A. tinctoria* and demonstrated antibacterial activity against drug-resistant bacterial strains (395) at a 300 mg/mL concentration.

Among different NQs mentioned above, lapachol (E in <u>Table 1</u>) has been regularly reported to have antibacterial and anticancer properties ^[51]. Lapachol has been derivatized to produce a pharmacologically important molecule. Zani et al. ^[38] had isolated lapachol from *Tabebuia ochracea* (Bignoniaceae family), which was further derivatized by Souza et al. ^[39] to produce thiosemicarbazone and semicarbazone. In an antibacterial assay, *E. faecalis* and *S. aureus* were found to be susceptible to thiosemicarbazone lapachol (0.05 µmol/mL) and semicarbazone lapachol (0.10 µmol/mL). Along with Bignoniaceae, many other families like Verbenaceae, Proteaceae, Leguminosae, Sapotaceae, Scrophulariaceae, and Malvaceae show broad scope to extract lapachol.

An interesting report by Balachandran et al. ^[52] documented the isolation of an antibacterially bioactive 1,4 NQ molecule from *Streptomyces* sp., named bluemomycin, using EA. The EA extract displayed potent antibacterial activity against both Gram-negative bacteria as well as Gram-positive bacteria.

In the view of the present scenario, it is also vital to communicate that the microorganisms used for antimicrobial assays are *S. aureus*, *P. aeruginosa*, and *E. coli*, which belong to the ESKAPE category. At lower MICs, most NQs of natural origin have demonstrated better activities against Gram-positive (*Staphylococcus* spp., *Bacillus* spp.) than Gram-negative bacteria. Among Gram-negative bacteria, *Pseudomonas spp.* and *E. coli* have been frequently tested to determine the antibacterial potency of NQs (Figure 2).



Figure 2. Percentage-wise distribution of bacterial species used to evaluate the antibacterial activity of (**A**): naturally derived and (**B**): chemically synthesized 1,4-naphthoquinones.

3.2. Chemically Synthesized 1,4-Naphthoquinone Derivatives

Just like those of a natural origin, chemically derivatized 1,4-NQs are widely investigated for antimicrobial potential. Literature suggests that most of the additions or deletions of the chemical groups have been carried out at 2nd and 3rd positions in NQ moieties. The groups preferred for additions are H, OH, CH₃, Cl, Br, N, and S. Most of the chemical elements chosen for derivatization are placed from 14 to 17 in the periodic table. These atoms are typically reactive non-metals and strong oxidizing agents.

Chemical modifications in the NQ moieties have been facilitated to explore them for a wide range of biological applications. Ravichandiran et al. ^[53] synthesized a series of NQs containing phenylamino-phenylthio moieties and evaluated their antibacterial activity against *S. aureus*, *Listeria monocytogenes*, *E. coli*, *P. aeruginosa*, and *K. pneumonia* with MICs ranging between 15.6 to 500 μ g/mL. Most of the synthesized NQ derivatives were able to inhibit *S. aureus* and *E. coli*. However, NQ derivatives were found to be less effective against the other three

strains. The structure–activity relationship (SAR) suggested that the introduction of the thiophenol group in one of the structures can exhibit moderate antibacterial activity as compared to its ligand. This study revealed that the introduction of a substituted amide group, acid chlorides (aliphatic and aromatic), in one of the compounds shows superior antibacterial properties as compared with its parent compound. Likewise, a compound containing a 3,5-dinitro aryl moiety displayed enhanced antibacterial activity against selected pathogens. Contrary to this, synthesized compounds having a bulky moiety or electron-withdrawing groups (NO₂ and methyl) showed low inhibitory actions against test cultures. This can be further justified by the fact that the bulky groups cannot enter or penetrate easily into the bacterial cell and thus, encounter challenges to fit at the target sites ^[53]. The synthesis of quinones substituted with N-, S-, O- and the evaluation of their antimicrobial and anticancer activities was performed by Kurban et al. ^[18]. For the first time, researchers documented the efficiency of benzo- and NQ derivatives to inhibit an enzyme, catalase (which is responsible for the decomposition of H₂O₂ into H₂O and O₂ to protect cells from oxidative damage).

From the literature, it is noteworthy that the bacterial species of the ESKAPE category are extensively tested to assess the efficacies of chemically synthesized NQ derivatives. NQs can impose greater effects against Grampositive bacteria. However, it is also important to highlight the exceptional cases wherein Gram-negative strains like *K. pneumoniae* and *E. coli* are also susceptible to some of the NQ derivatives ^{[53][54]}. The mechanism of action might be influenced by the constituents of the cell wall. Also, the interaction of the functional groups of NQ derivatives at the target position is another parameter which cannot be neglected. The addition or deletion of a single chemical group (for example OH, Cl, Br, S, N, or O) to or from a specific position in a NQ structure can affect the efficacies of the NQ against targeted pathogens. <u>Table 3</u> presents the antibacterial potential of chemically synthesized NQ derivatives along with their MICs.

Type of Naphthoquinone	Antibacterial Activity Against	MIC (µg/mL)	Reference
Plumbagin	Mycobacterium tuberculosis	4	
	Mycobacterium smegmatis	4	[<u>55</u>]
Lawsone	Mycobacterium tuberculosis	>16	
	Mycobacterium smegmatis	>32	
Lawsone methyl ether	Staphylococcus aureus (MRSA)	62.5–125	[<u>56</u>]

Table 3. Antibacterial potential of chemically synthesized naphthoquinones along with their minimum inhibitory concentration (MIC).

Type of Naphthoquinone	Antibacterial Activity Against	MIC (µg/mL)	Reference
Lawsone derivatives	Staphylococcus aureus (MRSA)	32–128 & >128	[57]
	Staphylococcusaureus (MSSA)	0.6 -128 & >128	
Lapachol	Staphylococcus aureus	1.25 mM	[<u>58]</u>
Lapachol, nor lapachol	Staphylococcus aureus (MRSA)	30-500 & >500	[<u>59</u>]
	Staphylococcus aureus	8–512	
	Bacillus subtilis	32–512	
Imidazole derivatives of 1,4- naphthoquinone	Pseudomonas aeruginosa	8–256	[<u>60</u>]
	Proteus vulgaris	16-256	
	Escherichia coli	32–256	
	Staphylococcus aureus	4–256	
1,4-naphthoquinone derivatives	Bacillus subtilis	128–512	
	Pseudomonas aeruginosa	32–512	[<u>61</u>]
	Proteus vulgaris	128–512	
	Escherichia coli	256–512	
	Staphylococcus aureus	31.25–500	
Phenylamino-phenylthio derivatives of 1,4-naphthoquinone (no. of derivatives synthesized)	Listeria monocytogenes	62.5–500	
	Pseudomonas aeruginosa	62.5–500	[53]
	Escherichia coli	15.6–500	
	Klebsiella pneumoniae	62.5–500	
Sulfide derivatives of 1,4-	Staphylococcus aureus	7.8–250 & >250	[24]
naphthoquinone	Escherichia coli	31.3–250 & >250	
Menadione	Staphylococcus aureus	128	[62]

Type of Naphthoquinone	Antibacterial Activity Against	MIC (µg/mL)	Reference
	Pseudomonas aeruginosa	64	
	Escherichia coli	128	
	Klebsiella pneumoniae	128	
	Staphylococcus aureus	32–256	
	Staphylococcus epidermidis	16–256	
Arylsulfanylmethyl-[1,4]- naphthoquinone derivatives	Staphylococcus simulans	32–256	[<u>63</u>]
	Escherichia coli	256 (less activity)	
	Enterococcus faecalis	32–256	
	Staphylococcus aureus	7.8–500	
	Salmonella bongori	125–500	
	Pseudomonas aeruginosa	125–500	
1,4-naphthoquinone substituted at	Proteus vulgaris	250–500	[<u>64]</u>
	Escherichia coli	125–500	
	Klebsiella pneumoniae	62.5–500	
	Enterococcus faecalis	125–500	
	Enterobacter cloacae	250–500	
	Staphylococcus aureus	1.1–45.4	
Shikonin derivatives	Bacillus subtilis	2–50 & >50	[05]
	Escherichia coli	3.1–50 & >50	[<u>C0]</u>
	Pseudomonas aeruginosa	4–50 & >50	
Naphthoquinone derivatives	Staphylococcus aureus	16–256	[<u>19</u>]
	Pseudomonas aeruginosa	64–256	

Type of Naphthoquinone	Antibacterial Activity Against	MIC (µg/mL)	Reference
	Escherichia coli	64–256	
	Enterococcusfaecalis	64–256	
2-bromo-5-hydroxy-1,4-NQ	Staphylococcus aureus	16	
	Staphylococcus aureus	32–256	
	Staphylococcus epidermidis	128	
	Bacillus cereus	256	
	Salmonella enterica	256	
Juglone	Listeria monocytogenes	256	[<u>66</u>]
	Pseudomonas aeruginosa	128	
	Escherichia coli	128	
	Enterococcus feacalis	256	
	Vibrio alginolyticus	256	
Nitrogen and sulfur derivatives of 1,4-	Bacillus subtilis	1.4–19.3	[<u>67</u>]
naphthoquinone	Proteus vulgaris	2.7–39	
Plumbagin, juglone, lawsone,	Staphylococcusaureus (MRSA)	3.9–125	[<u>68]</u>
menadione and their analogues	Pseudomonas aeruginosa	No significant activity	
	Staphylococcus aureus	3.1	[<u>69</u>]
	Bacillus anthracis	6.25	
Menadione	Streptococcus pyogenes	25	
	Streptococcus agalactiae	6.25	
1,4-naphthoquinone	Staphylococcus aureus	6.25	
	Bacillus anthracis	12.5	

Streptococcus pyogenes50Streptococcus agalactiae12.5Streptococcus agalactiae16-512 & >512Listeria monocytogenes512 & >512Pseudomonas aeruginosa512 & >512Pseudomonas aeruginosa256-512Acinetobacter baumannii512 & >512Plumbagin derivativesMycobacterium smegmatis13.3-30.4Plumbagin derivatives6.25-50 & >50Nitrogen, sulfur groups substitution at 2, 3 positions of 1,4-naphthoquinoneEscherichia coli6.25-50 & >50Klebsiella pneumoniae1.56-50 & >50	[70]
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Nitrogen, sulfur groups substitution at 2, 3 positions of 1,4-naphthoquinone Escherichia coli 6.25–50 & >50 Klebsiella pneumoniae 1.56–50 & >50	
Klebsiella pneumoniae 1.56–50 & >50	[<u>72</u>]
5-Hydroxy-2-methyl-1,4-NQ Clostridium perfringens Antibacterial activity observed	[<u>30]</u>
Lactobacillus casei mentioned specifically for the	
Bifidobacterium bifidum	
1,4-naphthoquinone Bifidobacterium breve	
[74] Clostridium perfringens	io
Escherichia coli	ar
1,2-naphthoquinone Clostridium perfringens	
Bifidobacterium bifidum	اد

propose their extraordinary candidature against pathogens. The percentage-wise distribution of bacterial species evaluated to demonstrate antibacterial potential of 1,4-NQs is depicted in <u>Figure 2</u>. *Staphylococcus* spp. belonging to the ESKAPE and MDR pathogen groups has been reported frequently for its susceptibility towards natural

Type of Naphthoquinone	Antibacterial Activity Against	MIC (µg/mL)	Reference	bacteria,
E. CUII (~11%) and MSEUDUINUNAS SP	Bifidobacterium breve	_useu iui iii viliu assays. iii a	นนแบบ เบ แ	bacteria, ie above-
5-Amino-8-Hydroxy-1,4-NQ	Staphylococcus aureus	50	[<u>73</u>]	ural NQs,
1,4-naphthoquinone	Staphylococcus aureus	10		coli (both aumannii
	Staphylococcus aureus	12.5-50 & >50		<u>e 2</u>). The
	Streptococcus faecalis	12.5–50 & >50		ens.
(L)-a-amino acid methyl ester, heteroalkyl and aryl substituted1,4-	Klebsiella pneumoniae	6.25–50 & >50	[54]	
naphthoquinone derivatives	Escherichia coli	12.5-50 & >50		
	Pseudomonas aeruginosa	50 & >50		ise.

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