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Topological Index Nclass as a factor determining the antibacterial activity of quinolones against Escherichia coli.

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ABSTRACT.

Background: Due to antibiotic resistance and the lack of investment in antimicrobial R&D, QSAR methods appear as an ideal approach for the discovery of new antibiotics. **Result/Methodology:** Molecular topology and LDA were used to construct a model to predict activity against *Escherichia coli*. This model establishes new SARs, of which, molecular size and complexity (*Nclass*), stand out for their discriminant power. This model was used for the virtual screening of the Index Merck database, with results showing a high success rate as well as a moderate restriction. **Conclusions:** The model efficiently finds new active compounds. The topological index *Nclass* can act as a filter in other QSAR models predicting activity against *E. coli*.

KEY WORDS. Quinolones, Antibacterial activity, QSAR, Molecular Topology, *Escherichia* coli

INTRODUCTION.

In the early 60s, a group of experts in infectious diseases announced the need to accelerate the discovery of new antibacterial drugs due to the increasing resistance of some bacteria, such as pneumococci or staphylococci, against traditional antibiotics [1]. Since then, the scientific community has witnessed how the number of antibiotic resistant bacteria has increased exponentially during the last few decades. In fact, according to WHO (World Health Organization) estimates, if resistances continue to increase at the present rate, infectious diseases will become the first cause of death worldwide with more than 10,000,000 deaths, exceeding those caused by cancer, diabetes and cardiovascular diseases [2].

Escherichia coli is one of the bacterial species against which the development of new antibiotics is crucial. In fact, WHO classifies it as a Priority 1 (Critical) pathogen in their Priority Pathogens List of R&D of New Antibiotics [3]. E. coli is a usual commensal in the gastrointestinal tract, which coexists symbiotically with its host [4]. However, certain strains have acquired virulence factors that increase their ability to adapt to new niches, allowing them to cause a broad spectrum of diseases [5]. Generally, these diseases are grouped in three types: enteric, urinary tract infection or septicemia. The

development of resistances in this species is especially serious, having detected *E. coli* strains resistant to all the antibiotics currently available, including colistin [6].

This increase in resistant strains necessitates the research of new antimicrobial agents in order to increase the therapeutic arsenal. However, WHO has emphasized repeatedly the insufficient investment currently dedicated to the development of antimicrobial drugs [7]. The fact that antibiotics are used for acute treatments of short duration and that, due to resistances, they lose efficacy a few years after they are introduced in the market, has led the pharmaceutical industry to abandon all efforts of developing new antimicrobial compounds. As a result, the discovery of new antibacterial therapies relies on research centers and universities [8]. This is where QSAR (Quantitative Structure-Activity Relationship) methods play a key role, seeing as they provide useful information for the rational design of new molecules at a minimum cost [9]. With these methods, one can predict the pharmacological activity of a compound without synthesizing it. Therefore, Computational Chemistry and Virtual Combinatorial Chemistry have experienced a great surge for the last few years [10, 11]. Since virtual libraries are usually big, filters are usually applied to discard molecules with low probability of becoming drugs. Lipinski was the first to apply these filters by describing the "rule of 5" in 1997 [12].

QSAR methods have been used to study antibacterial activity against *E. coli* in different ways. Some studies use known antibiotic families to obtain models for a better understanding of the structural features that most contribute to antibacterial activity against this microorganism [13,14]. For example, Cooper et al. developed a model to identify physicochemical descriptors that best classify the potencies of ß-lactams and fluoroquinolones against *E. coli* [13]. Similarly, Kawczack et al. revealed influential features of fluoroquinolones for their activating against *E. coli* by performing PCA and MLR on a set of these compounds [14]. On the other hand, the most common application of QSAR models against *E. coli* is for the development of new antibacterial compounds [15-17]. Along these lines, Merzoug et al. applied virtual screening to a library of peptide deformylase inhibitors and found three peptides with significant antibacterial activity and a novel mechanism of action [15]. Other studies use QSAR models to obtain new compounds from families with known antibiotic

activity, such as cinnamaldehyde-amino acid Schiff bases [16] and N-substituted rhodanine synthons [17].

The aim of this paper is to develop a prediction model to find new antibacterial molecules and provide additional information regarding the possible structure-activity relationships of quinolones.

MATERIALS AND METHODS.

Compound selection

Quinolones constitute a well-known therapeutic group commonly used for the treatment of different infections, both in hospital and community settings. This group is ideal for the development of a QSAR model because there is a great amount of information published regarding active and inactive molecules within this family, which is often the most difficult to find when conducting QSAR studies. This, along with the fact that quinolones are generally safe and well tolerated drugs, is why they are ideal candidates for the development of an antibacterial activity prediction model.

A group of 43 compounds with proven activity against *E. coli* was selected with a literature search in ISI Web of Science, Medline and SciFinder. Furthermore, a group of 39 inactive compounds was also selected. **All 82 compounds had quinolone structure** (the structure of all compounds as well as references regarding their activity can be found in the Supporting Information file).

A compound was classified as active when it had a Minimum Inhibitory Concentration (MIC) against $E.\ coli \le 1 \text{mg/L}$. On the other hand, a compound was classified as inactive when it had a MIC $\ge 16 \text{mg/L}$. Compounds with intermediate activities (MIC > 1 mg/L and MIC < 16 mg/L) were not included in the compound database. Regarding stereoisomers, only those compounds for which all stereoisomers had the same activity were included in the database seeing as the descriptors used do not include 3D parameters.

Molecular connectivity and topological descriptors

The topological descriptors used are described in the Supporting Information, as well as their definitions and references. A total of 148 descriptors were used to characterize each compound. They were calculated using the hydrogen-depleted

graph to obtain the adjacency topological matrix with MOLCONN-Z [18] and DESMOL13 [19] software.

Model construction

i. Linear Discriminant Analysis

Stepwise Linear Discriminant Analysis (LDA) was used in order to calculate 7 Discriminant Functions (DFs) using different combinations of the topological indexes (i.e. electrotopological, electrotopological and charge, connectivity and charge, etc.). The software used for the LDA was the BMDP 7M Biomedical package [20], which randomly selects molecules for the training and test groups. The topological indexes used to calculate the linear classification functions were chosen in a stepwise manner using the Fisher-Snedecor parameter, F, which determines the relationship between the variance explained by the equation and the residual variance [21].

The discriminant power of the resulting equations was assessed by the percentage of correct classifications for each group. The classification criterion is the minimal Mahalanobis distance, which is the measure of the separation between the active and inactive groups, this allowing to have an idea of the ability of the discriminatory power of the final LDA model. The quality of the discriminant function was assessed using the Wilk's U-statistical parameter, λ . LDA was then applied to the training groups to obtain discriminant functions linking topological indexes and antibacterial activity.

Different combinations of the 7 DFs obtained were analyzed in order to obtain a prediction model with one DF or the combination of several DFs with the best prediction capability, especially of inactive compounds.

ii. Pharmacological Activity Distribution Diagrams

The pharmacological distribution diagrams (PDD) of the selected DFs were built in order to determine the intervals of the DF with the maximum expectancy of finding an active compound. PDDs are a type of histogram plot where connectivity functions are represented with the expectancies on the ordinate axis. They allow the visualization of regions of minimum overlap, allowing the selection of the intervals in which the probability of finding active molecules is maximum [21].

Model validation (virtual screening)

A virtual library was constructed based on the Index Merck database, which is made up of more than 10000 compounds including drugs, biological products and chemical reagents. All inorganic compounds and polymers were excluded. In the case of stereoisomers, a single graph was drawn to represent all the different options. When an organic cation corresponded to a conjugate acid, the corresponding base was drawn. Anionic organic compounds were drawn as free acids.

Finally, the chemical graphs of 6384 compounds were drawn using ChemBioDraw Ultra 12.0 and 148 topological indices were calculated for each compound using DESMOL13 and MOLCONN-Z 3.0.

RESULTS/DISCUSSION.

Prediction Model

To obtain the DFs, a training group with 31 active and 32 inactive compounds and a test group with 12 active and 7 inactive compounds were formed. The test group allows the quality evaluation of the selected DFs. The selected discriminant functions, DF1 and DF2, along with their corresponding statistical parameters are depicted in equations 1 and 2 (the values for the indexes, DFs and classification probability along with the activity classification of the compounds used for the construction of the model can be found in the Supporting Information):

148 **DF1** = -137.77814 + 13.66578
$$^{5}\chi_{p}$$
 - 9.46572 $^{0}\chi_{v}$ + 14.55307 $^{SH}_{tCH}$ - 3.10866 $^{S}_{>CH}$ +

149 0.94222**S_{CI}** - 6.32603**4G^V**

150 N = 82
$$\lambda$$
 = 0.1230521 F = 66.515 (Eq. 1)

151 **DF2** =
$$-108.72509 - 55.72224\chi_{c}v - 0.91239N_{class} + 156.844^{0}C - 45.40381^{2}C$$

152 N = 82
$$\lambda$$
 = 0.2786756 F = 37.532 (Eq. 2)

Both equations have low values for λ , which highlights that the linear dependence between independent variables is not relevant. Furthermore, F presents

high values, indicating that the selected independent variables contribute largely to the classification of both groups.

In order for a compound to be considered active against *E. coli*, it had to have a DF1 value between 0 and 32 and a DF2 value between 0 and 12. On the other hand, to avoid false actives, compounds with values for DF1 > 32 and/or DF2 > 12 are considered unclassifiable These intervals were determined using the corresponding PDDs (Figure 1).

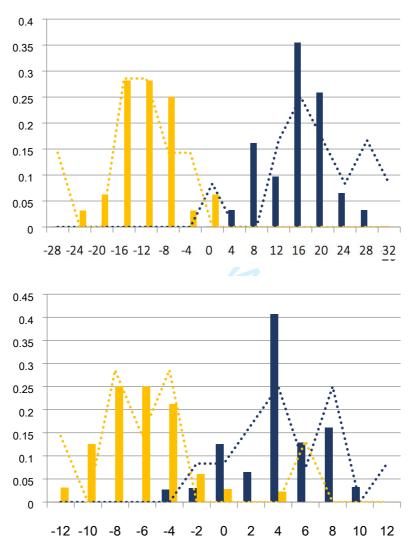


Figure 1. PDDs for DF1 (top) and DF2 (bottom) used to determine activity intervals. Yellow bars: active training, yellow dotted lines: active test, blue bars: inactive training, blue dotted lines: inactive test.

Compounds that only satisfied one or none of the DFs were considered inactive. The model classified correctly all inactive compounds, both in the training and the test sets. As for active compounds, 29 (93.5%) were correctly classified in the training set

and 11 (91.7%) in the test set. The total accuracy of the model was of 96.3% (Table 1) and the Matthew's correlation coefficient is 0.89.

Table 1. Classification accuracy for combination of DF1 and DF2 for the prediction of antibacterial activity against *E. coli*.

Group	Active	Inactive	Accuracy (%)
Training active	29	2	93.5
Training inactive	0	32	100
Test active	11	1	91.7
Test inactive	0	7	100
TOTAL	40	20	96.3

Regarding the indexes affecting the values of DF1, there is a positive sign associated with the presence of ramifications ($^5\chi_p$), the electrotopological index for hydrogen atoms bonded to sp² carbons ($^5H_{tCH}$), and the presence of CI atoms ($^5H_{tCH}$). On the other hand, the order 0 valence connectivity index ($^0\chi v$), directly associated with molecular volume [22], the presence of methanetriyl groups ($^5H_{tCH}$), and the valence charge index $^4G^v$, directly related with the dipolar moment [23], have a negative effect on the values of DF1.

It is worth noting that the presence of chlorine atoms in the molecule favors its antibacterial activity. This was experimentally observed by Fouramadi *et al.* [24], who proved that the potency of a molecule against *E. coli* increased by substituting hydrogen with fluorine and was drastically superior if these were replaced with chlorine atoms (Figure 2).

Figure 2. MIC of a quinolone against *E. coli* depending on the substituents.

Regarding the values of DF2, a positive correlation can be observed with the quotient between valence and non-valence order 0 connectivity index (${}^{O}C$). On the other hand, this correlation is negative with the quotient between valence and non-valence order 2 connectivity index (${}^{2}C$). This indicates the importance of an appropriate polarity

of the molecule, to which these indexes are associated [20]. Similarly, the values of DF2 are negatively affected by valence order 4 cluster connectivity index ($^4\chi_C v$), directly related to molecular volume [25], and the number of types of topologically equivalent vertices (Nclass), which is associated to molecular size and complexity.

When analyzing the indices in both DFs, the presence of a discrete index was detected in DF2, *Nclass*. It was observed that active compounds were grouped within a certain range of values for this index. Following, the empirical distribution of the active and inactive compounds depending on their value for *Nclass* was studied. Analyzing the probability distribution of Figure 3, it was determined that the range of values in which there is a higher probability of finding an active compound is when *Nclass* has values between 11 and 16. This range includes 95.3% of the active compounds. In view of this distribution, *Nclass* can be considered as a new SAR of the antibacterial quinolones against *E. coli*.

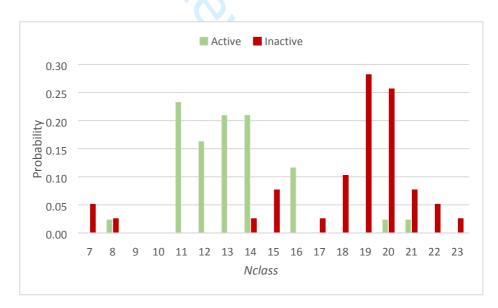


Figure 3. Probability distribution for active and inactive compounds according to their *Nclass* values.

Regarding the provided definition for *Nclass*, number of classes of topologically (symmetry) equivalent graph vertices [18], it must be noted that we could not find any relationship between the term "symmetry" that appears in the definition and the values of *Nclass* calculated for the molecules used to build the model. The only reference we have found that discusses the nature of this index is an article by Stanton [26] in which

he defines *Nclass* as the number of classes of connectivity types identified in a structure. He suggests that this index measures the size and complexity of the molecule, increasing its value proportionally to these factors. Stanton explains that each type of topological substructure (first-order path, second-order path, etc.) are each considered a separate topological class.

Studying this index, we have concluded that, for practical purposes, it **should be defined** as the topological distance (number of axis) between the two farthest vertices of a graph through the shortest path (Figure 4).

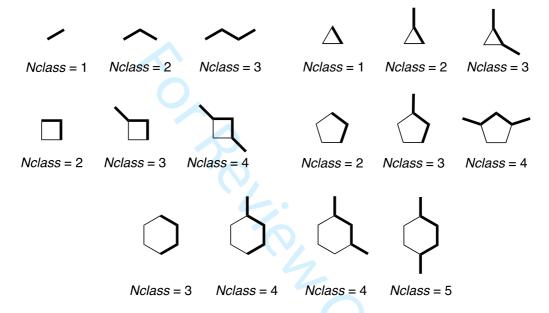


Figure 4. Examples of structures with different Nclass values.

Once both groups were defined, it was established that all compounds with values for Nclass outside the 11-16 range would be directly considered inactive. As for those with values within the range 11-16, DF1 would be applied to determine their activity. Thus, in order for a compound to be classified as active it must have a value for Nclass within 11-16 and a value for DF1 within 0-32. As in the previous case, to avoid false actives, compounds with values for DF1 > 32 are considered unclassifiable.

As it can be observed in Table 2, this model appears to have a lower accuracy percentage for active compounds, however, this is due to the fact that the model is more restrictive, which also means that it will be more efficient in discarding inactive compounds and, therefore, avoiding false actives. Moreover, the discriminatory power

of the model remains unchanged, as shown by its Matthew's correlation coefficient, 0.89.

Table 2. Classification accuracy for the Prediction Model of antibacterial activity against *E. coli*.

Group	Active	Inactive	Accuracy (%)
Training active	20	11	64.5
Training inactive	0	32	100
Test active	11	1	91.7
Test inactive	0	7	100
TOTAL	31	51	89.1

Model validation (Virtual screening)

Traditionally, the search of new active compounds using QSAR methods consists in applying the model to the same family of compounds used for its construction. This is obligatory when the model is obtained using fragmentary parameters [27]. However, the use of overall connectivity parameters along with LDA [28] allows its application to compounds with large structural diversity, yielding excellent results [21, 29, 30].

The model was used for the screening of the virtual library created using the Index Merck database. The model selected 113 theoretically active compounds, of which 47 (42%) had already been described as active against *E. coli* in bibliography. On the other hand, when the combination of DF1 and DF2 is used as a model for the virtual screening of Index Merck 174 molecules are selected as theoretically active but only 27% are described as active against *E. coli* in bibliography (the theoretically active compounds and their values for the DFs and *Nclass* can be found in the Supporting Information).

These results show that the model that combines DF1 and *Nclass* results more restrictive and increases the success rate.

CONCLUSIONS.

The prediction model obtained highlights the importance of molecular size and complexity, represented with *Nclass*, for the antibacterial activity of quinolones against

E. coli. In fact, the *Nclass* index could be included as a filter in existing prediction models for antibacterial activity against *E. coli*. The results obtained prove that molecular topology is a cost-effective, potent and useful tool for the discovery and selection of new antibacterial compounds against *E. coli*.

FUTURE PERSPECTIVES.

Antibiotic resistance has rapidly become a global threat to the public health. The increasing rate of antibiotic-resistant bacteria calls for the constant and urgent introduction of new antibacterial treatments in the market. However, R&D from the pharmaceutical industry in this field is scarce and continues to decrease. Therefore, any breakthrough, however small, made in this field will be of great help to minimize this problem. It is here where QSAR methods play a key role in the future of these treatments. These methods allow the identification of theoretically active molecules in a time- and cost-effective manner. Moreover, we currently live in the world of Big Data, with a constantly growing number of databases containing thousands of chemical compounds. The prediction model developed in this article could be applied to any of these chemical databases in order to find new active compounds against E. coli. Furthermore, the topological index Nclass can act as a filter in other QSAR models predicting activity against E. coli in the same way that the Lipinski's Rule of 5 is used for selecting drug-like molecules. The methodology used to develop this prediction model could also be used to construct models against other resistant bacteria that pose a threat to the public health. On the other hand, an exhaustive search for quinolones has been carried out and structural and activity data are collected. These data can be used by other groups to develop QSAR models with other methodologies. Overall, we consider that QSAR prediction models will continue to evolve and be used for the development of new antibacterial compounds in the fight against drug-resistant bacteria.

SUMMARY POINTS

- Antibiotic resistance is a serious problem that affects worldwide
- There is a lack of research from the pharmaceutical industry

- The search and development of new antibiotics falls on universities and research centers. QSAR methods appear as one of the most effective tools for this purpose.
- A mathematical-topological model with good statistical parameters, capable of selecting compounds with activity against *E. coli*, has been developed.
- The prediction model obtained highlights the importance of molecular size and complexity, represented with *Nclass*, for the antibacterial activity of quinolones against *E. coli*.
- The topological index *Nclass* could be considered as a new SAR of the antibacterial quinolones against *E. coli*.
- The model has been applied to a database of more than 6000 compounds selecting 113 compounds as theoretically active, of which 42% have proven antibacterial activity against *E. coli*.
- The remaining 66 compounds are strong candidates to be developed as new drugs against *E. coli*.
- Molecular topology is a cost-effective, potent and useful tool for the discovery and selection of new antibacterial compounds against *E. coli*.

SUPPLEMENTARY INFORMATION

Supporting information contains definitions of the topological indexes used to obtain the DFs, results for the LDA of the chosen DFs and the compounds selected by the model as theoretically active. It also contains information regarding the quinolones used for the construction of the DFs.

FINANCIAL & COMPETING INTERESTS DISCOLSURE

The authors have no relevant affiliations or financial involvement with any organization or entity with financial interest in or financial conflict with the subject matter or materials discussed in the manuscript. This includes employment, consultancies, honoraria, stock ownership or options, expert testimony, grants or patents received or pending, or royalties.

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SUPPORTING INFORMATION

Table S1. Symbols and definitions of the topological indexes used with DESMOL13¹ and MOLCONN-7² programs

Symbol	Name	Definition	Refs.
Connectivity indexes			
k = 0.10 t = p, c, pc	Kier-Hall indexes Kier-Hall of order k y type path (p), cluster (c) and path-cluster (pc)	${}^k\chi_t = \sum_{j=1}^{k_{n_t}} \left(\prod_{i \in S_j} \mathcal{S}_i\right)^{-1/2}$ \mathcal{S}_i : number of bonds, σ or π of the atom i to non-hydrogen atoms. S_i : j th substructure of order k and type t .	3
$k \times t^{v}$ $k=0-10$ $t=p,c,pc$	Kier-Hall indexes of order k and type path (p), cluster (c) and path-cluster (pc)	${}^{k}\chi_{t}{}^{v} = \sum_{j=1}^{k_{n_{t}}} \left(\prod_{i \in S_{j}} \delta_{i}{}^{v}\right)^{-1/2}$ $\delta_{i}{}^{v}: \text{ Kier-Hall valence of the atom } i.$ $S_{j}: $	2
^k D _t k=0-4 t=p,c,pc	Connectivity differences of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	$^{k}\mathbf{D}_{t}=^{k}\chi_{t}-^{k}\chi_{t}^{\mathbf{v}}$	2
k=0-4 t=p,c,pc	Connectivity ratios of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	$^{k}C_{t} = ^{k}\chi_{t}/^{k}\chi_{t}^{v}$	2
		arge indexes	
G _k k=1-5	Topological charge index of order <i>k</i>	$G_k = \sum_{i=1}^{N-1} \sum_{j=i+1}^{N} \left \mathbf{M}_{ij} - \mathbf{M}_{ji} \right \delta(\mathbf{k}, \mathbf{D}_{ij})$ $\mathbf{M} = \mathbf{A}. \mathbf{Q}: \text{ product of the adjacency and inverse squared distance matrixes for the hydrogen-depleted molecular graph.}$ $\mathbf{D}: \text{ distance matriz.}$ $\delta: \text{ Kronecker delta.}$	4
G _k v k=1-5	Valence topological charge index of order <i>k</i>	$G_k^{\ \ v} = \sum_{i=1}^{N-1} \sum_{j=i+1}^{N} \left \mathbf{M}_{ij}^{\ \ v} - \mathbf{M}_{ji}^{\ \ v} \right \delta(k, \mathbf{D}_{ij})$ $\mathbf{M}^{v} = \mathbf{A}^{v}.\mathbf{Q}$: product of the electronegativity-modified adjacency and inverse squared distance matrixes for the hydrogen-depleted molecular graph. \mathbf{D} : distance matrix. δ : Kronecker delta.	4
J _k , J _k ^v k=1-5	Pondered topological charge indexes of order <i>k</i>	$J_k = \frac{G_k}{N-1} \qquad J_k^{\ v} = \frac{G_k^{\ v}}{N-1}$	4

	Electroto	ppological indexes	
S [⊤] (i)	Sum of electrotopological indexes for a type of atom <i>i</i>	$S_i = I_i + \Delta I_i$ I_i : intrinsic state value of atom i . ΔI_i : delta value for the intrinsic topological state for the atom type i .	5
Gmaxpos	Gmaxpos index	Maximum positive value for the electrotopological state of non-hydrogen atoms in the molecule.	6
Hmaxpos	Hmaxpos index	Maximum positive value for the electrotopological state of hydrogen atoms in the molecule.	7
	Molec	cular properties	
N	Molecular size	Number of non-hydrogen atoms.	4
L	Length	Maximum distance between atoms in terms of bonds	4
PR_i	PRO to PR3	Number of pairs of ramifications separated by <i>i</i> atoms.	4
R	Ramification	Number of simple structural branches.	1
V _k k=3,4	Grade k vertexes	Number of atoms with $k \sigma$ or π bonds, with other atoms (hydrogens not included).	4
knotp	Knotp index	Difference between ${}^3\chi_c$ and ${}^4\chi_{pc}$ indexes	8
knotpv	knotpv index	Difference between ${}^3\chi^{\rm v}{}_c$ and ${}^4\chi^{\rm v}{}_{pc}$ indexes	8
numhbd	Numhbd index	Number of hydrogen-donating atoms in a molecule.	8
numhba	Numhbd index	Number of hydrogen-accepting atoms in a molecule.	8
	Infori	mation indexes	
I _{Shannon}	Shannon index	Index based on the atomic diversity of the molecule.	9
NI, NI2	NI, NI2 index	Indexes based on information theory.	9
BonldW	Bonchev IdW(G) index	Index based on the distribution of topological distances in the molecule.	10
BonldG	Bonchev Id(G) index	Index based on the numer of order 2 subgraphs in the molecule.	10
	Molecul	lar shape indexes	
ⁿ к n=1-3	Kappa index of order <i>n</i>	${}^{1}\kappa = N(N-1)^2/({}^{1}P)^2$ ${}^{2}\kappa = (N-1)(N-2)^2/({}^{2}P)^2$ ${}^{3}\kappa = (N-3)(N-2)^2/({}^{3}P)^2 [N \text{ even; N>3}]$ ${}^{3}\kappa = (N-1)(N-3)^2/({}^{3}P)^2 [N \text{ odd; N>3}]$ N: number of atoms (hydrogens not included) ${}^{1,2,3}P : \text{number of path type subgraphs of orders 1, 2 and 3.}$	11
ⁿ K _ℤ n=1-3	Kappa-alpha index of order	$^{1}\kappa = N + \alpha(N + \alpha - 1)^{2}/(^{1}P + \alpha)^{2}$ $^{2}\kappa = (N + \alpha - 1)(N + \alpha - 2)^{2}/(^{2}P + \alpha)^{2}$ $^{3}\kappa = (N + \alpha - 3)(N + \alpha - 2)^{2}/(^{3}P + \alpha)^{2}$ [N even; N>3]	11

$^{3}\kappa = (N+\alpha-1)(N+\alpha-3)^{2}/(^{3}P+\alpha)^{2}[N+\alpha-3]^{2}$	odd; N>31
N: number of non-hydrogen ato	· -
1,2,3P: number of path type su	
orders 1, 2 and 3.	
$\alpha = \Sigma[(R_i/R_{Csp3})-1]$	
R _i : covalent radius for atom <i>i</i> .	
R _{Csp3} : covalent radius for atom o	Csp³.
$\Phi = (1_{V_{-}}, 2_{V_{-}}) / N$	
φ Phia flexibility index N: number of non-hydrogen ato	oms.
Global topological indexes	
$\delta^{v}_{\cdot} + 1$	
$I_{i} = \frac{\delta_{i}^{v} + 1}{\delta_{i}}; \text{Sum} - I = \sum I_{i}$	
Sum of the intrinsic state	the atom i 5
values 0, number of 0 of 1 boliusot	the atom /
con to other non-hydrogen ato	
δ_i^{v} , Kier-Hall valence of the ator	
Sum of the change in $\Delta I = \frac{I_i - I_j}{r_{ii}^2} \; ; \qquad \text{Sum} - \Delta I = \sum_{i=1}^{n} \frac{I_i - I_j}{r_{ii}^2} \; ;$	$\frac{\mathbf{I_i} - \mathbf{I_j}}{2}$
Sum- ΔI Sum of the change in r_{ij}^2	r_{ij}^2 5
intrinsic state values r _{ij} , number of vertexes between	
atoms.	
Total electrotopological Sum of all the electrotopologic	al indexes
state index in the molecule.	8
Sum of the intrinsic states of all	the atoms
TTd4 index in the molecule.	8
Number of symmetrically equiv	alent types 12
nclass Symmetry index of atoms.	13
Sum of the topological distance	s between
W Wiener index all the non-hydrogen atom	s by the 14
shortest path.	
We Wiener polarity index Number of pairs of atoms with	a distance
Wp Wiener polarity index of 3 bonds.	15
Sum of the topological distanc	es and the
Ww Hyper-Wiener Index squared topological distances b	etween all 15
the non-hydrogen atoms by the	e shortest
path.	
Sum of the topological distance	s between
NAGE Total NAGE on an involve and an analysis of a second	
Wt Total Wiener index all the non-hydrogen atoms by	the longest 15
path.	the longest 15

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Table S2. Structure and references on *E. coli* activity of the compounds used to develop the model.

Compound	IUPAC Name	Structure	References
Act1	(E)-7-(3-amino-4- (methoxyimino)piperi din-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 8-methoxy-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	Р О О О О О О О О О О О О О О О О О О О	Liu H, Huang J, Wang J et al. Synthesis, antimycobacterial and antibacterial evaluation of I-[(1R, 2S)-2- fluorocyclopropyl]fluoroquinol one derivatives containing an oxime functional moiety. Eur J Med Chem 2014; 86:628-638.
Act2	(E)-7-(2- (aminomethyl)-3- (methoxyimino)azetid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-8- methoxy-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N O O O O O O O O O O O O O O O O O O O	Idem
Act3	(E)-7-(3-amino-4- (methoxyimino)piperi din-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	P O O O O O O O O O O O O O O O O O O O	Idem
Act4	(Z)-7-(3- (aminomethyl)-4- (methoxyimino)pyrrol idin-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N N N N F	Idem
Act5	(Z)-7-(3- (aminomethyl)-4- (ethoxyimino)pyrrolid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Act6	(Z)-7-(3- (aminomethyl)-4- (phenoxyimino)pyrrol idin-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N N N F	Idem

Act7	(E)-7-(2- (aminomethyl)-3- (methoxyimino)azetid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N O O O O O O O O O O O O O O O O O O O	Idem
Act8	(E)-7-(2- (aminomethyl)-3- (ethoxyimino)azetidin -1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N N N OH	Idem
Act9	(E)-7-(2- (aminomethyl)-3- (phenoxyimino)azetid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N O O O O O O O O O O O O O O O O O O O	Idem
Act13	(<i>Z</i>)-7-(3- (aminomethyl)-4- (methoxyimino)pyrrol idin-1-yl)-8-chloro-6- fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N N CI N F	Idem
Act14	(Z)-7-(3- (aminomethyl)-4- (ethoxyimino)pyrrolid in-1-yl)-8-chloro-6- fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H ₂ N N CI N F	Idem
Act15	(Z)-7-(3- (aminomethyl)-4- (phenoxyimino)pyrrol idin-1-yl)-8-chloro-6- fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H_2N N CI F F O	Idem

Act16	(E)-7-(3-amino-4- (methoxyimino)piperi din-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4-dihydro- 1,8-naphthyridine-3- carboxylic acid	Р О О О О О О О О О О О О О О О О О О О	Idem
Act17	(Z)-7-(3- (aminomethyl)-4- (methoxyimino)pyrrol idin-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4-dihydro- 1,8-naphthyridine-3- carboxylic acid	H ₂ N N N N F	ldem
Act18	(Z)-7-(3- (aminomethyl)-4- (ethoxyimino)pyrrolid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	H_2N N N N N N N N N N	ldem
Act19	(Z)-7-(3- (aminomethyl)-4- (phenoxyimino)pyrrol idin-1-yl)-6-fluoro-1- (2-fluorocyclopropyl)- 4-oxo-1,4-dihydro- 1,8-naphthyridine-3- carboxylic acid	H_2N N N N N N N N N N	ldem
Act20	(E)-7-(2- (aminomethyl)-3- (methoxyimino)azetid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	H_2N N N N N N N N N N	Idem
Act21	(E)-7-(2- (aminomethyl)-3- (ethoxyimino)azetidin -1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	H_2N N N N N N N N N N	ldem
Act22	(E)-7-(2- (aminomethyl)-3- (phenoxyimino)azetid in-1-yl)-6-fluoro-1-(2- fluorocyclopropyl)-4- oxo-1,4-dihydro-1,8-	H ₂ N N N N F	ldem

	naphthyridine-3- carboxylic acid		
Act26	6,8-difluoro-1-(4-fluorophenyl)-7-(4-(2-((4-(2-fluorophenyl)-5-oxo-2,5-dihydrofuran-3-yl)oxy)ethyl)piperazin -1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F 0 0 0 O O O O O O O O O O O O O O O O	Wang XD, Wei W, Wang PF et al. Novel 3-arylfuran-2(5H)-one-fluoroquinolone hybrid: design, synthesis and evaluation as antibacterial agent. Bioorg Med Chem 2014, 22(14):3620-3628.
Act27	1-cyclopropyl-6,8-difluoro-7-(4-(2-((4-(2-fluorophenyl)-5-oxo-2,5-dihydrofuran-3-yl)oxy)ethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Wang XD, Wei W, Wang PF et al. Novel 3-arylfuran-2(5H)-one-fluoroquinolone hybrid: design, synthesis and evaluation as antibacterial agent. Bioorg Med Chem 2014, 22(14):3620-3628.
Act32	(Z)-7-(3- (aminomethyl)-4- (methoxyimino)-3- methylpiperidin-1-yl)- 1-cyclopropyl-6,8- difluoro-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	O O O O O O O O O O	Chai Y, Liu M, Wang B, et al. Synthesis and in vitro antibacterial activity of novel fluoroquinolone derivatives containing substituted piperidines. Bioorg Med Chem Lett 2010; 20(17):5195-5198.
Besifloxacin	7-(3-aminoazepan-1- yl)-8-chloro-1- cyclopropyl-6-fluoro- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	CLSI Performance Standards for Antimicrobial Susceptibility Testing, 26th Edition.
Ciprofloxacin	1-cyclopropyl-6- fluoro-4-oxo-7- (piperazin-1-yl)-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	ldem
Clinafloxacino	7-(3-aminopyrrolidin- 1-yl)-8-chloro-1- cyclopropyl-6-fluoro- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	ldem
Delafloxacin	1-(6-amino-3,5-difluoropyridin-2-yl)-8-chloro-6-fluoro-7-(3-hydroxyazetidin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	$\begin{array}{c} O & O \\ F & O \\ O & O \\$	ldem

Enoxacin	1-ethyl-6-fluoro-4- oxo-7-(piperazin-1- yl)-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	ldem
Finafloxacin	8-cyano-1- cyclopropyl-6-fluoro- 7- (hexahydropyrrolo[3, 4-b][1,4]oxazin-6(2H)- yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	ldem
Fleroxacin	6,8-difluoro-1-(2- fluoroethyl)-7-(4- methylpiperazin-1-yl)- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	O O O O O O O O O O O O O O O O O O O	Idem
Garenoxacin	1-cyclopropyl-8- (difluoromethoxy)-6- fluoro-7-(1- methylisoindolin-5- yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	HN FO DH	Idem
Gatifloxacin	1-cyclopropyl-6- fluoro-8-methoxy-7- (3-methylpiperazin-1- yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Gemifloxacin	(Z)-7-(3- (aminomethyl)-4- (methoxyimino)pyrrol idin-1-yl)-1- cyclopropyl-6-fluoro- 4-oxo-1,4-dihydro- 1,8-naphthyridine-3- carboxylic acid	O O O O O O O O O O O O O O O O O O O	ldem
Grepafloxacin	1-cyclopropyl-6- fluoro-5-methyl-7-(3- methylpiperazin-1-yl)- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	ldem
Levofloxacin	9-fluoro-3-methyl-10- (4-methylpiperazin-1- yl)-7-oxo-2,3-dihydro- 7 <i>H</i> - [1,4]oxazino[2,3,4- <i>ij</i>]quinoline-6- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem

Lomefloxacin	1-ethyl-6,8-difluoro- 7-(3-methylpiperazin- 1-yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Moxifloxacin	1-cyclopropyl-6- fluoro-8-methoxy-7- (octahydro-6 <i>H</i> - pyrrolo[3,4- <i>b</i>]pyridin- 6-yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Norfloxacin	1-ethyl-6-fluoro-4- oxo-7-(piperazin-1- yl)-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Ofloxacin	9-fluoro-3-methyl-10- (4-methylpiperazin-1- yl)-7-oxo-2,3-dihydro- 7H- [1,4]oxazino[2,3,4- ij]quinoline-6- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Pradofloxacin	8-cyano-1- cyclopropyl-6-fluoro- 7-(octahydro-6 <i>H</i> - pyrrolo[3,4- <i>b</i>]pyridin- 6-yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Pridmore A, Stephan B, Griefe HA. In vitro activity of pradofloxacin against clinical isolates from European field studies In: Proceedings of the American Society for Microbiology (ASM) 105th General Meeting, Atlanta, GA, USA p617, 2005. Abraham J, Ewet K, de Jong A. Pradofloxacin: Comparative in vitro activity against selected pathogens from the US. In: 42nd Interscience Conference on Antimicrobial Agents and Chemotherapy, San Diego, CA, USA p.189.
Sparfloxacin	5-amino-1- cyclopropyl-7-(3,5- dimethylpiperazin-1- yl)-6,8-difluoro-4-oxo- 1,4-dihydroquinoline- 3-carboxylic acid	NH ₂ O O F OH N N HN F ∆	Idem

Trovafloxacin	7-(6-amino-3- azabicyclo[3.1.0]hexa n-3-yl)-1-(2,4- difluorophenyl)-6- fluoro-4-oxo-1,4- dihydro-1,8- naphthyridine-3- carboxylic acid	H ₂ N F	ldem
Ulifloxacin	6-fluoro-1-methyl-4- oxo-7-(piperazin-1- yl)-2a,3-dihydro- 1H,4H- [1,3]thiazeto[3,2- a]quinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
PGE-8367769	1-cyclopropyl-6- fluoro-7,8-dimethyl- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Macinga DR, Renick PJ, Makin KM, et al. Unique biological properties and molecular mechanism of 5,6-bridged quinolones. Antimicrob Agents Chemother. 2003; 47(8):2526-2537.
Inact1	1-ethyl-6,8-difluoro- 4-oxo-7-(4-(((3,4,5- triacetoxy-6- (acetoxymethyl)tetra hydro-2 <i>H</i> -pyran-2- yl)oxy)methyl)-1 <i>H</i> - 1,2,3-triazol-1-yl)-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Azad CS, Bhunia SS, Krishna A, Shukla PK, Saxena AK. Novel Glycoconjugate of 8-Fluoro Norfloxacin Derivatives as Gentamicin-resistant Staphylococcus aureus Inhibitors: Synthesis and Molecular Modelling Studies. Chem Biol Drug Des. 2015; 86(4):440-446.
Inact2	7-(4-(((3-acetamido- 4,5-diacetoxy-6- (acetoxymethyl)tetra hydro-2 <i>H</i> -pyran-2- yl)oxy)methyl)-1 <i>H</i> - 1,2,3-triazol-1-yl)-1- ethyl-6,8-difluoro-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	O O O O O O O O O O O O O O O O O O O	Idem
Inact3	1-ethyl-6,8-difluoro- 4-oxo-7-(4-(2-((5-oxo- 4-phenyl-2,5- dihydrofuran-3- yl)oxy)ethyl)piperazin -1-yl)-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Wang XD, Wei W, Wang PF et al. Novel 3-arylfuran-2(5H)-one-fluoroquinolone hybrid: design, synthesis and evaluation as antibacterial agent. Bioorg Med Chem. 2014, 22(14):3620-3628.

Inact4	7-(4-(2-((4-(4- chlorophenyl)-5-oxo- 2,5-dihydrofuran-3- yl)oxy)ethyl)piperazin -1-yl)-1-ethyl-6,8- difluoro-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	CI O O O O O O O O O O O O O O O O O O O	Idem
Inact5	1-cyclopropyl-6,8-difluoro-7-(4-(2-((4-(4-methoxyphenyl)-5-oxo-2,5-dihydrofuran-3-yl)oxy)ethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact6	1-cyclopropyl-6- fluoro-7-(4-(2-(4- methylpiperazin-1- yl)acetyl)piperazin-1- yl)-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	Б К К К К К К К К К К К К К	Suresh N, Nagesh HN, Renuka J, et al. Synthesis and evaluation of 1-cyclopropyl-6-fluoro-1,4-dihydro-4-oxo-7-(4-(2-(4-substitutedpiperazin-1-yl)acetyl)piperazin-1-yl)quinoline-3-carboxylic acid derivatices as anti-tubercular and antibacterial agents. Eur J Med. 2014; 71:324-332.
Inact7	1-cyclopropyl-7-(4-(2-(4-(3,4-difluorophenyl)piperazin-1-yl)acetyl)piperazin-1-yl)-6-fluoro-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact8	7-(4-(2-(4-(bis(4-fluorophenyl)methyl) piperazin-1-yl)acetyl)piperazin-1-yl)-1-cyclopropyl-6-fluoro-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F N O O O O O O O O O O O O O O O O O O	ldem
Inact9	1-ethyl-6-fluoro-7-(4- (5-(2-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Foroumadi A, Soltani F, Moshafi MH, Ashraf-Askari R. Synthesis and in vitro antibacterial activity of some N-(5-aryl-1,3,4-thiadiazole-2- yl)piperazinyl quinolone derivatives. Farmaco. 2003; 58(10):1023-1028.

Inact10	1-ethyl-6-fluoro-7-(4- (5-(2-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact11	1-cyclopropyl-6- fluoro-7-(4-(5-(3- nitrophenyl)-1,3,4- thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact12	1-ethyl-6-fluoro-7-(4- (5-(3-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact13	1-ethyl-6-fluoro-7-(4- (5-(3-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact14	1-cyclopropyl-6- fluoro-7-(4-(5-(4- nitrophenyl)-1,3,4- thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact15	1-ethyl-6-fluoro-7-(4- (5-(4-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact16	1-ethyl-6-fluoro-7-(4- (5-(4-nitrophenyl)- 1,3,4-thiadiazol-2- yl)piperazin-1-yl)-4- oxo-1,4-dihydro-1,8- naphthyridine-3- carboxylic acid	O_2N	Idem

Inact17	1-cyclopropyl-6-fluoro-7-(4-(2-((6-nitrobenzo[d]thiazol-2-yl)amino)-2-oxoethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	O_2N N S NH N N N O	Sharma PC, Kumar R, Chaudhary M, Sharma A, Rajak H. Synthesis and biological evaluation of novel benzothiazole clubbed fluoroquinolone derivatives. J Enzyme Inhib Med Chem. 2013; 28(1):1-10.
Inact18	1-cyclopropyl-6-fluoro-7-(4-(2-((6-methylbenzo[d]thiazo l-2-yl)amino)-2-oxoethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	N F O O O O O O O O O O O O O O O O O O	Idem
Inact19	1-cyclopropyl-6-fluoro-7-(4-(2-((6-fluorobenzo[d]thiazol-2-yl)amino)-2-oxoethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	F-_N F-_O O O O O O O O O O O O O O O O O O O	Idem
Inact20	7-(4-(2-((6- chlorobenzo[d]thiazol -2-yl)amino)-2- oxoethyl)piperazin-1- yl)-1-ethyl-6-fluoro-4- oxo-1,4- dihydroquinoline-3- carboxylic acid		Idem
Inact21	7-(4-(2-((5,7-dichlorobenzo[d]thiaz ol-2-yl)amino)-2-oxoethyl)piperazin-1-yl)-1-ethyl-6-fluoro-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	CI N F O O O O O O O O O O O O O O O O O O	Idem
Inact22	1-ethyl-6-fluoro-7-(4-(2-((6-nitrobenzo[d]thiazol-2-yl)amino)-2-oxoethyl)piperazin-1-yl)-4-oxo-1,4-dihydroquinoline-3-carboxylic acid	O ₂ N-√N F OH	Idem
Inact23	1-ethyl-6-fluoro-7-(4- (2-((4- methylbenzo[d]thiazo l-2-yl)amino)-2- oxoethyl)piperazin-1- yl)-4-oxo-1,4-	N F O O O O O O O O O O O O O O O O O O	Idem

	dihydroquinoline-3-			
	carboxylic acid			
	1-ethyl-6-fluoro-7-(4-			
	(2-((4-			
	fluorobenzo[d]thiazol	F oo		
	-2-yl)amino)-2-	F OH		
Inact24	oxoethyl)piperazin-1-		Idem	
	yl)-4-oxo-1,4-	S NH N N		
	dihydroquinoline-3-			
	carboxylic acid			
	7-(4-(2-((4-			
	chlorobenzo[d]thiazol			
	-2-yl)amino)-2-			
	oxoethyl)-3-	CI		
	methylpiperazin-1-yl)-	N FYNOH		
Inact25	1-cyclopropyl-6-	S"NH (N"N"	Idem	
	fluoro-8-methoxy-4-			
	oxo-1,4-	'		
	dihydroquinoline-3-			
	carboxylic acid			
	1-cyclopropyl-7-(4-(2-			
	((6,7-			
	dichlorobenzo[d]thiaz			
	ol-2-yl)amino)-2-	0 0		
	oxoethyl)-3-	CI—N F		
Inact26	methylpiperazin-1-yl)-	CI S NH N N	Idem	
	6-fluoro-8-methoxy-	O N Y O A		
	4-oxo-1,4-			
	dihydroquinoline-3-			
	carboxylic acid			
	1-cyclopropyl-6-			
	fluoro-8-methoxy-7-			
	(3-methyl-4-(2-((4-	NO ₂		
	nitrobenzo[d]thiazol-	F TOH		
Inact27	2-yl)amino)-2-	S NH NH N	Idem	
	oxoethyl)piperazin-1-	S NH N N		
	yl)-4-oxo-1,4-			
	dihydroquinoline-3-			
	carboxylic acid			
	1-cyclopropyl-6-			
	fluoro-8-methoxy-7-			
	(3-methyl-4-(2-((4-	0 0		
	methylbenzo[<i>d</i>]thiazo	F OH		
Inact28	l-2-yl)amino)-2-	Shuh Vutt	Idem	
	oxoethyl)piperazin-1-			
	yl)-4-oxo-1,4-			
	dihydroquinoline-3-			
	carboxylic acid			
	1-cyclopropyl-6-	 		
	fluoro-7-(4-(2-((4-	0 0 F ~ Å Å		
Inact29	fluorobenzo[d]thiazol	\(\sum_N \) \(ldem	
mact29	-2-yl)amino)-2-			
	oxoethyl)-3-			
	methylpiperazin-1-yl)-			

Inact30	8-methoxy-4-oxo-1,4-dihydroquinoline-3-carboxylic acid (Z)-7-(3-(aminomethyl)-4-(ethoxyimino)-3-methylpiperidin-1-yl)-1-ethyl-6,8-difluoro-4-oxo-1,4-dihydroguinoline 2	H_2N F O	Chai Y, Liu M, Wang B, et al. Synthesis and in vitro antibacterial activity of novel fluoroquinolone derivatives containing substituted piperidines. Bioorg Med Chem
Inact31	dihydroquinoline-3-carboxylic acid (Z)-7-(3- (aminomethyl)-4- (ethoxyimino)-3- methylpiperidin-1-yl)- 6,8-difluoro-1-(2- fluoroethyl)-4-oxo- 1,4-dihydroquinoline- 3-carboxylic acid	H_2N F O O O F O O F F	Lett. 2010; 20(17):5195-5198.
Inact32	(Z)-7-(3- (aminomethyl)-4- (ethoxyimino)-3- methylpiperidin-1-yl)- 1-cyclopropyl-6- fluoro-8-methoxy-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	H_2N F O	Idem
Inact33	7-(4- cyclopentylpiperazin- 1-yl)-1-ethyl-6-fluoro- 4-oxo-1,4- dihydroquinoline-3- carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Abuo-Rahma Gel-D, Sarhan HA, Gad GF. Design, synthesis, antibacterial activity and physicochemical parameters of novel N-4-piperazinyl derivatives of norfloxacin. Bioorg Med Chem. 2009; 17(11):3879-3886.
Inact34	7-(4-(5,6-diphenyl- 1,2,4-triazin-3- yl)piperazin-1-yl)-1- ethyl-6-fluoro-4-oxo- 1,4-dihydroquinoline- 3-carboxylic acid	F O O O O O O O O O O O O O O O O O O O	Idem
Inact35	7-(4- ((dimorpholinoamino) methyl)piperazin-1- yl)-1-ethyl-6-fluoro-4- oxo-1,4- dihydroquinoline-3- carboxylic acid	O O O O O O O O O O O O O O O O O O O	Idem
Inact36	7-(4-((bis(1,3-dioxoisoindolin-2-yl)amino)methyl)pipe razin-1-yl)-1-ethyl-6-fluoro-4-oxo-1,4-	F O O O O O O O O O O O O O O O O O O O	Idem

	dihydroquinoline-3- carboxylic acid		
PGE-5215205	1-methyl-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	O O O O O O O O O O O O O O O O O O O	Macinga DR, Renick PJ, Makin KM, et al. Unique biological properties and molecular mechanism of 5,6-bridged quinolones. Antimicrob Agents Chemother. 2003; 47(8):2526-2537.
PGE-6116532	7,8-dimethyl-10-oxo-7,10-dihydro-1 <i>H</i> -[1,3]dioxino[5,4- <i>f</i>]quinoline-9-carboxylic acid	O O O O	Idem
PGE-9604297	1-ethyl-4-oxo-1,4- dihydroquinoline-3- carboxylic acid	O O N OH	Idem

Table S3. LDA results and classification of the compounds for DF1.

Compound	⁵ χ ^p	⁰ х ^ν	SH _{tCH}	S _{>CH-}	S _{Cl}	4 G [∨]	DF1	Prob.	Mahalanobis	Clas.
Act2	8.159	16.263	15.196	-2.304	0	5.023	16.321	1.000	3.3	+
Act3	7.579	15.639	14.907	-2.153	0	3.751	17.658	1.000	2	+
Act5	7.651	16.347	15.006	-1.819	0	4.097	10.166	1.000	1.9	+
Act6	8.833	18.026	15.205	-1.914	0	4.352	11.989	1.000	2.7	+
Act8	7.464	15.639	14.894	-2.074	0	3.877	14.862	1.000	1.2	+
Act9	8.649	17.319	15.093	-2.169	0	4.133	16.726	1.000	3.0	+
Act13	7.779	16.696	15.116	-2.037	6.558	4.406	15.117	1.000	5.9	+
Act15	9.120	19.083	15.381	-2.131	6.654	4.773	12.766	1.000	5.8	+
Act16	7.579	15.509	14.845	-2.296	0	3.992	16.902	1.000	1.3	+
Act17	7.492	15.509	14.877	-1.962	0	4.172	14.016	1.000	0.7	+
Act19	8.833	17.896	15.142	-2.056	0	4.539	11.575	1.000	1.8	+
Act20	7.299	14.802	14.766	-2.239	0	3.855	19.311	1.000	2.0	+
Act21	7.464	15.509	14.832	-2.239	0	3.966	15.132	1.000	1.1	+
Act22	8.649	17.189	15.031	-2.334	0	4.222	16.996	1.000	2.7	+
Act26	12.371	23.565	16.194	0	0	5.890	6.631	0.999	16.9	+
Act32	8.121	17.322	15.802	-0.120	0	5.229	6.501	0.999	5.5	+
Besifloxacin	7.123	15.507	15.023	0.007	6.633	3.933	12.765	1.000	5.4	+
Clinafloxacin	6.532	14.092	14.758	0.032	6.526	3.613	16.058	1.000	5.5	+
Delafloxacin	8.007	15.621	14.759	-0.705	6.358	3.525	24.452	1.000	10.8	+
Enoxacin	6.099	12.674	14.432	0	0	3.612	12.801	1.000	1.5	+

Fleroxacin	6.833	14.137	15.183	0	0	4.370	15.105	1.000	2.7	+
Gatifloxacin	7.517	15.289	15.102	0.291	0	4.947	7.809	1.000	4.3	+
Gemifloxacin	7.216	15.261	14.875	-0.023	0	4.177	6.489	0.998	3.0	+
Grepafloxacin	6.875	14.881	15.147	0.426	0	4.671	4.865	0.992	6.8	+
Lomefloxacin	6.625	13.975	15.193	0.075	0	4.215	14.678	1.000	2.7	+
Norfloxacin	6.099	12.804	14.495	0	0	3.492	13.239	1.000	1.4	+
Ofloxacin	7.953	14.659	14.984	-0.158	0	4.715	20.874	1.000	6.1	+
Pradofloxacin	8.563	15.819	15.249	0.709	0	4.714	19.388	1.000	5.3	+
Sparfloxacin	7.493	15.629	15.645	-0.104	0	4.874	13.855	1.000	3.9	+
Ulifloxacin	6.959	13.976	14.618	-1.466	0	3.326	21.277	1.000	3.1	+
PGE-8367769	5.496	10.174	12.289	0	0	2.886	1.606	0.833	21.3	+
Inact1	12.108	26.131	15.833	-7.289	0	7.440	-13.655	1.000	23.2	-
Inact2	12.108	16.223	15.841	-6.715	0	7.409	-15.991	1.000	19.8	-
Inact3	10.359	21.284	15.689	0	0	6.169	-8.384	1.000	2.2	-
Inact4	10.595	22.341	15.704	0	5.942	6.347	-10.479	1.000	4.6	-
Inact5	11.041	22.899	15.925	-0.128	0	6.454	-12.327	1.000	2.2	-
Inact6	8.869	19.374	15.133	0.128	0	5.743	-16.446	1.000	1.8	-
Inact7	10.807	22.362	14.258	0.086	0	6.582	-21.603	1.000	2.6	-
Inact10	10.006	20.313	15.078	0	0	6.222	-13.239	1.000	0.6	-
Inact12	9.864	20.443	15.134	0	0	6.046	-14.485	1.000	0.2	-
Inact13	9.864	20.313	15.072	0	0	6.166	-14.923	1.000	0.4	-
Inact14	9.999	20.728	15.242	0.105	0	6.027	-13.978	1.000	0.4	-
Inact15	9.661	20.443	15.129	0	0	6.100	-17.686	1.000	0.9	-
Inact16	9.661	20.313	15.066	0	0	6.220	-18.124	1.000	1.2	-
Inact17	10.579	21.818	15.214	0.084	0	5.569	-13.815	1.000	5.2	-
Inact19	10.249	20.933	15.184	0.097	0	5.489	-9.911	1.000	4.0	-
Inact20	9.911	21.404	15.086	0	6.018	5.596	-15.126	1.000	4.8	-
Inact21	10.076	22.461	15.105	0	12.24	5.869	-18.458	1.000	20.0	-
Inact22	10.241	21.534	15.101	0	0	5.642	-17.589	1.000	3.9	-
Inact23	9.874	21.270	15.098	0	0	5.999	-22.417	1.000	3.9	-
Inact25	11.449	23.889	15.606	-0.094	6.208	7.119	-17.936	1.000	5.8	-
Inact27	11.951	24.019	15.713	-0.194	0	7.173	-17.932	1.000	2.1	-
Inact28	11.449	23.756	15.705	-0.066	0	7.494	-24.839	1.000	5.5	-
Inact29	11.449	23.134	15.678	-0.147	0	7.022	-16.101	1.000	1.3	-
Inact30	7.887	17.744	15.655	0	0	5.413	-4.360	0.987	8.6	-
Inact31	8.027	17.829	15.669	0	0	5.408	-3.019	0.953	8.8	-
Inact32	8.783	19.059	15.565	0.041	0	6.145	-10.641	1.000	4.5	-
Inact33	7.254	16.157	14.899	0.669	0	3.813	-0.949	0.721	12.6	-

Inact34	11.763	22.366	15.333	0	0	7.261	-11.529	1.000	6.1	-
Inact35	10.137	21.273	15.249	0	0	6.101	-17.285	1.000	1.0	-
Inact36	14.238	25.051	15.478	0	0	8.708	-10.164	1.000	27.6	-
PGE-6116542	5.534	11.096	12.509	0	0	3.868	-9.599	1.000	16.7	-
PGE-9604297	3.866	8.805	11.844	0	0	2.026	-8.748	1.000	21.9	-
Act1	8.447	16.970	15.275	-2.358	0	5.128	14.206	1.000	2.8	+
Act4	7.492	15.639	14.939	-1.819	0	3.986	14.431	1.000	0.8	+
Act7	7.299	14.932	14.828	-2.075	0	3.766	19.042	1.000	1.7	+
Act14	7.938	17.403	15.182	-2.036	6.584	4.518	10.876	1.000	6.3	+
Act18	7.651	16.216	14.943	-1.961	0	4.283	9.750	1.000	1.6	+
Act27	11.059	21.869	15.892	-0.143	0	6.153	-0.862	0.297	11.0	-
Ciprofloxacin	6.438	13.088	14.608	0.188	0	3.419	16.691	1.000	1.5	+
Finafloxacin	8.563	15.521	15.208	-0.038	0	4.689	24.096	1.000	8.4	+
Garenoxacin	9.017	16.993	15.369	-3.330	0	4.279	31.564	1.000	16.2	+
Levofloxacin	7.953	14.659	14.984	-0.158	0	4.715	20.874	1.000	6.1	+
Moxifloxacin	8.563	16.281	15.320	0.887	0	4.973	13.866	1.000	3.5	+
Trovafloxacin	8.393	15.349	14.853	0.565	0	3.318	24.779	1.000	15.3	+
Inact8	13.099	26.326	15.409	-0.094	0	7.518	-30.988	1.000	14.4	-
Inact9	10.006	20.443	15.141	0	0	6.102	-12.801	1.000	0.4	-
Inact11	10.203	20.728	15.247	0.101	0	5.973	-10.763	1.000	1.1	-
Inact18	10.249	21.555	15.207	0.115	0	5.773	-17.312	1.000	3.0	-
Inact24	9.874	20.648	15.071	0	0	5.527	-13.928	1.000	2.1	-
Inact26	11.681	24.046	15.712	-0.121	12.36	7.308	-19.835	1.000	22.8	-
PGE-5215205	3.637	8.098	11.728	0	0	1.724	-4.952	0.993	24.8	-

Table S4. LDA results and classification of the compounds for DF2.

Compound	⁴ χ _c ^ν	n _{class}	°C	² C	DF2	Prob.	Mahalanobis	Clas.
Act2	0	13	1.337	1.833	5.859	0.997	0.4	+
Act3	0	14	1.334	1.794	6.334	0.998	0.6	+
Act5	0	14	1.319	1.763	5.500	0.996	0.3	+
Act6	0	16	1.330	1.786	4.250	0.986	1.1	+
Act8	0	14	1.334	1.809	5.619	0.996	0.4	+
Act9	0	16	1.344	1.829	4.421	0.988	1.4	+
Act13	0	13	1.302	1.721	5.509	0.996	0.2	+
Act15	0	16	1.302	1.742	1.852	0.864	1.3	+
Act16	0	14	1.346	1.834	6.311	0.998	0.9	+
Act17	0	13	1.346	1.809	8.305	1.000	1.6	+
Act19	0	16	1.340	1.819	4.258	0.986	1.3	+
Act20	0	13	1.362	1.863	8.481	1.000	2.2	+
Act21	0	14	1.346	1.851	5.532	0.996	0.7	+
Act22	0	16	1.354	1.865	4.387	0.988	1.8	+
Act26	0	21	1.351	1.920	-3.228	0.038	10.9	-
Act32	0.125	14	1.299	1.660	-0.122	0.469	17.5	-
Besifloxacin	0	11	1.244	1.572	5.006	0.993	3.5	+
Clinafloxacin	0	11	1.269	1.629	6.265	0.998	2.0	+
Delafloxacin	0	12	1.402	1.977	10.478	1.000	6.2	+
Enoxacin	0	11	1.307	1.307	1.475	0.814	4.8	+
Fleroxacin	0	12	1.345	1.344	8.171	1.000	1.3	+
Gatifloxacin	0	11	1.262	1.262	3.154	0.959	2.7	+
Gemifloxacin	0	13	1.311	1.310	5.241	0.995	0.0	+
Grepafloxacin	0	11	1.249	1.249	3.599	0.973	3.2	+
Lomefloxacin	0	11	1.309	1.309	5.153	0.994	0.8	+
Norfloxacin	0	11	1.293	1.293	1.781	0.856	3.8	+
Ofloxacin	0	12	1.268	1.268	1.386	0.800	3.2	+
Pradofloxacin	0	12	1.282	1.282	9.296	1.000	4.4	+
Sparfloxacin	0	11	1.301	1.300	9.621	1.000	3.0	+
Ulifloxacin	0	11	1.217	1.217	7.474	0.999	9.0	+
PGE-8367769	0	8	1.334	1.334	4.057	0.983	8.1	+
Inact1	0	19	1.357	2.083	-7.856	1.000	8.7	-
Inact2	0	19	1.352	2.053	-7.238	0.999	7.0	-
Inact3	0	20	1.301	1.835	-6.318	0.998	1.0	-
Inact4	0	21	1.278	1.791	-8.756	1.000	1.8	-
Inact5	0	22	1.290	1.778	-7.164	0.999	1.7	-

Inact6	0	18	1.238	1.613	-4.228	0.986	1.5	-
Inact7	0	22	1.289	1.714	-4.435	0.988	2.8	-
Inact10	0	18	1.293	1.787	-3.447	0.969	0.5	-
Inact12	0	19	1.285	1.759	-4.382	0.988	0.3	-
Inact13	0	19	1.293	1.788	-4.434	0.988	0.4	-
Inact14	0	20	1.281	1.693	-2.934	0.949	2.0	-
Inact15	0	20	1.285	1.758	-5.266	0.995	0.5	-
Inact16	0	20	1.293	1.787	-5.318	0.995	0.6	-
Inact17	0	21	1.289	1.694	-2.597	0.931	3.4	-
Inact19	0	20	1.268	1.657	-3.254	0.963	2.1	-
Inact20	0	20	1.227	1.643	-9.080	1.000	2.7	-
Inact21	0	20	1.208	1.616	-10.864	1.000	5.2	-
Inact22	0	21	1.293	1.755	-4.758	0.991	1.4	-
Inact23	0	19	1.235	1.653	-7.417	0.999	1.5	-
Inact25	0	19	1.214	1.581	-7.435	0.999	2.8	-
Inact27	0	20	1.273	1.675	-3.379	0.967	1.8	-
Inact28	0	19	1.221	1.590	-6.802	0.999	2.3	-
Inact29	0	19	1.254	1.637	-3.776	0.978	1.4	-
Inact30	0.125	15	1.292	1.739	-5.337	0.995	13.8	-
Inact31	0.125	15	1.325	1.758	-1.316	0.789	16.3	-
Inact32	0.125	15	1.255	1.639	-7.002	0.999	14.4	-
Inact33	0	14	1.228	1.592	-1.222	0.772	4.5	-
Inact34	0	19	1.275	1.835	-9.459	1.000	3.3	-
Inact35	0	17	1.219	1.687	-9.587	1.000	5.3	-
Inact36	0	18	1.323	1.862	-2.154	0.896	1.9	-
PGE-6116542	0	8	1.301	1.899	1.853	0.136	14.0	+
PGE-9604297	0	7	1.329	1.949	4.929	0.007	18.6	+
Act1	0	14	1.323	1.807	3.920	0.981	0.3	+
Act4	0	13	1.334	1.771	8.302	1.000	1.5	+
Act7	0	13	1.350	1.820	8.538	1.000	1.9	+
Act14	0	14	1.289	1.715	2.959	0.951	0.6	+
Act18	0	14	1.331	1.799	5.472	0.996	0.3	+
Act27	0	20	1.319	1.795	-1.668	0.159	6.5	-
Ciprofloxacin	0	11	1.287	1.701	5.858	0.997	0.9	+
Finafloxacin	0	12	1.307	1.689	8.595	1.000	1.9	+
Garenoxacin	0	13	1.338	1.825	6.511	0.999	0.5	+
Levofloxacin	0	12	1.268	1.713	1.386	0.800	3.2	+
Moxifloxacin	0	12	1.246	1.579	4.004	0.982	3.1	+

Trovafloxacin	0	13	1.389	1.849	13.288	1.000	10.3	+
Inact8	0	23	1.274	1.695	-6.917	0.999	2.8	-
Inact9	0	18	1.285	1.757	-3.391	0.967	0.4	-
Inact11	0	19	1.281	1.694	-2.048	0.886	2.0	-
Inact18	0	20	1.232	1.603	-6.532	0.999	1.9	-
Inact24	0	19	1.272	1.709	-4.165	0.985	0.5	-
Inact26	0	20	1.197	1.561	-10.064	1.000	5.3	-
PGE-5215205	0	7	1.358	1.995	7.379	0.001	22.6	+

Table S5. Theoretically active compounds selected by the prediction model of antibacterial activity against *E. coli* (DF1= 0-32 and *Nclass* = 11-16).

Compound	Activity	DF1	Nclass
Aconine	Alcaloid	29.800868	11
Aconitine	Antipyretic	23.4464385	14
Adlumidine	Plant extract	5.37152537	11
Albofungin	Antibacterial agent	17.1846	14
9-aminocamptothecin	Antitumoral	4.64123362	11
Anabsinthin	Plant extract	3.22992963	11
Asperlicin	Plant extract	4.05486582	14
Asperuloside	Anti-inflammatory	4.6412	12
Baknkosin	Plant extract	1.91637848	11
Balofloxacin	Antibacterial agent	10.6166	12
Baptigenin	Plant extract	5.8222	11
Bicuculline	GABA Antagonist	5.37152537	11
Bis(4-amino-1-anthraquinonyl)amine	Plant extract	0.6841	14
Brequinar	Antitumoral	11.2589198	14
Brucine	Plant extract	22.8943611	11
Camptothecin	10.3Antitumoral	7.46488877	11
Carminic acid	Pigment	13.3912435	13
Cervicarcin	Antitumoral	1.43479591	11
Cevine	Plant extract	8.4642941	12
Chlorophacinone	Pesticide	14.7436	11
Cichorin	Plant extract	24.5077405	11
Cinolazepam	Sedative	10.5092	11
Clocortolone	Corticoid	28.5745441	12
Daidzein	Antioxidant	20.5348	11
Danofloxacin	Antibacterial agent	4.3332	12
Daphinin	Plant extract	2.50387559	11
Delsoline	Plant extract	18.9492644	11
Delcosine	Plant extract	27.2274329	11
Demeclocycline	Antibacterial agent	19.2969	11
Destomycin A	Antibacterial agent	9.82704032	14
Difloxacin	Antibacterial agent	3.7894	13
Diflufenican	Pesticide	17.0194	14

Difluprednate	Anti-inflammatory	28.6290376	14
Dolasetron	Antiemetic	3.6360	13
Doxacycline	Antibacterial agent	3.90377797	11
Droxicam	Anti-inflammatory	6.0101	11
Elliptone	Antitumoral agent	14.2946905	12
Enrofloxacin	Antibacterial agent	0.3662	13
Florantyrone	Choleretic	6.1612	11
Flumetsulam	Herbicide	7.5798	11
Flumiclorac	Pesticide	6.2391	11
Flumioxazin	Herbicide	16.1175	11
Flunoxaprofen	Anti-inflammatory	27.9084	12
Fluocortolone	Glucocorticoid	1.1966	12
Fluperolone	Glucocorticoid	30.4742166	14
Flurprednisolone	Corticosteroid	23.4515	12
Frangulin A	Laxative	26.6393	13
Genistein	Antitumoral	7.1546	11
Genitopicrin	Plant extract	5.44208396	11
Germine	Plant extract	23.624289	12
Glucofrangulin	Laxantive	2.0069	14
a-Glucogallin	Plant extract	3.2521793	11
b-Glucogallin	Plant extract	3.2521793	11
Halopredone	Anti-inflammatory	25.8426115	14
Halophytine	Plant extract	31.8583805	13
Hygromycin B	Anthelmintic	9.82704032	14
Indaconitine	Plant extract	10.940243	14
Justicidin B	Plant extract	6.40057596	11
Laccaic acid	Pigment	5.20682652	14
Leucodrin	Plant extract	6.94981692	11
Linezolid	Antibacterial agent	31.7005098	14
Loganin	Plant extract	9.65111285	12
Lyxoflavine	Vitamin	23.6352	11
, Marbofloxacin	Antibacterial agent	31.7005098	12
Medicagol	Plant extract	24.6561522	11
Melezitose	Carbohydrate	22.8365434	12
Meprednisone	Glucocorticoid	10.5627254	12
Metacycline	Antibacterial agent	30.0751418	11
Minocycline	Antibacterial agent	7.8405858	12
Mometasone	Anti-inflammatory	0.87176752	14
Moxifloxacin	Antibacterial agent	7.8406	12
Nalbuphine	Analgesic	28.3357209	11
Narcotoline	Plant extract	10.1735761	12
Noscapine	Antitussive	15.7870044	12
Ochratoxin A	Mycotoxin	0.5502	14
Paroxetine	Antidepressant	2.00305135	13
Pefloxacin	Antibacterial agent	9.81034608	12
Pipebuzone	Anti-inflammatory	5.8395182	12
Pirazolac	Anti-inflammatory	4.67725356	12
i ii azolac	Anti-iiiiaiiiiiatoi y	4.07723330	14

Pirenoxine	Cataract treatment	6.7677	11
Pirenzepine	Anti-ulcer	1.10982539	11
Plumieride	Plant extract	9.5018	12
Prednyldiene	Glucocorticoid	2.8233	12
Pseudobaptigenin	Plant extract	2.94054093	11
Quatrimycin	Antibacterial agent	15.2208	11
Quininone	Anti-malaria	1.7458218	12
Quinoxyfen	Fungicide	13.5283125	11
Rhamnetin	Plant extract	5.1892	11
Rhodomycin B	Vitamin	4.8605	14
Riboflavin	Vitamin	4.1640	11
Ribostamycin	Antibacterial agent	2.43943805	13
Rotenone	Antiparasitic	17.0583372	14
Rubitecan	Topoisomerase inhibitor	3.1031	12
Rufloxacin	Antibacterial agent	30.4049	12
Sancycline	Antibacterial agent	2.21199125	11
Scoparin	Reagent	18.8206	13
Scutellarein	Plant extract	2.5795	11
Sitafloxacin	Antibacterial agent	25.0367	11
Skimmin	Plant extract	5.9765	11
Sporidesmin A	Mycotoxin	18.7458006	11
Swertiamarin	Plant extract	8.0637	12
Tectorigenin	Plant extract	0.7760	11
Temafloxacin	Antibacterial agent	23.3822	12
Tephrosin	Plant extract	24.7192314	14
Tetracycline	Antibacterial agent	13.2993	11
Tetrahydrocortisone	Steroid	10.1895718	12
Topotecan	Antitumoral	5.55463332	13
Tosufloxacin	Antibacterial agent	27.1346	12
Tritoqualine	Antihistaminic	2.46150506	13
Tropisetron	Antiemetic	0.07165887	11
Trovafloxacin	Antibacterial agent	26.8002	13
Verbenalin	Plant extract	14.237474	12
Vindoline	Plant extract	3.73294636	11

Table S6. Theoretically active compounds selected by DF1=0-32 and DF2 = 0-12.

Compound	Activity	DF1	DF2
Aflatoxin B1	Mycotoxin	9.618	1.958
Aflatoxin G1	Mycotoxin	11.559	0.934
Aflatoxin M1	Mycotoxin	12.535	2.979
Afloqualone	Sedative	9.819	2.292

Allo a firmacina	A satisfication	17.105	1.020
Albofungin	Antibiotic	17.185	1.938
Alizarine Blue	Pigment	22.699	10.918
Amarogentin	Plant extract	2.291	7.186
Amarolide	Plant extract	1.043	4.068
1-Aminoanthraquinone	Plant extract	10.378	9.906
1-Aminoanthraquinone-2-	Plant extract	4.241	13.995
carboxylic Acid	A 122	4.644	0.705
9-Aminocamptothecin	Antitumoral	4.641	0.705
Anagyrine	Antitumoral	0.654	0.925
Anilinephthalein	Laxative	8.941	0.814
Anisindione	Anticoagulant	0.406	1.861
Anthralin	Epithelium recovery	4.225	11.037
Anthraquinone	Plant extract	11.361	7.597
Anthrone	Pigment	8.565	2.751
Apigetrin	Plant extract	4.021	8.722
Ascorbigen	Analgesic	12.955	8.936
Asperuloside	Anti-inflammatory	10.617	12.004
Balofloxacin	Antibiotic	5.822	2.159
Baptigenin	Plant extract	0.684	10.734
Benzanthrone	Pigment	24.204	1.888
Benzotrifluoride	Reagent	2.722	9.268
Bis(4-amino-1-	Plant extract	14.744	3.637
anthraquinonyl)amine			
Bostycoidin	Antibiotic	2.152	5.667
Cacotheline	PH indicator de pH	8.485	11.067
Capecitabine	Antitumoral	3.260	2.162
Carbamazepine	Anticonvulsivo	2.293	1.411
Catalposide	Plant extract	16.829	9.558
Chartreusin	Antibiotic	18.747	0.597
p-Chlorobenzotrifluoride	Disolvente	0.036	4.983
Chlorophacinone	Pesticida	10.509	2.361
Chrysophanic Acid	Antitumoral	3.613	11.098
Cinolazepam	Sedative	20.535	2.909
Clorindione	Anticoagulant	7.082	2.688
Coriamyrtin	Plant extract	10.060	2.745
Coumestrol	Estrógeno	18.284	4.224
Cyclic AMP	Bioquímica	0.566	8.623
Cyclic GMP	Bioquímica	5.844	11.099
Cytisine	Deshabituación tabáquica	2.143	5.982
Daidzein	Antioxidante	4.333	3.781
Daidzin	Plant extract	10.111	7.826
Danofloxacin	Antibiotic	19.297	5.084
Demeclocycline	Antibiotic	3.789	13.391
DFDD	Reagent	9.992	2.475
Difloxacin	Antibiotic	17.019	2.198
Diflufenican	Pesticida	3.636	1.121
p-Difluorobenzene	Reagent	13.717	11.182
4,4'-Difluorodiphenyl	Reagent	13.834	4.032
Diphenadione	Pesticida	11.710	5.941
Dolasetron	Antiemético	6.010	1.029
Doxefazepam	Sedative	23.886	3.308
Doxenitoin	Anticonvulsivo	1.347	0.065
Droxicam	Anti-inflammatory	0.366	10.286
Efavirenz	Antirretroviral	15.277	6.834
		10.277	J.JJ-∓

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Emodin	Plant extract	3.495	13.836
Enrofloxacin	Antibiotic	6.161	0.562
Fialuridine	Antiviral	21.908	2.047
Florantyrone	Colerético	7.579	2.298
Fluconazole	Antifúngico	0.171	2.875
Flucytosine	Antifúngico	13.759	9.461
Fludioxonil	Antifúngico	23.124	5.194
Fluindione	Anticoagulant	18.360	10.631
Flumequine	Antibiotic	21.524	12.749
Flumetsulam	Herbicida	6.239	8.265
Flumiclorac	Pesticida	16.117	5.789
Flumioxazin	Herbicida	27.908	7.576
Flunitrazepam	Sedative	13.288	4.558
Flunoxaprofen	Anti-inflammatory	1.197	0.873
Fluocortolone	Glucocorticoide	23.452	0.267
Fluorescamine	Reagent	18.442	1.999
Fluorescein	Pigment	25.072	3.622
p-Fluoroaniline	Reagent	13.212	8.286
p-Fluorophenylacetic Acid	Reagent	6.264	12.133
o-Fluorotoluene	Reagent	9.259	1.824
m-Fluorotoluene	Reagent	5.390	2.597
p-Fluorotoluene	Reagent	9.311	3.750
Fluprednisolone	Corticosteroide	26.639	2.575
Flurbiprofen	Anti-inflammatory	14.648	5.300
Fluroxypyr	Herbicida	2.846	1.372
Flutriafol	Antifúngico	13.144	3.853
Frangulin A	Laxative	7.155	9.696
Frenolicin	Antiparasitario	4.828	8.867
Fustin	Neuroprotector	7.058	13.472
Gallein	Pigment	24.412	10.141
Genistein	Antitumoral	2.007	6.992
Gentisin	Antiviral	5.527	1.451
Gibberellic Acid	Fertilizante	30.002	5.712
Glitoxin	Inmunosupresor	9.067	3.585
Glucofrangulin A	Laxative	23.635	11.657
Hematein	Pigment	3.157	11.289
Hydromorphone	Opioide	20.049	2.054
Ibafloxacin	Antibiotic	18.183	9.465
Indanthrene®	Pigment	31.605	0.605
Indigo	Pigment	13.425	3.415
Isatide	Plant extract	24.274	10.064
Isoapo-b-erythroidine	Plant extract	6.432	0.785
Isolysergic Acid	Mycotoxin	4.916	3.877
Kaempferitrin	Plant extract	14.571	7.671
Kaempferol	Plant extract	0.769	10.628
Lufenuron	Pesticida	1.580	3.299
Luteolin	Plant extract	5.221	10.116
Lyxoflavine	Vitamina	4.753	8.181
Mefloquine (Hydrochloride)	Antimalárico	8.442	11.944
Minocycline	Antibiotic	7.841	8.446
Mitomycin C	Antitumoral	18.701	8.687
Morphine	Opioide	17.945	2.699
Moxifloxacin	Antibiotic	24.496	4.004
Naloxone	Antagonista opioide	23.532	3.191
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Naltrexone	Antagonista onioido	21.426	0.985
1,4-Naphthoquinone	Antagonista opioide Plant extract	1.274	10.704
Naringin	Edulcorante	10.094	6.304
Nevirapine	Antirretroviral	6.738	0.304
Niflumic Acid	Anti-inflammatory	1.552	6.326
Nivalenol	Mycotoxin	23.403	12.953
Normorphine	Opioide	16.243	6.101
Nybomycin	Antiviral	11.526	1.667
Ochratoxin A	Mycotoxin	0.550	1.469
Parvaguone	Antimalárico	3.785	9.887
Phanquinone	Antibiotic	3.144	7.081
Phenanthrenequinone	Pigment	11.628	7.597
Phenindione	Anticoagulant	8.812	7.509
Phenolphthalein	pH indicator	13.187	2.809
Phenytoin	Anticonvulsant	6.540	3.795
Phloridzin	Plant extract	8.333	10.775
Phorbol	Reagent	8.466	2.659
Picrotin	Antidote	27.510	1.635
Pirenoxine	Cataract treatment	6.768	4.632
Pleurotin(e)	Antibiotic	23.089	6.937
Plumieride	Plant extract	9.502	9.842
Polyoxin A	Antibiotic	9.308	13.504
Porfiromycin	Antitumoral	10.050	6.506
Prenylidene	Glucocorticoid	2.823	0.664
Praquiloside	Phytotoxin	15.896	4.538
Quatrimycin	Antibiotic	15.221	11.019
Quercetin	Nutritional supplement	10.659	13.791
Quinoline Yellow Spirit Soluble	Pigment	19.567	3.226
Rhamnetin	Plant extract	5.189	5.750
Rheochrysdin	Plant extract	12.872	10.868
Rhodomycin B	Plant extract	4.860	2.477
Riboflavin	Vitamin	4.164	8.181
Rolitetracycline	Antibiotic	11.944	1.408
Ruberythric Acid	Pigment	31.628	9.565
Rubitecan	Topoisomerase inhibitor	3.103	2.746
Rufloxacin	Antibiotic	30.405	1.920
Samaderin A	Plant extract	27.932	3.512
Santonic Acid	Reagent	9.248	7.186
Scoparin	Reagent	18.821	11.310
Scutellarein	Plant extract	2.579	9.906
Silymarin (Silybin)	Hepatoprotector	9.201	2.780
Sitafloxacin	Antibiotic	25.037	5.695
Skimmin	Plant extract	5.976	12.474
Sophorabioside	Plant extract	20.341	3.322
Sorbinil	Aldose reductase inhibitor	30.107	10.312
Spectinomycin	Antibiotic	9.085	8.145
Swetiamarin	Plant extract	8.064	13.759
Tazobactam	Antibiotic	7.542	13.804
Tectorigenin	Plant extract	0.776	3.647
Tegafur	Antitumoral	6.778	12.566
Temafloxacin	Antibiotic	23.382	6.824
Tetracycline	Antibiotic	13.299	11.019
Tetrantoin	Anticonvulsant	9.523	5.549
Theaflavine	Antioxidant	14.521	10.273

Thiazopyr	Herbicide	3.995	10.359
Thioxanthone	Printing product	10.137	3.000
Tosufloxacin	Antibiotic	27.135	10.909
Trovafloxacin	Antibiotic	26.800	13.288
Truxillic acid	Reagent	8.467	13.039
Vellosimine	Plant extract	9.002	0.168
Viridin	Antifungal	21.757	4.967
Vomitoxin	Mycotoxin	4.228	7.937
Voriconazole	Antifungal	7.369	5.230



Group	Active	Inactive	Accuracy (%)
Training active	29	2	93.5
Training inactive	0	32	100
Test active	11	1	91.7
Test inactive	0	7	100
TOTAL	40	20	96.3



Group	Active	Inactive	Accuracy (%)
Training active	20	11	64.5
Training inactive	0	32	100
Test active	11	1	91.7
Test inactive	0	7	100
TOTAL	31	51	89.1

