Antibacterial Quinolones

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Contributor: Ioana-Andreea Lungu , Octavia-Laura Moldovan , Victoria Biriş , Aura Rusu

The antibacterial quinolones (QNs) and further developed fluoroquinolones (FQNs) represent one of the most important classes of antimicrobial agents from many points of view: activity spectrum, administrations, and tissue distribution, being primarily used to fight bacterial infections.

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1. Antibacterial Quinolones (QNs)

From the discovery of nalidixic acid until the synthesis of the newest fluoroquinolones (FQNs), these synthetic antibacterial agents have proved to be a valuable tool in the fight against infections ^{[1][2][3][4][5]}. Following Lesher's discovery of nalidixic acid in 1962 as an antibacterial agent, the discovery of 6-fluoro analogues has given rise to the FQNs class, one of the most commonly used antibiotic classes ^{[1][6][7][8][9][10]}. Thus, fluorinated compounds opened the road to further generations, improved pharmacokinetic and pharmacodynamic profiles, and provided a broader antibacterial spectrum. As a result, the class is generically called "fluoroquinolones" because all representatives are mostly fluorinated structures ^{[2][11]}.

Third-generation representatives (e.g., levofloxacin) are active against streptococci [12]. Noting that the introduction of a fluorine atom increased the efficiency of flumequine (first generation), the following optimized compounds had a fluorine atom at the C6 position. More efficient FQNs of the second generation (norfloxacin, ciprofloxacin, and ofloxacin) and third generation (levofloxacin) were obtained. Exceptionally, temafloxacin (second generation) presented three fluorine atoms in its chemical structure. Unfortunately, temafloxacin was withdrawn due to cardiotoxicity. Additionally, the producer withdrew clinafloxacin (a third-generation chlorofluoroguinolone) due to the adverse effects of phototoxicity and hypoglycemia ^{[13][14]}. Fourth-generation FONs can have more than one fluorine atom or chlorine atom [15][16]. The advantages of FQNs are essential: slower development of bacterial resistance given the action on DNA gyrase and Topoisomerase IV, and activity against anaerobic bacteria in addition to Gramnegative and Gram-positive strains [12][17]. Over the last two decades, representatives such as besifloxacin, finafloxacin, delafloxacin, and zabofloxacin have received approval for therapy. Presently, the classification by the generation of these new compounds is controversial. They are often reported as belonging to the fourth generation and less frequently to a new generation (the fifth). Even though the mechanism of action for these new compounds does not bring essential new elements, there is the question of shaping the fifth generation, considering the broad spectrum of activity (including resistant bacterial strains) and higher potency. Moreover, these new representatives present a low risk for bacterial resistance development [18].

2. Structural Characterization of Antibacterial QNs

What people now generically term "quinolones" are, in fact, derivatives of either 4-quinolone, 1,8-naphthyridine-4-one, or pyrido-pyrimidine-5-one structures (**Figure 1**) ^{[12][19]}.

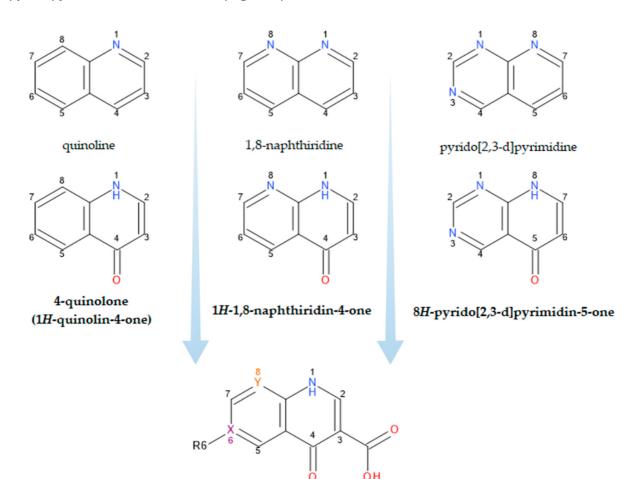


Figure 1. Structural precursors and general structure of QNs (FQNs if R6 = F); X, Y = C or N.

The classification according to the chemical structure of the basic nucleus (**Figure 1**) includes the following groups of compounds:

- Naphthyridine derivatives (nalidixic acid, enoxacin, trovafloxacin, zabofloxacin);
- Quinoline derivatives (cinoxacin);
- Pyrido-pyrimidine derivatives (pyromidic acid, pipemidic acid);
- Quinoline derivatives (norfloxacin, ciprofloxacin, enrofloxacin, moxifloxacin, besifloxacin, delafloxacin, finafloxacin, lascufloxacin, nemonoxacin);
- Compounds with different structures (flumequine, ofloxacin, marbofloxacin, nadifloxacin, and levonadifloxacin).

A contradiction between the generic name "quinolones" and the exact name of the compounds belonging to this class is observed [1][13][18][20][21][22].

An attempt at classification by the number of fluorine atoms in the chemical structure of FQNs includes:

- Non-fluorinated quinolones (nemonoxacin);
- Monofluoroquinolones (ciprofloxacin, enoxacin, marbofloxacin, moxifloxacin, finafloxacin, pradofloxacin, nadifloxacin and levonadifloxacin, zabofloxacin);
- Difluoroquinolones (lomefloxacin, sarafloxacin, sparfloxacin, garenoxacin);
- Trifluoroquinolones (fleroxacin, temafloxacin, trovafloxacin, lascufloxacin);
- Monochloro- and monofluoroquinolones (besifloxacin);
- Monochloro- and difluoroquinolones (sitafloxacin);
- Monochloro- and trifluoroquinolones (delafloxacin) [18][21][22].

The chemical structure of this class of compounds is based on the 1,4-dihydro-pyridine-4-one nucleus, essential for antibacterial activity. The biological activity of a QN is determined by the following important structural elements: (a) the pyridinic ring, unsaturated between the C2 and C3 positions, the presence of a 4-oxo functional group, substitution at the N1 position; and (b) an aromatic B ring. Positions C2, C3, and C4 determine the antibacterial activity (influences the affinity towards bacterial enzymes) ^{[11][20]}. In addition, positions C3 and C4 are involved in metal chelation and other interactions with di- and trivalent cations ^[23]. The newest FQNs' structural characteristics are described in detail by Rusu A. et al. (2021) ^[18].

FQN derivatives are amphoteric compounds whose chemical structure has a carboxyl group at the C3 position (essential for antibacterial activity on the DNA gyrase target). Most commonly, FQNs contain a heterocycle with nitrogen at the C7 position ^[22]. However, various other radicals have been linked to the general structure over time; the chemical structures of relevant representatives are presented in **Figure 2**.

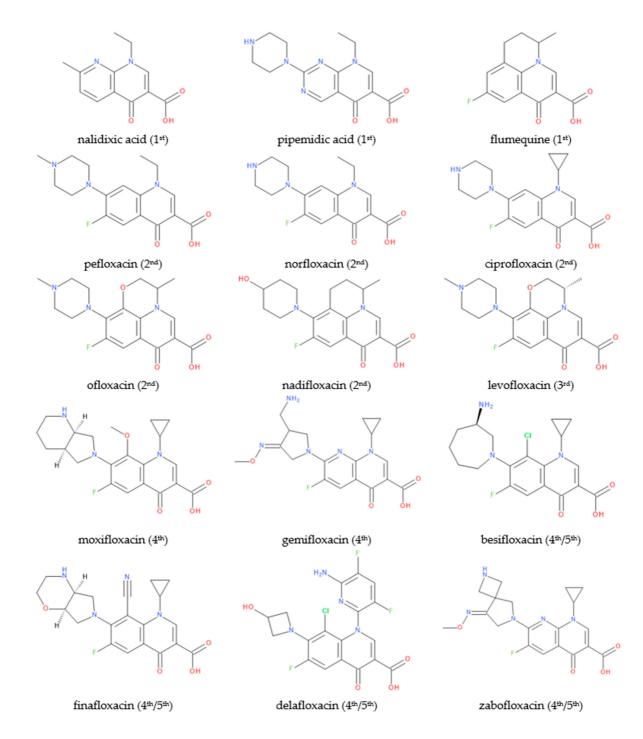


Figure 2. Chemical structures of (F)QNs; the generation is mentioned in parentheses.

There are essential structural elements (C2, C3, and C4 positions) of FQNs closely related to the mechanism of action. In addition, the C1, C5, C7, and C8 positions can serve as targets for various potential substituents (**Table 1**) ^[2].

Table 1. Essential structure-activity relationship aspects in the antibacterial QNs class.

Position on the Chemical Structure	Requirements and Possible Implications	References	
2	Optimal is a hydrogen moiety; larger moieties may hinder the C3 and C4 positions.	[2]	
3	A carboxyl group is required (essential for interacting with the DNA bases and DNA gyrase).	[2][24][25][26]	
4	Oxo-(keto) moiety is required; essential for interacting with the DNA bases and DNA gyrase.	[<u>27</u>]	
6	Small moiety is required (optimal—fluorine); fluorine increases the potency by between 5- and 100-fold compared to any other potential halogen moiety.	[2]	
1	It is involved in the pharmacokinetic properties and overall potency. A cyclopropyl moiety confers activity against Gram-negative bacteria. A 2,4-difluorophenyl substituent determines less potency but heightens activity against anaerobes (e.g., temafloxacin; it was withdrawn shortly after approval due to severe adverse reactions).	[<u>2][28][29]</u>	
5	Specific radicals substituted at this position (- NH_2 , - CH_3) may increase activity against Gram-positive bacteria.	[<u>2][28]</u>	
7	It is involved in pharmacokinetic properties and the spectrum of activity. A five- or six-membered nitrogen heterocycle at this position improves the activity and pharmacokinetic profile. For example, amino pyrrolidine or an alkyl moiety determines enhanced activity against Gram-positive bacteria. On the other hand, piperazine determines better activity against Gram-negative bacteria.	[<u>2][28]</u>	
8	It is involved in the pharmacokinetic properties and activity against anaerobic bacteria.	[2]	

Over time, researchers have explored numerous possibilities for modifying the basic structure of QNs. Researchers have tested various substitutions to obtain a molecule with potent antibacterial activity, a broad spectrum of activity, and superior pharmacokinetic properties. Numerous new compounds have been synthesized and tested for their biological activity, with the ultimate goal being the perfect "X-floxacin".

3. Physicochemical Properties of FQNs

The antibacterial QNs class contains crystalline substances or crystalline powders, generally white or yellowish-white in color, tasteless, with a slightly bitter or bitter taste, odorless, insoluble in water and slightly soluble in common organic solvents, and soluble in dimethyl sulfoxide. Their solubility increases in acidic and basic environments (QNs form water-soluble salts). Many QNs are conditioned as salts: hydrochlorides (ciprofloxacin, moxifloxacin, besifloxacin, lascufloxacin), malate hemihydrates (nemonoxacin), methanesulfonates (pefloxacin), and toluenesulfonates (tosufloxacin) ^{[18][30][31][32][33][34]}.

The low water solubility of FQNs (except their salts) is due to the crystalline structure with condensed aromatic nuclei. FQNs present high melting points (greater than 200 °C) due to a stable crystalline structure. Their water solubility depends on pH (as amphoteric compounds): in acidic or basic environments, they dissolve, forming salts; in the range of pH 6–8, water solubility is low ^{[11][21]}. Increasing the solubility of FQNs is very important for their parenteral administration. For this purpose, new strategies were developed, such as obtaining prodrug formulations ^{[35][36][37]}.

Some FQNs form hydrates depending on the temperature and relative humidity. Lambert A. et al. (2007) confirmed the predominance of the zwitterion form of levofloxacin in water and its lipophilic character, providing models of hydrated molecules with five water molecules ^[38].

FQNs present one or two chiral centers in the chemical structure and are available as racemates (ofloxacin, gemifloxacin, nadifloxacin), enantiomers (levofloxacin, moxifloxacin), or diastereoisomers (besifloxacin) ^{[11][18][39]}. It is known that spatial conformation influences the physical properties of molecules. Thus, an increase in the water solubility of the enantiomers was noted, with the racemates having reduced solubility (e.g., levofloxacin versus ofloxacin) ^{[40][41]}.

FQNs derivatives are amphoteric compounds with four different chemical species in solution (cationic, anionic, zwitterionic, and neutral). These distinct molecular species have different properties in terms of solubility and lipophilicity. The ionized forms are much more soluble in water, and the neutral forms are more lipophilic. Two pKa values are most frequently reported. A value is conferred by the three-carboxyl group, which gives the molecule an acidic character. The nitrogen atoms confer another value from the heterocyclic substituents in C7 position (piperazine, pyrrolidine, etc.). Rusu et al. (2011) established three protonation centers by 1H NMR-pH titrations (the carboxylate moiety, the N-1' and N'-4-piperazine nitrogens) for six FQNs. Additionally, macro- and microprotonation schemes and species-specific diagrams have been outlined ^{[42][43]}. Knowledge of the intimate protonation processes of FQNs is crucial in facilitating diffusion through membranes under particular conditions, increasing the distribution and accumulation in different target tissues, binding to structural components of membranes or specific intracellular ligands, and interpreting chemical structure–biological activity relationships ^{[42][44]}.

The fluorine atom is often identified in the lead optimization studies as a strategy to increase the lipophilicity (log P) of the compound, block the metabolism, or optimize the pharmacokinetic properties ^{[45][46][47][48]}. The introduction of the fluorine atom at the C6 position led to an increased antimicrobial activity versus non-fluorinated QNs; it increased the degree of penetration into the bacterial cell and the activity against Gram-negative bacteria ^[49]. Several FQNs from the second generation were found to be lipophilic compounds (e.g., pefloxacin), intermediate lipophilic compounds (e.g., norfloxacin) based on the true partition coefficients ^[50].

The physicochemical parameters of lomefloxacin, levofloxacin, and moxifloxacin, as potential bioavailability descriptors, were determined (in vitro and in silico) in a study performed by Kłosińska-Szmurło, E. et al. (2014) ^[51]. These published data concerning the lipophilic character are in agreement with the following order enrofloxacin > levofloxacin > ciprofloxacin > norfloxacin, established later by Blokhina S.V. et al. (2016) ^[41]. Some new FQNs are

more lipophilic than others based on the experimental or calculated log P values. An ordering of the new compounds according to their lipophilic character is as follows: nadifloxacin > lascufloxacin > delafloxacin > besifloxacin > nemonoxacin > finafloxacin > zabofloxacin ^[18]. Currently, numerous structural modifications of FQNs are being studied to increase the lipophilicity of the molecule (e.g., new derivatives, FQNs hybrids, prodrugs) ^[52].

Another characteristic property of FQNs is the ability to be complexed by metal ions. Due to the carboxylic group at the C3 position, the piperazinyl ring (or another N-heterocycle) at the C7 position, and the carbonyl oxygen atom at the C4 position, FQNs could form metal complexes. FQNs act as bidentate, unidentate, or bridging ligand. The stoichiometry of the chelated forms depends on several factors: the relative concentrations of the chelating agents (FQNs), the metal ions, the valence of the metal ion, and the pH value. FQNs can form 1:1, 2:1, or 3:1 chelates with metal ions. Over time, research on metal ion complexation has focused on numerous chemical elements ^{[23][32][53]}.

4. Mechanism of Action of Antibacterial FQNs

Antibacterial FQNs have a bactericidal effect involving a particular mechanism of action, namely the inhibition of DNA replication and transcription ^{[23][54]}. This mechanism is carried out by interacting with complexes of DNA and the enzymes DNA gyrase (a type II Topoisomerase) and Topoisomerase IV. These are two essential enzymes involved in DNA cleavage and ligation reactions ^{[17][26][55]}. The two enzymes have a similar tetrameric structure (A2B2). Gyrase's subunits are GyrA and GyrB while Topoisomerase IV's subunits are ParC and ParE. The GyrA subunit of gyrase contains the active site with tyrosine residuals. In contrast, the subunit GyrB contains the TOPRIM domain where the divalent ions bind, making the processes of DNA cleavage and ligation reaction. The ParC subunits are responsible for DNA binding and the cleavage and re-ligation reaction. The ParE subunits are responsible for ATP binding and hydrolysis. These two enzymes ensure the double helix passes through a temporary double-stranded break to make DNA replication possible. The genomic integrity through this process is maintained by connecting the enzymes (the active site contains tyrosine residues) to the DNA strands through covalent bonds, forming complexes known as "cleavage complexes" ^{[23][55][56]}.

Although similar in structure and mechanics, the two enzymes' particular function in DNA replication differs ^{[26][57][58]} ^[59]. DNA gyrase is a unique enzyme in bacterial cells but not in the higher eukaryotes. It is the only Topoisomerase that can introduce negative supercoils into DNA using energy from ATP hydrolysis. It is primarily responsible for releasing the tension that accumulates in front of the replication forks ^{[24][26][55][60]}. Topoisomerase IV plays a role in relaxing positive supercoils in the DNA. This enzyme removes knots forming in the chromosome and decatenates the two chromosomes that result from replication ^{[23][55]}. Even though these DNA gyrase and Topoisomerase IV are essential for cell survival, they have the potential to fragment the genome, and this is the characteristic that QNs use to destroy the bacterial cell ^[26].

FQNs exert their action by binding to one or both target enzymes and the DNA, stabilizing the cleavage complexes ^[25]. Most commonly, DNA gyrase represents the main target of QN in Gram-negative bacteria. In contrast, Topoisomerase IV represents the main target of QN in Gram-positive bacteria, with DNA gyrase being the secondary target in this case ^[61]. The exact binding of the (F)QNs to the target enzymes is partially elucidated. X-ray

crystallography facilitated the discovery of the localization of the amino acids involved in the F(QN)–target interaction. These are located near the active-site tyrosine, involved in DNA breakage ^[25]. More detailed studies have been conducted to investigate the active site of the Topoisomerase IV–DNA cleavage complex for *Streptococcus pneumoniae* with new 7,8-bridged FQNs. The new 7,8-bridge compounds have proven antibacterial activity and offer an alternative to design new FQNs substituted on the C1, C7, and C8 positions to increase activity against resistant bacteria ^[62].

The drug is intercalated between the DNA substrate and the enzyme. Interestingly, FQNs have a greater affinity for enzyme–DNA complexes than enzymes. The structural model of the drug–enzyme–DNA complexes has been discovered using X-ray crystallography. An essential part of connecting the FQN and the enzyme is the presence of a noncatalytic magnesium ion (Mg²⁺) coordinated with four water molecules and the C3/C4 FQNs' carbonylic oxygens. Two water molecules coordinated with Mg²⁺ interact with the residues in the GyrA subunit of the enzyme. The magnesium ion is essential for forming a bridge between the enzyme and the drug. Interactions between the GyrB subunit and the C7 substituent of the FQN are also crucial for binding. Once the big complexes are formed (drug–Mg²⁺–enzyme–DNA), the enzymes become toxic to the cell, with the replication and transcription processes being blocked ^{[24][26][27]}.

However, the FQNs or the complexes they form do not kill the bacterial cell alone, especially since the genesis of complexes is reversible. There are a few critical factors that lead to either a slow death or to a rapid one.

- The slow death is caused by the unprocessed complexes that block replication and transcription;
- The immediate death occurs when the complexes are processed (by dissociation of the gyrase subunits or by removal of the gyrase from the DNA). In this case, the cell is killed due to the fragmentation of the chromosome, which results when the broken DNA is not repaired.

Additionally, more DNA breaks are caused by an accumulation of reactive oxygen species induced by damaged DNA and possibly by the cleavage complexes ^{[24][26][27]}.

5. Indications, Spectrum of Activity, and Pharmacokinetics Data

The representatives from the first generations were used mainly in treating urinary tract infections caused by Gramnegative bacteria ^{[2][19]}. The second-generation representatives have expanded this utility spectrum to include the respiratory, urogenital and gastric tract, bone and joint infections, septicemia and surgical infections, some *Staphylococcus* spp., and venereal diseases ^{[11][63][64]}. Moreover, the representatives from the second generation have longer half-lives, less protein binding, and improved activity on Gram-negative bacteria ^[19]. Thirdgeneration FQNs have markedly enhanced activity. Finally, the fourth generation is indicated in treating communityacquired pneumonia, skin and skin structure infections, bacterial conjunctivitis, and otitis externa ^{[3][13][16][65][66][67][68]} [^{69][70][71]}. The usual doses and indications of the most used antibacterial QNs are summarized in **Table 2**. **Table 2.** Doses and therapeutic indications (US Food and Drug Administration (FDA) and European Medicines

 Agency (EMA) approved).

Compounds (Generation)	Usual Doses	Indications and Administration	References
Nalidixic acid (1st)	4 g daily (every 6 h); 7 to 14 days in acute infections, reducing after that to half this dose in chronic infections.	Uncomplicated urinary tract infections; Oral administration.	[<u>11][13][63]</u> [<u>72</u>]
Norfloxacin (2nd)	400 mg twice a day (every 12 h); 3–7– 21–28 days depending on the severity and nature of the infection.	Uncomplicated and complicated urinary tract infections; Acute or chronic prostatitis; Uncomplicated gonorrhea; Oral administration.	[<u>11][14][63]</u> [73][74]
Ciprofloxacin (2nd)	250–500 mg (every 12 h); 7 to 14 days or more, depending on the severity and nature of the infection.	Uncomplicated and complicated urinary tract infections, pyelonephritis, sexually transmitted diseases, prostatitis, skin and tissue infections; Oral (as the hydrochloride or base) and parenteral administration (lactate), eye drops, eye ointment, or ear drops (as the hydrochloride).	[<u>11][63][75]</u>
Ofloxacin (2nd)	200–400 mg twice a day (every 12 h); 3 days to 6 weeks, depending on the severity and nature of the infection.	Similar to ciprofloxacin. In addition, <i>Chlamydia</i> or <i>Chlamydophila</i> infections include nongonococcal urethritis and mycobacterial infections (leprosy and tuberculosis); Oral (as a base) and parenteral administration (as a hydrochloride salt).	[<u>11][63]</u>
Pefloxacin (2nd)	400 mg twice daily (every 12 h); similar to norfloxacin.	Uncomplicated gonococcal urethritis in males, Gram- negative bacterial infections in the gastrointestinal system and the genitourinary tract; Oral and parenteral administration (as a mesylate salt).	[<u>11][14][73]</u> [<u>76]</u>
Nadifloxacin (<i>topical use</i>) (2nd)	Twice a day as cream or ointment (1%).	Acne vulgaris and other skin infections; Topical use.	[<u>39][63][64]</u> [<u>77]</u>
Levofloxacin (3rd)	250–500 mg (once or twice daily); 7 to 14 days, depending on the severity and nature of the infection.	Acute and chronic bronchitis, exacerbated forms, acquired pneumonia (nosocomial), and other susceptible infections, including tuberculosis; Oral and parenteral administration (as a hemihydrate); Ophthalmic use (0.5% ophthalmic solution).	[<u>11][63][65]</u> [<u>78][79]</u>
Gatifloxacin (ophthalmic	Day 1:1 drop every 2 h in the affected	Bacterial conjunctivitis, ophthalmic use (0.3% or 0.5% ophthalmic solution).	[80][81][82]

Compounds (Generation)	Usual Doses	Indications and Administration	References			
use) (3rd)	eye(s) while awake, up to 8 times Day 2 to 7:1 drop twice to 4 times daily in the affected eye(s) while awake.					
Moxifloxacin (4th)	Oral: 400 mg once a day; 5–10 days depending on the severity and nature of the infection; Ophthalmic administration: one drop in the affected eye 3 times daily for 7 days.	Sexually transmitted diseases, prostatitis, skin and tissue infections, acute and chronic bronchitis, exacerbated forms, acquired pneumonia (nosocomial), intra-abdominal infections, gynecological infections, bacterial conjunctivitis; Oral, parenteral, and ophthalmic administration (0.5%) as a hydrochloride salt.	[<u>66][68][83]</u> [<u>84][85]</u>			
Delafloxacin (4th)	Intravenous: 300 mg over 60 min, every 12 h; Oral: 450 mg every 12 h; 5 to 14 days.	Bacterial skin and skin structure infections; Oral and intravenous administration.	[<u>16][86][87]</u>			
Besifloxacin (topical, ophthalmic use) (4th)	Ophthalmic administration: 1 drop in the affected eye 3 times daily, 4 to 12 h apart for 7 days.	Bacterial conjunctivitis; Ophthalmic suspension (0.6%).	[<u>15][88][89]</u>			
Finafloxacin (topical, ophthalmic use) (4th)	Optic administration: 4 drops in the affected ear(s) twice daily for 7 days.	Acute otitis externa; Optic suspension (0.3%).	[<u>70][90][91]</u>			
 Balofloxacin, third generation—approved in South Korea (2001) ^[92]; 						

- Prulifloxacin, fourth generation—approved in Japan (2002) [93];
- Sitafloxacin, fourth generation—approved in Japan (2008) [94], Thailand (2012) [95];
- Nemonoxacin, fourth generation—approved in Taiwan (2014) [96];
- Zabofloxacin, fourth generation—approved in South Korea (2015) [97].

The pharmacokinetic properties of the several QNs are listed in **Table 3**. Some representatives have the potential to be incorporated into dual antibiotic hybrids.

Table 3. Pharmacokinetic data of some representative QNs.

FQNs	Single Dose p.o. ¹ (g)	Plasmatic Concentration (µg/mL)	Half- Life (Hours)	Binding to Plasma Proteins (%)	Elimination Route	References
Avarofloxacin	0.25	2	14	65	renal	[<u>98]</u>
Ciprofloxacin	0.2	0.8	4–6	20–50	renal, hepatic, feces	[6][11][99][100]
Delafloxacin	0.45	5.80-7.17	4.2– 14.9	84	renal	[<u>86][87][101]</u>
Enoxacin *	0.20	1.0	5	40–60	renal, hepatic	[<u>6][11][18][100]</u>
Fleroxacin *	0.4	5.0	10–12	23	renal, hepatic	[<u>18][102]</u>
Gatifloxacin *	0.20	2.0	7.8	20	renal	[6][18][66][100]
Gemifloxacin *	0.32	1.6	6.9	60–70	renal and others	[<u>6][18][66]</u>
Grepafloxacin *	0.40	0.93	12	50	hepatic, renal	[<u>6][18]</u>
Lomefloxacin *	0.2	0.7	3–4	10	renal	[11][18][100]
Levofloxacin	0.50	6.2-8.7	6–7	24–40	renal	[<u>6][11][99]</u>
(Ala)Levonadifloxacin	1	16.5	4.5	85	-	[<u>103</u>]
Moxifloxacin	0.40	4.5	12	30–50	hepatic, renal	[<u>6][66][100]</u>
Nalidixic acid	1.00	20–40	6–7	93–97	renal	[<u>6][11][100</u>]
Nemonoxacin	0.5	7.02	15	16	renal	[<u>104</u>]
Norfloxacin	0.40	1.5–2	4—8	15	renal, hepatic, feces	[<u>11][73][100</u>]
Ofloxacin	0.20	1.5	4.5–9	32–40	renal	[<u>6][11][100]</u>
Pefloxacin	0.40	3.9–5.8	8–13	20–30	hepatic, renal, feces	[<u>105]</u>
Sparfloxacin *	0.40	1.1–1.3	20	40–50	renal, hepatic	[6][11][18][99] [100]
Temafloxacin *	0.60	2.43	8	25	hepatic,	[<u>6][18][106]</u>

FQNs	Single Dose p.o. ¹ (g)	Plasmatic Concentration (µg/mL)	Half- Life (Hours)	Binding to Plasma Proteins (%)	Elimination Route	References	
					renal		
Trovafloxacin *	0.10	1.0	9.1	76–85	hepatic	[6][18][100]	[:] QNs
Zabofloxacin	0.4	2.0	8.24– 8.32	NA ²	NA ²	[<u>97][101][104]</u> [<u>107</u>]	ne of the

However, some advantages counterbalance the potential side-effects that might have otherwise driven scientists to search for other options in addition to FQNs. This class of antibacterial agents has advantages such as the mechanism of action that confers a bactericidal effect, their effectiveness and potency, and slower development of antimicrobial resistance, especially for the newer representatives, because of their dual activity against both target enzymes ^{[7][20][118]}.

In addition to the advantages of antibacterial activity, FQNs also have an advantage from a chemical point of view. Their structures are relatively easy to synthesize, thus offering the possibility of developing numerous potential derivatives with various advantageous particularities ^{[2][6][7][65]}. Furthermore, FQNs have excellent complexing properties with metal ions due to their chemical structure and can form combinations with other active molecules ^[23]. ^[119]. The advantages mentioned above most likely counterbalance any drawbacks of the potential side effects. For this reason, FQNs have been the target of numerous attempts at hybridization and the development of new antibacterial agents ^[52].

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