

Article Similarity-Based Virtual Screening to Find Antituberculosis Agents Based on Novel Scaffolds: Design, Syntheses and Pharmacological Assays

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Abstract: A method to identify molecular scaffolds potentially active against the Mycobacterium tuberculosis complex (MTBC) is developed. A set of structurally heterogeneous agents against MTBC was used to obtain a mathematical model based on topological descriptors. This model was statistically validated through a Leave-n-Out test. It successfully discriminated between active or inactive compounds over 86% in database sets. It was also useful to select new potential antituberculosis compounds in external databases. The selection of new substituted pyrimidines, pyrimidones and triazolo[1,5-*a*]pyrimidines was particularly interesting because these structures could provide new scaffolds in this field. The seven selected candidates were synthesized and six of them showed activity in vitro.

Keywords: MTBC; virtual screening; topological indices; linear discriminant analysis; pharmacological activity distribution diagrams; antimicrobial drugs; drug design

1. Introduction

Tuberculosis is one of the deadliest infections in the world, killing almost 1.45 million people annually [1,2]. Most of these deaths occur in poor countries, and a more rational distribution of wealth could prevent them. However, the increasing incidence of drug-resistant MTBC has caused its resurgence in developed countries and makes therapeutic alternatives necessary [3,4]. Although new compounds potentially active against these bacteria are under active research, very few new treatments have been developed in recent years [5–9]. Therefore, more extensive investigations are needed to find new molecular scaffolds capable of generating new inhibitors of mycobacteria.

The α -acetylenic ketones of type A (Scheme 1) have been shown to be highly versatile building blocks. For instance, these conjugated ynones have proven to be very suitable substrates for the synthesis of (E)-3-acylpropenoic acids [10], and of a wide range of heterocyclic systems [11–21], including the synthesis of the natural product L-lathyrine and related analogues [22,23]. Furthermore, when properly functionalized, compounds of type A have also proven to be valuable substrates for combinatorial and parallel synthesis on solid support of highly molecular diverse 2,4,6-trisubstituted pyrimidines [24–27]. These α -acetylenic ketones are produced readily, mainly by direct palladium-catalyzed coupling of acyl chlorides with 1-alkynes (Sonogashira reaction), [28,29] by reaction of alkynylzinc chlorides with acid halides, [30,31] or by reaction of lithium or magnesium acetylides



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Scheme 1. Cyclocondensation of α -acetylenic ketones A with 3-amino-5-benzylsulfanyl-1,2,4-triazole.

In this paper, we have further expanded the synthetic utility of these conjugated ynones A, and we wish to report the synthesis of novel 2,5,7-substituted triazolo[1,5-*a*]pyrimidines B by cyclocondensation of different α -acetylenic ketones A with 3-amino-5-benzylsulfanyl-1,2,4-triazole (Scheme 1).

The triazolo[1,5-*a*]pyrimidine nucleus is of considerable chemical and pharmacological interest. Antibiotic, anticholesteremic, antidiabetic, antiallergic, anti-inflammatory, antipyretic, antiphlogistic, analgesic and anticancer activities have been described for these types of compounds. They also serve in the treatment and prevention of circulatory diseases such as hypertension, heart diseases, stroke, hypercholesterol, arteriosclerosis and are effective coronary vasodilators and bronchodilators [36–49].

On the other hand, several triazolo[1,5-*a*]pyrimidine derivatives have found applications in the agrochemical industry. Additionally, 1,2,4-triazolo[1,5-*a*]pyrimidinesulfonamides are used as herbicides and plant growth inhibitors, and they show activity against acetolactate synthase [46,50,51].

Currently, there are many methods available to synthesize triazolo[1,5-*a*]pyrimidines, mainly they are cyclocondensations between 3-amino-1,2,4-triazoles and 1,3-bifunctional synthons such as 1,3-dicarbonyl compounds or their equivalents [52–59], vinylogous iminium salts [47,60,61], ketene dithioacetals [48,62,63], and 3-ketovinyl compounds [46]. We used α -acetylenic Ketones of types A as 1,3-bifunctional synthons.

Although a few pyrimidines [64], pyrimidones [65] and triazolo[1,5-*a*]pyrimidines [66] have already been synthetized and tested for anti-TB activity, their substitution patterns and synthesis methods are different from the compounds described here.

The identification of new targets requires knowledge of the specific biochemical pathways of mycobacteria, but many metabolic processes are still unknown and the structurebased design of new anti-TB agents is a complex task [67].

On the other hand, extra-mechanistic virtual screening methodologies have demonstrated their ability to model the presence of activity within structurally heterogeneous groups of compounds in different therapeutic areas [68–77] as well as in predicting toxicological properties [78,79] and drug-like characteristics [80]. In these models, structural similarity is the key. Molecules are characterized through structural invariants, that is, by descriptors that are independent of molecular conformation. Many of them are topological indices (TI) [81–87], which are capable of characterizing most of the molecular structure [88–95].

The aim of this study was to develop new discriminant models based on the molecular structures of antituberculosis agents, shown in Table S1 [96–98] and inactive substances, shown in Table S2, characterized by topological indices [99] (Table 1), in order to screen structural databases to identify new potentially useful scaffolds against MTBC.

Symbol	Name	Definition	Reference
N	Molecular size	Number of non-hydrogen atoms.	[68]
V <i>k</i> <i>k</i> = 3, 4	Vertices of degree k	Number of atoms having k bonds, σ or π , to non-hydrogen atoms.	[68]
R	Ramification	Number of single structural branches.	[68]
W	Wiener path number	Sum of the distances between any two atoms in terms of bonds.	[100]
L	Length	Maximal distance between atoms in terms of bonds.	[68]
PRk $k = 0-3$	Pairs of ramifications at distance k	Number of pairs of single branches at distance <i>k</i> in terms of bonds.	[68]
k = 0 - 4 t = p, c, pc	Randić-like indices of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	${}^{k}\chi_{t} = \sum_{j=1}^{{}^{k}n_{t}} \left(\prod_{i \in S_{j}} \delta_{i}\right)^{\frac{-1}{2}}$ δ_{i} , number of bonds, σ or π , of the atom <i>i</i> to non-hydrogen atoms. S_{j} , <i>j</i> th sub-structure of order <i>k</i> and type <i>t</i> .	[71,81,82]
$k_{\chi t}^{v} k = 0 - 4$ t = p, c, pc	Kier-Hall indices of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	${}^{k}\chi_{t}{}^{V} = \sum_{j=1}^{{}^{k}n_{t}} \left(\prod_{i \in S_{j}} \delta_{i}{}^{V}\right)^{\frac{-1}{2}}$ $\delta_{i}{}^{v}, \text{ Kier-Hall valence of the atom } i.$ $S_{j, j}\text{ th sub-structure of order } k \text{ and type } t.$	[71,81,82]
G _k k = 1–5	Topological charge indices of order <i>k</i>	$G_k = \sum_{i=1}^{N-1} \sum_{j=i+1}^{N} \left M_{ij} - M_{ji} \right \delta\left(k, D_{ij}\right)$ M = AQ, product of the adjacency and inverse squared distance matrices for the hydrogen-depleted molecular graph. D, distance matrix. $\delta, \text{ Kronecker delta.}$	[68,101]
G_k^v k = 1-5	Valence topological charge indices of order <i>k</i>	$G_k^{\ V} = \sum_{i=1}^{N-1} \sum_{j=i+1}^{N} \left M_{ij}^{\ V} - M_{ji}^{\ V} \right \delta\left(k, D_{ij}\right)$ $M^V = A^V Q, \text{ product of the}$ electronegativity-modified adjacency and inverse squared distance matrices for the hydrogen-depleted molecular graph. D, distance matrix. δ , Kronecker delta.	[68,101]
J_k k = 1-5	Pondered topological charge indices of order <i>k</i>	$J_k = \frac{G_k}{N-1}$	[68,101]
$J_k^{\rm v}$ k = 1-5	Pondered valence topological charge indices of order <i>k</i>	$J_k{}^V = \frac{G_k{}^V}{N-1}$	[68,101]
k = 0-4 t = p, c, pc	Connectivity differences of order <i>k</i> and type path (p), cluster (c) and	${}^{k}D_{t} = {}^{k}\chi_{t} - {}^{k}\chi_{t}{}^{V}$	[68]

path-cluster (pc)

Table 1. Descriptors used.

Symbol	Name	Definition	Reference
E_k k = 1–5	Topological charge differences of order <i>k</i>	$E_k = G_k^{V} - G_k$	[102]
F_k k = 1–5	Pondered topological charge differences of order <i>k</i>	$F_k = J_k^{V} - J_k$	[102]
k = 0 - 4 t = p, c, pc	Connectivity quotients of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	${}^{k}C_{t} = \frac{{}^{k}\chi_{t}}{{}^{k}\chi_{t}{}^{V}}$	[68]
k = 0-4 t = p, c, pc	Inverse connectivity quotients of order <i>k</i> and type path (p), cluster (c) and path-cluster (pc)	${}^{k}Q_{t} = \tfrac{{}^{k}\chi_{t}{}^{V}}{{}^{k}\chi_{t}}$	[102]
CG_k k = 1–5	Topological charge quotients of order <i>k</i>	$CG_k = rac{G_k}{G_k^V}$	[102]
$\begin{array}{c} QG_k\\ k=1-5 \end{array}$	Inverse topological charge quotients of order <i>k</i>	$QG_k = rac{G_k^V}{G_k}$	[102]

Table 1. Cont.

2. Results and Discussion

2.1. Antituberculosis Activity Modelling

Figure 1 illustrates the proposed virtual screening procedure. In this method, thresholds on descriptor values were combined with Linear Discriminant Analysis (LDA) to give a qualitative discriminant model. Table S3 in the Supplementary Materials collects all these values.

Given a population, for example of molecules, that can be classified into several groups according to their experimental properties, for example a group of molecules with a pharmacological activity and another without this activity, LDA is a method to find linear combinations of independent variables (for example the aforementioned structural invariants) whose numerical values can be used to distinguish between these different categories. When two categories are defined, the classification is done by the so-called discriminant function (DF).





The following equation DF was obtained by stepwise LDA using the set shown in Table 2:

$$DF = 0.88 - 11.99 J_1 + 4.86 J_1^{v} - 11.11 J_3^{v} + 0.81 {}^{1}D - 0.2 {}^{4}C_c$$
$$N = 60 \lambda = 0.34 F = 21.12$$

where N = 60 represents 25 anti-MTBC drugs and 35 presumably inactive compounds.

Active Group				Inactive Group			
Compound	DF	Prob. ^a Active	Predicted Activity ^b	Compound	DF	Prob. ^a Inactive	Predicted Activity ^b
Streptonicozid	3.14	1.000	+	Butibufen	-3.14	1.000	_
Tobramycin	3.49	1.000	+	Aldicarb	-2.74	1.000	_
Kanamycin	3.88	1.000	+	Antrafenine	-2.55	1.000	_
Amikacin	4.14	1.000	+	Carprofen	-1.89	0.997	_
Dihydrostreptomycin	2.80	0.999	+	Beclobrate	-1.84	0.997	_
Streptomycin	2.32	0.997	+	Benzoctamine	-1.77	0.996	_
Ethambutol	2.23	0.996	+	Carmofur	-1.77	0.996	_
Pyrazinamide	1.97	0.992	+	Aminothiazole	-1.73	0.996	_
Énviomycin	1.92	0.991	+	Acifran	-1.72	0.996	_
Ofloxacin	1.82	0.988	+	Brilliant Blue	-1.72	0.996	_
Moxifloxacin	1.66	0.982	+	Amitraz	-1.68	0.995	_
Gatifloxacin	1.66	0.981	+	Clofibrate	-1.62	0.994	_
Rifampin	1.60	0.978	+	Paraoxon	-1.52	0.992	_
Azithromycin	1.49	0.971	+	Piroxicam	-1.39	0.989	_
Verazide	1.44	0.967	+	Carmustine	-1.11	0.977	_
Isoniazid	1.20	0.938	+	Ornithine	-1.11	0.977	_
Capreomycin	1.16	0.930	+	Alpidem	-1.10	0.976	_
Sparfloxacin	0.98	0.889	+	Alprazolam	-1.04	0.972	_
Clarithromycin	0.93	0.874	+	Bixin	-1.04	0.972	_
Salinazid	0.92	0.871	+	Benzoic Acid	-1.03	0.971	—
Tuberin	0.91	0.869	+	Amsacrine	-0.98	0.967	_
Clofazimine	0.13	0.432	NC	Azaserine	-0.96	0.965	_
Imipenem	0.09	0.404	NC	Theofibrate	-0.88	0.957	_
Ph-Aminosalicylate	-0.200	0.230	_	Azacosterol	-0.83	0.951	_
PAS	-1.000	0.031	_	Allicin	-0.77	0.943	_
				Prazepam	-0.77	0.943	_
				Camazepam	-0.71	0.933	_
				Aminopromazine	-0.57	0.904	—
				Bromazepam	-0.45	0.870	_
				Carnitine	-0.32	0.824	_
				Acipimox	-0.28	0.809	—
				Buspirone	-0.03	0.677	—
				Azapicyl	0.03	0.640	—
				Acronine	0.11	0.588	NC
				Captodiamine	0.28	0.470	NC

Table 2. Classification for each compound by DF.

^a Prob., probability; ^b When probability active or probability inactive > 0.60, the compound is classified as active "+" or inactive "-", respectively. In any other case, the compound is considered non-classified "NC".

The topological descriptors selected in this equation were: the charge indices (J_1, J_1^v, J_3^v) ; the difference index $({}^1D = {}^1\chi - {}^1\chi^v)$; and the quotient index $({}^4C_c = {}^4\chi_c/{}^4\chi_c^v)$, where ${}^m\chi_t$ and ${}^m\chi_t{}^v$ are, respectively, single and valence Randić-Kier-Hall indices of order m and type t. Topological charge-transfer indices, J_1, J_1^v and J_3^v , are measures of the contribution of molecular topological structure to the charge transfer at topological distance 1 and 3, respectively [68,101]. Difference and quotient indices are related to charge distributions within molecular fragments [68]. Thus, 1D is the net contribution of the heteroatoms to the electronic charge within fragments of order 1 (bonds). The 4C_c index can be related to electron densities of cluster-type fragments of order 4 (three atoms bonded to a central one) in which there is at least one heteroatom. The maximum correlation between pairs of these selected variables, for the group of actives, was as weak as 0.424, corresponding to the pair J_1^v and J_3^v . Thus, the variables can be considered as independent. Table S4 shows these intercorrelations.

Table 2 shows the results of the classification for each one of the compounds included in the LDA.

The linear equation gave good results since most compounds were classified with a probability over 86% (Table 2). When the probabilities are between 40% and 60%, the compounds were counted as non-classified (NC), and finally below 40%, they were considered as inactive. Under this framework, the error percentage in the active set was about 9%, whereas in the inactive was 0%. These probabilities are the so-called posterior probabilities, computed by the Bayes rule as the probability of classifying a case (molecule) conditioned to the model obtained. Let π_k the prior probability: $\pi_k = \frac{N \text{ of cases in class } k}{Total N \text{ of cases}}$. The posterior probability *P* is given by: $P = \frac{f_k(x)\pi_k}{\sum_{i=1}^k f_i(x)\pi_i}$ where $f_k(x)$ is the class-conditional density of the case x in the class k. Assuming that this density for x, given every class k, follows a normal distribution, the density formula for a multivariate Gaussian distribution is applicable. $\frac{\exp\left[-\frac{1}{2}(x-\mu_k)^T Cov_k^{-1}(x-\mu_k)\right]}{\sqrt{(2-x)^2 + k(2-x)}}$ where x and the mean μ_k are both column vectors, Thus, $f_k(x) =$ $\sqrt{(2\pi)^p \cdot \det(Cov)}$ *Cov* is the covariance matrix and *p* is its dimension. The denominator involves the square root of the determinant of this matrix. The result of the matrix multiplications in the numerator is a scalar number.

The results of the internal validation are illustrated in Table 3, with the percentage of correct classification within each group. Five runs were performed. A number of compounds ranging from 9 to 16 were randomly extracted from the training to a test set. Wilks λ values are shown for each equation. The lower the Wilks λ value, the better the discrimination. Correct classification percentages are shown for training and test sets, for active and inactive compounds. The number of compounds classified as active (+) or inactive (-) appears in parentheses, where (a/b) = number of (+) compounds/number of (-) compounds. Average values are also shown, as well as the performance of DF function.

Daras Ma		Trainin	g Group	Test Group		
Kun No.	Λ	(+)	(—)	(+)	(—)	
1	0.28	90% (18/2)	100% (0/26)	60% (3/2)	100% (0/9)	
2	0.31	87% (20/3)	96% (1/27)	50% (1/1)	100% (0/7)	
3	0.35	86% (19/3)	89% (3/24)	100% (3/0)	75% (2/6)	
4	0.33	82% (14/3)	100% (0/27)	88% (7/1)	100% (0/8)	
5	0.33	91% (20/2)	96% (1/27)	100% (3/0)	86% (1/6)	
Average	-	87%	96%	80%	92%	
DF	0.34	84% (21/4)	97% (1/34)	No	No	

Table 3. Results of the internal validation for DF.

(a/b) = number of (+) compounds/number of (-) compounds.

The results for the training and test groups are within the same range. The mean percentage of success obtained with the training group for DF (Table 3) was 87% for active (+) and 94% for inactive (-). For the test it was 80% and 92%, respectively. The results were similar to those obtained with the DF equation, which points out the validity of the LDA equation.

The results of the external validation test are shown in Table 4. As can be seen, for the active set there was a misclassified compound, namely ethionamide. The same is found in the opposite group, where glucosamine was misclassified.

Active Group				Inactive Group				
Compound	DF	Prob. ^a Active	Predicted Activity ^b	Compound	DF	Prob. ^a Inactive	Predicted Activity ^b	
Morphazinamide	4.32	1.000	+	Canthaxanthin	-3.46	1.000	_	
Neomycin	5.02	1.000	+	Genite	-2.57	1.000	_	
Tubercidin	2.85	0.999	+	Altretamine	-1.91	0.997	_	
Ciprofloxacin	2.15	0.995	+	Etifoxin	-1.86	0.997	_	
Viomycin	1.75	0.986	+	Dichlone	-1.78	0.996	_	
Rifabutin	0.73	0.799	+	Feprazone	-1.47	0.991	_	
Ethionamide	-1.30	0.014	_	Antipyrine	-0.44	0.868	_	
				Benorylate	-0.25	0.792	_	
				Chloropal	0.19	0.526	NC	
				Glucosamine	1.77	0.014	+	

Table 4. Results of the external validation for DF.

^a Prob., probability; ^b When probability active or probability inactive > 0.60, the compound is classified as active "+" or inactive "-", respectively. In any other case, the compound is considerate non-classified "NC".

Figure 2 shows the PDD obtained from DF. As can be inferred from Figure 2, the optimal range of DF to find active compounds can be established between 0 and 4.5.



Figure 2. PDD in the training set, obtained by DF equation. E, expectancy of activity or inactivity; black line, active group; grey line, inactive group.

2.2. Similarity-Based Virtual Screening

A virtual library containing new pyrimidine derivatives generated by combining the chemical scaffolds and substituents depicted in Table 5 was screened as described in Section 3.2. After building the database containing the substituted pyrimidines, the virtual screening was performed. Descriptors calculated for the entire database are available to the readers upon request to the corresponding author. Based on these parameters, a virtual screening was carried out so that those molecular structures with the values of the descriptors and DFs within the thresholds, shown in the Table S3, were selected as candidates. Table S5, under Supplementary Materials, shows the calculated descriptors for the selected compounds. Seven potentially active molecules were selected. These structures are presented in Figure 3. These compounds selected as possible candidates were synthesized and tested in vitro.



 Table 5. Scaffolds and fragments used for the generation of the virtual library.



Figure 3. Structures of the selected compounds.

2.3. Chemistry

Synthesis of 1,2,4-triazolo[1,5-*a*]pyrimidine derivatives. Reaction of the required magnesium acetylides, generated from alkynes 1a-b and *i*-PrMgCl in dry THF at 0 °C, with piperonal 2, afforded the expected propargylic alcohols 3a-b. Oxidation of 3a-b with MnO₂ afforded α -acetylene ketones 4a and 4b in 79% and 37% overall yield, respectively (Scheme 2).



Reagents and conditions: i) *i*-PrMgCl, THF, 0°C, 5-12h., ii) MnO₂, CH₂Cl₂, 0°C-r.t., 3-12h., iii) DBU, DMF, MgSO₄, 40°C, 1-2h.

Scheme 2. Synthesis of protected 1,2,4-triazolo[1,5-a]pyrimidine derivatives 6.

Cyclocondensation of 3-amino-5-benzylsulfanyl-1,2,4-triazole **5** with α -acetylene ketones **4a-b** in dry DMF at 40 °C, gave the corresponding triazolo[1,5-*a*]pyrimidines **6a-b** in moderate yields (73% and 38%, respectively) (Scheme 2).

Finally, deprotection of 6b in acidic conditions afforded 7 in 77% yield (Scheme 3).



Scheme 3. Deprotection of 6b to give 7.

Synthesis of pyrimidine derivatives. Two of us reported on the synthesis of novel 4-alkoxypyrimidines starting from 2-alkylsulfanylpyrimidinones of type 8 [103,104]. The method is based on a selective *O*-alkylation reaction with bulky aliphatic alcohols using the Mitsunobu conditions (method A, Scheme 4) or in basic medium with sterically demanding agents like α -haloketones (method B, Scheme 4).



Scheme 4. O-alkylation of 2-alkylsulfanylpyrimidinones 8.

Oxidation of the thioether moiety to the corresponding sulfone **10** using *m*-CPBA and nucleophilic displacement by different nucleophiles produced the corresponding highly molecular diverse pyrimidines of type **11**. In addition, when 4-isopropoxipyrimidine **11a** was treated with a 1:1 mixture of $H_2SO_4/AcOH$ at 90 °C for 15 min, the selective cleavage of the 4-isopropoxy group took place yielding 2-aryloxypyrimidinone **12** (Scheme 5).



Reagents and conditions: i) *m*-CPBA, CH₂Cl₂, 0°C to r.t., 2 h; ii) Ar-NH₂, dioxane, reflux, 30 h ; iii) PhOH, Cs₂CO₃, dioxane, 60 °C, 3 h; iv) H₂SO₄/AcOH 1:1, 90 °C, 15 min

Scheme 5. Oxidation of 9 to the corresponding sulfone 10, nucleophilic displacement to produce the corresponding pyrimidines 11, and displacement of the 4-isopropoxy group of 11a to yield 2-aryloxypyrimidinone 12.

On the other hand, the reaction of the deprotected amine **9e** with phenylboronic acid and glyoxylic acid (Petasis reaction) [105,106] gave the desired α -phenyl glycine derivative **13** in moderated yield (Scheme 6).



Scheme 6. Petasis reaction of the amine 9e to give α -phenyl glycine derivative 13.

Finally, we prepared the compound **14** by reducing the carbonyl group of compound **9d** with NaBH₄ in MeOH. This reduction afforded the hydroxy derivative **14** in 80% isolated yield (Scheme 7).



Scheme 7. Reduction of 9d to obtain the hydroxy derivative 14.

Microbiological study. To check the predicted antituberculosis activity of the selected candidates, microbiological tests were performed. The result of in vitro susceptibility test of conventional drugs is shown in Table 6, and the experimental results of the selected compounds are illustrated in Table 7. Two compounds, **7** and **12**, showed MIC₅₀ = MIC₉₀ = 32 mg/L, corresponding to 81.5μ M and 121.1μ M, respectively; four compounds, **6a**, **9c**, **11b** and **13**, showed MIC₉₀ = 64 mg/L against *M. tuberculosis*, which corresponds to 146, 167.8, 160.1, 131.8 μ M, respectively; and **14** was inactive at the assayed concentrations. The compound **11b** showed the best MIC₅₀ = 80μ M, but its MIC₉₀ raised to 160.1 μ M. Thus, the results pointed out that 7 was the most active agent, showing molar MIC₉₀ four times lower than Ethambutol but within the same magnitude order.

Compound	MIC Range/mg/L	MIC ₅₀ /mg/L	MIC ₉₀ /mg/L	Molecular Mass/Da	MIC ₅₀ /µM	MIC ₉₀ /μM
Ethambutol	1–8	4.0	4.0	204.3	19.6	19.6
Isoniazid	0.05-0.2	0.05	0.05	137.1	0.4	0.4
Rifampin	0.125 - 1	0.125	0.5	823.0	0.2	0.6
Streptomycin	0.125–0.5	0.25	0.5	581.6	0.4	0.9

Table 6. Experimental MIC of conventional drugs against MTBC.

MIC range: minimal and maximal inhibitory values found.

Compound	MIC Range/mg/L	MIC ₅₀ /mg/L	MIC ₉₀ /mg/L	Molecular Mass/Da	MIC ₅₀ /μM	MIC ₉₀ /µM
6a	64	64	64	438.5	146.0	146.0
7	32	32	32	392.4	81.5	81.5
9c	64	64	64	381.4	167.8	167.8
11b	16 - 64	32	64	399.8	80.0	160.1
12	32	32	32	264.3	121.1	121.1
13	32 - 64	64	64	485.6	131.8	131.8
14	>128	>128	>128	414.5	>308.8	>308.8

Table 7. Experimental MIC of the selected compounds against MTBC.

MIC range: minimal and maximal inhibitory values found.

3. Materials and Methods

3.1. Antituberculosis Activity Modelling

A group of 32 compounds with known activity against MTBC were compiled from various sources [96–98]. Their structures are shown in Table S1 in the Supplementary Material. A set of 45 compounds with a different pharmacological activity was also used as inactive group. Their structures are shown in Table S2 in the Supplementary Material. The compounds were characterized by a set of 90 descriptors calculated using the DesMol program [99]. The descriptors used with their symbols, definitions and references are shown in Table 1. These were used to build a model capable of discriminating between active and inactive antituberculous compounds.

DFs were calculated with randomly selected subsets of 25 out of 32 active and 35 out of 45 inactive compounds by using BMDP New System [107]. Descriptor selection was based on the Fisher-Snedecor F parameter. The variables were introduced step by step in the DF: in each step, the variable that added the most to the separation of the groups was entered in the equation, or the variable that contributed the least to improving said separation was eliminated. The classification criterion was the shortest Mahalanobis distance. The discriminant ability of the DF was evaluated by two parameters, Wilks λ , and the percentage of correct classification in each group. The independent variables in this study were the calculated structural invariants, and the discrimination property was the presence of activity against MTBC.

The validation of the selected DF was performed by two methods. One internal leave*n*-out test in which the program randomly chose and pulled out a ratio of compounds, and used them to evaluate the DF obtained with the rest; and another external test with a previously unused data set.

To choose the optimal ranges of values for this equation, the corresponding Pharmacological Distribution Diagram, PDD [108], was obtained. These diagrams are useful to determine the intervals of the equation in which the probability of finding new candidates is maximum. This is a graph similar to a histogram in which expectancies appear on the ordinate axis. For an arbitrary range of values of a given function, the activity expectancy is $E_a = a/(i + 100)$, where a is the percentage of active compounds in the range and i is the corresponding percentage of inactive within the same range. The expectancy of inactivity is similarly defined as $E_i = i/(a + 100)$. This plot provides a good visualization of the regions of minimal overlap between the active and inactive compounds and helps to select the optimal interval of the DF.

The 90 descriptors and the selected discriminant function were used as filters to select potential candidates in structural databases. For this, the maximum and minimum values of each descriptor were established as thresholds for the 90 variables, while for DF, the optimal interval was taken. Compounds exhibiting all 90 values within the thresholds and DF values within the optimal intervals were considered as candidates.

3.2. Similarity-Based Virtual Screening

A virtual library containing 320 structures of substituted pyrimidines was generated by combining the chemical scaffolds and substituents depicted in Table 5, using in-house software. The set of 90 descriptors and DFs were calculated for this library. A virtual screening was carried out based on these parameters to select as candidates those molecular structures whose values for the descriptors and DFs were within the thresholds.

3.3. Chemical Methods

DMF was dried over activated molecular sieves (4 Å). THF was dried over Na/ benzophenone prior use. All the other commercially available chemicals were used as purchased without further purification. Reactions involving magnesium acetylides and synthesis of triazolo[1,5-*a*]pyrimidines were run under a dry Ar atmosphere. Melting points (capillary tube) were measured with an electrothermal digital melting point apparatus IA 91,000 and are uncorrected. IR spectra were recorded on a Mattson-Galaxy Satellite FT-IR. Additionally, ¹H and ¹³C NMR spectra were recorded at 200 and 50 MHz, respectively, on a Brucker DPX200 Advance instrument with TMS as internal standard. MS spectra were recorded on a VG Quattro instrument in the positive ionization FAB mode, using 3-NBA or 1-thioglycerol as the matrix or in a Thermo Quest 2000 series apparatus for the EI (70 eV) mode. Analytical TLC was performed on precoated TLC plates, silica gel 60 F₂₅₄ (Merck). Flash-chromatography (FC) purifications were performed on silica gel 60 (230–400 mesh, Merck).

Synthesis of propargylic alcohols 3a-b. General procedure.

To a cooled (0 °C) solution of the corresponding alkyne **1a-b** in dry THF (2 mL/mmol), *i*-PrMgCl (2 M solution in THF) was added dropwise under Ar. The mixture was stirred at that temperature for 4 h. Then, a solution of the piperonal **2** (1.3 equiv) in dry THF (1 mL/mmol) was slowly added dropwise over a period of 15 min. The reaction mixture, under Ar, was stirred from 0 °C to r.t. until total consumption of **1a-b** (5–12 h., monitored by TLC). The reaction was quenched with saturated solution of NH₄Cl (3 mL/mmol) at r.t. and the organic solvent was eliminated under reduced pressure. The aqueous layer was extracted with AcOEt (3 × 3 mL/mmol) and the combined organic layers were dried over MgSO₄, the solvent was evaporated and the resulting residue purified by flash-chromatography (n-hexane:AcOEt).

1-Benzo[1,3]dioxol-5-yl-3-phenyl-prop-2-yn-1-ol (3a). According to the general procedure described above, the reaction between 1a (4.01 g, 39.26 mmol) and piperonal 2 (7.65 g, 50.9 mmol) afforded 8.02 g (81%) of 3a as a white solid. m.p.: 60–61 °C. IR (KBr): ν 3439 (br., OH). ¹H NMR (CDCl₃): δ 7.5–7.4 (m, 2*H*_{arom}), 7.3 (m, 3*H*_{arom}), 7.2–7.1 (m, 2*H*_{arom}), 6.85 (d, 1*H*_{arom}, *J* = 8.0 Hz), 6.00 (s, 2H, OCH₂O), 5.63 (s, 1H, CHC≡C), 2.59 (br., 1H, OH). ¹³C NMR (CDCl₃): δ 147.8, 147.6, 134.6 (3s, 3*C*_{arom}), 131.7, 128.6, 128.3 (3d, 5CH_{arom}), 122.2 (s, *C*_{arom}), 120.4, 108.1, 107.4 (3d, 3CH_{arom}), 101.2 (t, CH₂), 88.6 (s, *C*≡CPh), 86.5 (s, *C*≡CPh), 64.8 (d, CH). MS (FAB⁺) *m*/*z*: 253 ([M+1]⁺, 9). Anal. Calcd. for C₁₆H₁₂O₃: C, 76.18; H, 4.79. Found: C, 76.39; H, 4.61%.

1-Benzo[1,3]dioxol-5-yl-3-(tetrahydro-pyran-2-yloxy)-prop-2-yn-1-ol (3b). According to the general procedure described above, the reaction between **1b** (0.21 g, 1.48 mmol) and piperonal **2** (0.29 g, 1.92 mmol) afforded the crude of the compound **3b**. The reaction mixture was used in the next step of synthesis without further purification.

Synthesis of α -acetylenic ketones 4a-b. General procedure.

Over a cooled (0 °C), mechanically stirred suspension of MnO_2 (5 equiv) in CH_2Cl_2 (3 mL/mmol), a solution of the propargyl alcohol **3a** or of the reaction mixture **3b** in CH_2Cl_2 (3 mL/mmol) was added dropwise. The reaction mixture was stirred from 0 °C to r.t. until total consumption of **3a-b** (3–12 h., monitored by TLC) and then filtered through a Celite[®] pad. The solvents were removed under reduced pressure and the resulting residue was purified by flash-chromatography (n-hexane:AcOEt).

1-Benzo[1,3]dioxol-5-yl-3-phenyl-propynone (4a). According to the general procedure described above, the reaction between **3a** (1.98 g, 7.87 mmol) and MnO₂ (3.84 g,

39.35 mmol) afforded after crystallization of the residue with MeOH:H₂O instead of the chromatographic purification 1.91 g (97%) of **4a** as a yellow solid. m.p.: 100–101 °C. IR (KBr): v 1625 (s, C=O). ¹H NMR (CDCl₃): δ 7.92 (dd, 1H_{arom}, *J* = 8.2 Hz, *J*' = 1.8 Hz), 7.7–7.6 (m, 3H_{arom}), 7.5–7.4 (m, 3H_{arom}), 6.93 (d, 1H_{arom}, *J* = 8.2 Hz), 6.10 (s, 2H, OCH₂O). ¹³C NMR (CDCl₃): δ 176.0 (s, *C*=O), 152.8, 148.1 (2s, 2C_{arom}), 132.9 (d, 2CH_{arom}), 131.9 (s, C_{arom}), 130.6, 128.6, 127.2 (3d, 4CH_{arom}), 120 (s, C_{arom}), 108.2, 107.9 (2d, 2CH_{arom}), 102.1 (t, CH₂), 92.3 (s, C=CPh), 86.7 (s, C=CPh). MS (FAB⁺) *m*/*z*: 251 ([M+1]⁺, 100). Anal. Calcd. for C₁₆H₁₀O₃: C, 76.79; H, 4.03. Found: C, 76.53; H, 4.24%. IR and NMR spectra can be seen in Figure S1.

1-Benzo[1,3]dioxol-5-yl-4-(tetrahydro-pyran-2-yloxy)-but-2-yn-1-one (4b). According to the general procedure described above, the reaction between the reaction mixture of **3b** and MnO₂ (0.72 g, 7.4 mmol) afforded 157 mg (overall yield = 37%) of **4b** as a yellow oil. IR (NaCl): v 1640 (m, C=O). ¹H NMR (CDCl₃): δ 7.82 (dd, 1 H_{arom} , *J* = 7.4 Hz, *J'* = 1.8 Hz), 7.55 (d, 1 H_{arom} , *J* = 1.6 Hz), 6.89 (d, 1 H_{arom} , *J* = 8.2 Hz), 6.08 (s, 2H, OCH₂O), 4.9–4.8 (m, 1H, CH), 4.56 (s, 2H, CH₂OThp), 3.9–3.8 (m, 1H, CH₂), 3.6–3.5 (m, 1H, CH₂), 1.9–1.6 (m, 6H, 3CH₂). ¹³C NMR (CDCl₃): δ 175.6 (s, C=O), 152.9, 148.1 (2s, 2C_{arom}), 131.5 (s, C_{arom}), 127.4 (d, CH_{arom}), 108.2, 107.9 (2d, 2CH_{arom}), 102.1 (t, CH₂), 97.3 (d, CH), 89.6 (s, C≡CCH₂), 83.3 (s, C≡CCH₂), 62.0, 54.0, 30.1, 25.2, 18.8 (5t, 5CH₂). MS (FAB⁺) *m/z*: 289 ([M+1]⁺, 100). Anal. Calcd. for C₁₆H₁₆O₅: C, 66.66; H, 5.59. Found: 66.94; H, 5.76%. IR and NMR spectra can be seen in Figure S2.

Synthesis of triazolo[1,5-a]pyrimidines 6a-b. General procedure.

A mixture of 3-amino-5-benzylsulfanyl-1,2,4-triazole **5**, DBU and anhydrous MgSO₄ (5 g/g) in dry DMF (3 mL/mmol) was heated at 40 °C under Ar for 30 min. Then, a solution of the corresponding α -acetylenic ketone **4a-b** in dry DMF (3 mL/mmol) was slowly added using a *syringe pump* over a period of 5h. The reaction mixture, under Ar, was stirred at 40 °C until total consumption of **5** (1–2 h., monitored by TLC). The reaction was filtered and the organic solvent was eliminated under reduced pressure. The residue was dissolved with CH₂Cl₂ (15 mL/mmol) and was washed with saturated solution of NH₄Cl (3 × 3 mL/mmol). The organic layer was dried over MgSO₄, the solvent was evaporated and the resulting residue was purified by flash-chromatography (n-hexane:AcOEt).

5-Benzo[1,3]dioxol-5-yl-2-benzylsulfanyl-7-phenyl-1,2,4-triazolo[1,5-*a*]pyrimidine (6a). According to the general procedure described above, reaction between 5 (106 mg, 0.52 mmol), 4a (303 mg, 1.21 mmol, 2.5 equiv), DBU (81 μL, 0.53 mmol, 1.1 equiv) and anhydrous MgSO₄ (0.5g) afforded 166 mg (73%) of 6a as a white solid. m.p.: 143–144 °C. ¹H NMR (DMSO-*d*₆): δ 8.3–7.1 (m, 14*H*_{arom}), 6.21 (s, 2H, OC*H*₂O), 4.59 (s, 2H, PhC*H*₂S). ¹³C NMR (DMSO-*d*₆): δ 166.4, 159.4, 158.1, 150.1, 148.1, 146.1, 137.7 (7s, 8C_{arom}), 131.5 (d, CH_{arom}), 130.1 (s, C_{arom}), 129.7, 128.9, 128.4, 128.3, 127.2, 123.0, 108.5, 107.3, 105.6 (9d, 13CH_{arom}), 101.8, 34.4 (2t, 2CH₂). MS (FAB⁺) *m*/*z*: 439 ([M+1]⁺, 100). Anal. Calcd. for C₂₅H₁₈N₄O₂S: C, 68.48; H, 4.14; N, 12.78; S, 7.31. Found: C, 68.29; H, 4.27; N, 13.01; S, 7.02%. IR and NMR spectra can be seen in Figure S3.

5-Benzo[1,3]dioxol-5-yl-2-benzylsulfanyl-7-(tetrahydro-pyran-2-yloxymethyl)-1,2, 4-triazolo[1,5-*a*]pyrimidine (6b). According to the general procedure described above, the reaction between **5** (255 mg, 1.24 mmol), 4b (698 mg, 2.42 mmol, 2 equiv), DBU (12 μL, 0.082 mmol, 5% mol.) and anhydrous MgSO₄ (1.25 g,) afforded 225 mg (38%) of 6b as a yellow solid. m.p.: 62–64 °C. ¹H NMR (CDCl₃): δ 7.8–6.9 (m, 9*H*_{arom}), 6.09 (s, 2H, OCH₂O), 5.24 (d, 1H, *J* = 17 Hz, CH₂OThp), 5.01 (d, 1H, *J* = 17 Hz, CH₂OThp), 4.9 (m, 1H, CH), 4.60 (s, 2H, PhCH₂S), 4.0–3.9 (m, 1H, CH₂), 3.7–3.6 (m, 1H, CH₂), 1.9–1.7 (m, 6H, 3CH₂). ¹³C NMR (CDCl₃): δ 168.3, 160.1, 155.8, 150.4, 148.5, 146.5, 137.2, 130.7 (8s, 8C_{arom}), 129.1, 128.5, 127.4, 122.7, 108.4, 107.8, 102.7 (7d, 9CH_{arom}), 101.7 (t, CH₂), 99.14 (d, CH), 62.6, 62.5, 35.6, 30.2, 25.1, 19.3 (6t, 6CH₂). MS (FAB⁺) *m*/*z*: 477 ([M+1]⁺, 100). Anal. Calcd. for C₂₅H₂₄N₄O₄S: C, 63.01; H, 5.08; N, 11.76; S, 6.73. Found: C, 63.22; H, 4.84; N, 11.92; S, 6.50%.

Synthesis of (5-benzo[1,3]dioxol-5-yl-2-benzylsulfanyl-1,2,4-triazolo[1,5-*a***]pyri midin-7-yl)-methanol (7). A solution of 5-benzo[1,3]dioxol-5-yl-2-benzylsulfanyl-7-(tetrahydro-pyran-2-yloxymethyl)-1,2,4-triazolo[1,5-***a***]pyrimidine 6b** (99 mg, 0.21 mmol) in (AcOH:THF:H₂O) (4:2:1) (4 mL/mmol, 1 mL) was heated at 45 °C for 19 h. The organic

solvent was eliminated under reduced pressure, was added saturated solution of NaHCO₃ (3 mL/mmol) and the aqueous layer was extracted with AcOEt (3×3 mL/mmol). The combined organic layers were dried over MgSO₄, the solvent was evaporated and the resulting residue purified by flash-chromatography (n-hexane:AcOEt) afforded 64 mg (77%) of 7 as a yellow solid. m.p.: 175–177 °C. IR (KBr): v 3199 (br., OH). ¹H NMR (DMSO-*d*₆): δ 7.9–7.2 (m, 9*H*_{arom}), 6.25 (s, 2H, OCH₂O), 6.17 (t, 1H, *J* = 5.5 Hz, CH₂OH), 5.05 (d, 2H, *J* = 5.5 Hz, CH₂OH), 4.63 (s, 2H, PhCH₂S). ¹³C NMR (DMSO-*d*₆): δ 166.6, 159.3, 155.2, 150.5, 150.1, 148.3, 137.7, 130.2 (8s, 8C_{arom}), 128.9, 128.4, 127.2, 122.7, 108.7, 107.0, 102.5 (7d, 9CH_{arom}), 101.9, 57.7, 34.4 (3t, 3CH₂). MS (FAB⁺) *m*/*z*: 393 ([M+1]⁺, 100). Anal. Calcd. for C₂₀H₁₆N₄O₃S: C, 61.21; H, 4.11; N, 14.28; S, 8.17. Found: C, 60.94; H, 4.36; N, 14.09; S, 8.39%.

Method A for the preparation of the alkoxypyrimidines 9a and 9e. Mitsunobu reaction.

A solution of DIAD (1.2 equiv) in dry THF (1 mL/mmol) is added dropwise to a solution of Ph_3P (1.2 equiv), the appropriate 2-benzylsulfanylpyrimidinone **8a-b** (1 equiv) and different alcohols in tetrahydrofuran (2 mL/mmol) at room temperature. The reaction mixture was stirred at room temperature until total disappearance of **8a-b** (TLC monitoring). The solvent was evaporated until dryness and the crude product adsorbed over silica purified by flash chromatography (n-hexane:EtOAc).

2-Benzylsulfanyl-4-isopropoxy-6-phenyl-pyrimidine (9a). According to the general procedure described above, the reaction between **8b** (500 mg, 1.70 mmol), TPP (675 mg, 2.55 mmol), and DIAD (0.50 mL, 2.55 mmol) in dry THF (6 mL), afforded 526 mg (92%) of **9a** isolated as colorless solid after 2 h. m.p.: 81–82 °C. ¹H NMR (CDCl₃): δ 8.1–8.0 (s, 2*H*_{arom}), 7.7–7.2 (m, 8*H*_{arom}), 6.77 (s, 1H, *H*_{pyrim}), 5.46 (hept, 1H, *J* = 6.2 Hz, *CH*(CH₃)₂), 4.54 (s, 2H, PhCH₂S), 1.38 (d, 6H, *J* = 6.2 Hz, CH(CH₃)₂). ¹³C NMR (CDCl₃): δ 170.8, 169.4, 164.6 (3s, 3*C*_{pyrim}), 138.0, 136.8 (2s, 2*C*_{arom}), 130.5, 128.8, 128.7, 128.5, 128.4, 127.0 (6d, 10CH_{arom}), 99.7 (d, CH_{pyrim}), 69.5 (d, CH), 35.4 (t, CH₂), 21.9 (q, 2CH₃). MS (EI) *m*/*z*: 336 ([M]⁻⁺, 90). Anal. Calcd. for C₂₀H₂₀N₂OS: C, 71.40; H, 5.99; N, 8.33; S, 9.53. Found: C, 71.51; H, 6.17; N, 8.14; S, 9.25%.

[1-Benzyl-2-(2-benzylsulfanyl-pyrimidin-4-yloxy)-ethyl]-carbamic acid *tert*-butyl ester (9e). According to the general procedure described above, the reaction between 8a (1.00 g, 4.59 mmol), TPP (1.58 g, 5.96 mmol), N-Boc-phenylalaninol (1.50 g, 5.96 mmol) and DIAD (1.15 mL, 5.96 mmol) in dry THF (15 mL), afforded 1.47 g (71%) of 9e isolated as colorless solid after 5 h. m.p.: 108–110 °C. IR (KBr): v 3388 (br., NH), 1685 (s, C=O). ¹H NMR (CDCl₃): δ 8.30 (d, 1H, *J* = 4.8 Hz, *H*_{pyrim}), 7.5–7.4 (m, 10*H*_{arom}), 6.48 (d, 1H, *J* = 4.8 Hz, *H*_{pyrim}), 4.81 (br., 1H, NH), 4.37 (s, 2H, CH₂), 4.30 (s, 2H, CH₂), 4.25 (br., 1H, CH), 2.90 (m, 2H, PhCH₂C), 1.39 (s, 9H, (CH₃)₃). ¹³C NMR (CDCl₃): δ 171.3, 168.4 (2s, 2C_{pyrim}), 157.5 (d, CH_{pyrim}), 152.2 (s, C=O), 137.8, 137.3 (2s, 2C_{arom}), 129.3, 128.8, 128.5, 127.1, 126.6, 126.4 (6d, 10CH_{arom}), 103.7 (d, CH_{pyrim}), 79.6 (s, C), 64.1 (t, CH₂), 50.8 (d, CH), 37.8, 35.2 (2t, 2CH₂), 28.3 (q, 3CH₃). MS (FAB⁺) *m*/*z*: 452 ([M+1]⁺, 17). Anal. Calcd. for C₂₅H₂₉N₃O₃S: C, 66.49; H, 6.47; N, 9.31; S, 7.10. Found: C, 66.30; H, 6.68; N, 9.07; S, 6.08%.

Method B for the preparation of the alkoxypyrimidines 9b-d.

To a solution of the corresponding 2-benzylsulfanylpyrimidinone **8a-b** (1 equiv) in DMF (3 mL per mmol), 1.1 equiv of TMG were added. Then, the corresponding phenacyl bromide (1.1. equiv) was added dropwise. The reaction mixture was until total disappearance of **8a-b** (TLC monitoring). The solvent was evaporated until dryness and the crude product adsorbed over silica purified by flash chromatography (n-hexane:EtOAc).

2-(2-Benzylsulfanyl-pyrimidin-4-yloxy)-1-(4-chloro-phenyl)-ethanone (9b). According to the general procedure described above, the reaction between **8a** (600 mg, 2.8 mmol), (4-chloro-phenyl)-acetyl bromide (850 mg, 3.6 mmol) and TMG (0.45 mL, 3.6 mmol) in dry DMF (8 mL), afforded 856 mg (83%) of **9b** isolated as colorless solid after 6 h. m.p.: 111–112 °C. IR (KBr): v 1698 (s, C=O). ¹H NMR (CDCl₃): δ 8.34 (d, 1H, *J* = 5.6 Hz, *H*_{pyrim}), 7.9–7.3 (m, 9*H*_{arom}), 6.64 (d, 1H, *J* = 5.6 Hz, *H*_{pyrim}), 5.54 (s, 2H, CH₂O), 4.25 (s, 2H, PhCH₂S). ¹³C NMR (CDCl₃): δ 191.7 (s, C=O), 171.0, 167.6 (2s, 2C_{pyrim}), 157.8 (d, CH_{pyrim}), 140.3, 137.0,

132.5 (3s, 3C_{arom}), 129.2, 128.5, 128.4, 127.1, (4d, 9CH_{arom}), 103.8 (d, CH_{pyrim}), 67.4, 35.1 (2t, 2CH₂). MS (EI) *m*/*z*: 372 ([M+2]⁺, 19), 370 ([M]⁺, 52). Anal. Calcd. for C₁₉H₁₅ClN₂O₂S: C, 61.53; H, 4.08; N, 7.55; S, 8.65. Found: C, 61.32; H, 3.98; N, 7.76; S, 8.46%.

2-(2-Benzylsulfanyl-pyrimidin-4-yloxy)-1-(3-nitro-phenyl)-ethanone (9c). According to the general procedure described above, the reaction between **8a** (1.00 g, 4.6 mmol), (3-nitro-phenyl)-acetyl bromide (1.64 g, 5.7 mmol) and TMG (0.72 mL, 5.7 mmol) in dry DMF (15 mL), afforded 1.31 g (75%) of **9c** isolated as colorless solid after 6 h. m.p.: 97–98 °C. IR (KBr): v 1705 (s, C=O). ¹H NMR (CDCl₃): δ 8.74 (s, 1*H*_{arom}), 8.46 (d, 1*H*_{arom}, *J* = 8.2 Hz), 8.35 (d, 1H, *J* = 5.6 Hz, *H*_{pyrim}), 8.24 (d, 1*H*_{arom}, *J* = 7.8 Hz), 7.71 (t, 1*H*_{arom}), 7.3–7.2 (m, 5*H*_{arom}), 6.65 (d, 1H, *J* = 5.6 Hz, *H*_{pyrim}), 5.60 (s, 2H, CH₂O), 4.26 (s, 2H, PhCH₂S). ¹³C NMR (CDCl₃): δ 191.1 (s, *C*=O), 171.1, 167.4 (2s, 2*C*_{pyrim}), 158.0 (d, CH_{pyrim}), 103.8 (d, CH_{pyrim}), 67.5, 35.0 (2t, 2CH₂). MS (EI) *m*/*z*: 381 ([M]⁺⁺, 18). Anal. Calcd. for C₁₉H₁₅N₃O₄S: C, 59.83; H, 3.96; N, 11.02; S, 8.41. Found: C, 59.57; H, 4.11; N, 11.16; S, 8.14%.

2-(2-Benzylsulfanyl-6-phenyl-pyrimidin-4-yloxy)-1-phenyl-ethanone (9d). According to the general procedure described above, the reaction between **8b** (500 mg, 1.7 mmol), phenyl-acetyl bromide (415 mg, 2.0 mmol) and TMG (0.26 mL, 2.0 mmol) in dry DMF (5 mL), afforded 512 mg (73%) of **9d** isolated as colorless solid after 4 h. m.p.: 134–135 °C. IR (KBr): 1706 (m, C=O). ¹H NMR (CDCl₃): δ 8.1–8.0 (m, 4*H*_{arom}), 7.6–7.5 (m, 6*H*_{arom}), 7.4–7.1 (m, 5*H*_{arom}), 7.05 (s, 1H, *H*_{pyrim}), 5.67 (s, 2H, OCH₂), 4.39 (s, 2H, PhCH₂S). ¹³C NMR (CDCl₃): δ 192.9 (s, C=O), 170.7, 168.9, 165.2 (3s, 3C_{pirim}), 137.5, 136.4, 134.3 (3s, 3C_{arom}), 133.8, 130.7, 128.8, 128.7, 128.6, 128.4, 127.7, 127.1, 127.0 (9d, 15CH_{arom}), 99.1 (d, CH_{pyrim}), 67.7, 35.2 (2t, 2CH₂). MS (EI) *m*/*z*: 412 ([M]⁺, 25). Anal. Calcd. for C₂₅H₂₀N₂O₂S: C, 72.79; H, 4.89; N, 6.79; S, 7.77. Found: C, 72.57; H, 5.00; N, 6.56; S, 8.09%.

Oxidation of 9a,b to Sulfones 10a,b. General Procedure.

To a cooled (0 °C) solution of pyrimidine derivatives **9a,b** (1 equiv) in CH₂Cl₂ (5 mL per mmol), 2.5 equiv of *m*-CPBA (60% purity) were added in small portions. The mixture was then stirred at 0 °C during 2 h, then diluted with CH₂Cl₂ (20 mL per mmol) and washed with aq. satd. NaHCO₃ solution (2×5 mL per mmol) and brine (5 mL per mmol). The separated organic layer was dried (MgSO₄), filtered and evaporated to give a residue which was purified by flash chromatography using n-hexane/EtOAc.

4-Isopropoxy-6-phenyl-2-phenylmethanesulfonyl-pyrimidine (10a). According to the general procedure described above, the reaction of **9a** (450 mg, 1.34 mmol) and *m*-CPBA (1.15 g, 3.34 mmol) in CH₂Cl₂ (7 mL), afforded to **10a** (432 mg, 88%) isolated as a colorless solid. m.p.: 124–125 °C. ¹H NMR (CDCl₃): δ 8.1–8.0 (m, 2H_{arom}), 7.6–7.2 (m, 8H_{arom}), 6.99 (s, 1H, H_{pyrim}), 5.65 (hept, 1H, J = 6.0 Hz, $CH(CH_3)_2$), 4.88 (s, 2H, PhCH₂S), 1.43 (d, 6H, J = 6.2 Hz, $CH(CH_3)_2$). ¹³C NMR (CDCl₃): δ 170.9, 165.5, 164.7 (3s, 3C_{pyrim}), 135.0 (s, C_{arom}), 131.6, 131.3, 128.8, 128.7, 128.2, 127.2 (6d, 10CH_{arom}), 127.1 (s, C_{arom}), 106.0 (d, CH_{pyrim}), 71.6 (d, CH), 57.3 (t, CH₂), 21.7 (q, 2CH₃). MS (EI) *m*/*z*: 368 ([M]⁺, 67). Anal. Calcd. for C₂₀H₂₀N₂O₃S: C, 65.20; H, 5.47; N, 7.60; S, 8.70. Found: C, 65.29; H, 5.60; N, 7.48; S, 8.88%.

1-(4-Chloro-phenyl)-2-(2-phenylmethanesulfonyl-pyrimidin-4-yloxy)-ethanone (**10b**). According to the general procedure described above, the reaction of **9b** (675 mg, 1.82 mmol) and *m*-CPBA (1.31 g, 4.55 mmol) in CH₂Cl₂ (9 mL), afforded to **10b** (613 mg, 84%) isolated as a colorless solid. m.p.: 141–142 °C. IR (KBr): v 1692 (s, C=O). ¹H NMR (CDCl₃): δ 8.65 (d, 1H, *J* = 5.8 Hz, *H*_{pyrim}), 7.92 (d, 2*H*_{arom}, *J* = 6.8 Hz), 7.52 (d, 2*H*_{arom}, *J* = 6.8 Hz), 7.3–7.2 (m, 5*H*_{arom}), 7.12 (d, 1H, *J* = 5.6 Hz, *H*_{pyrim}), 5.72 (s, 2H, CH₂O), 4.54 (s, 2H, PhCH₂S). ¹³C NMR (CDCl₃): δ 190.9 (s, C=O), 169.3, 163.9 (2s, 2C_{pyrim}), 158.3 (d, CH_{pyrim}), 140.8, 132.2 (2s, 2C_{arom}), 131.0, 129.4, 129.2, 128.8, 128.7 (5d, 9CH_{arom}), 126.4 (s, C_{arom}), 111.4 (d, CH_{pyrim}), 68.3, 57.6 (2t, 2CH₂). MS (FAB⁺) *m*/*z*: 405 ([M+3]⁺, 24), 403 ([M+1]⁺, 62). Anal. Calcd. for C₁₉H₁₅ClN₂O₄S: C, 56.65; H, 3.75; N, 6.95; S, 7.96. Found: C, 56.81; H, 3.54; N, 7.18; S, 8.15%.

Ipso-substitution reaction of pyrimidinyl sulfone derivatives 10a,b.

Synthesis of 4-isopropoxy-2-phenoxy-6-phenyl-pyrimidine (11a). To a solution of phenol (55 mg, 0.57 mmol) in dioxane (2 mL), the Cs_2CO_3 (210 mg, 0.59 mmol) was added.

The reaction mixture was stirred at r.t. for 15–20 min. Then, the sulfone 10a (200 mg, 0.54 mmol) was added. After stirring at 60 °C for 3 hours, the solvent was removed *in vacuo* and the mixture was acidified with 2 N hydrochloric acid and extracted with ethyl acetate. The residue was purified using flash chromatography to give 11a (131 mg, 80%) as a colorless solid. m.p.: 68–69 °C. ¹H NMR (CDCl₃): δ 8.0–7.9 (m, 2*H*_{arom}), 7.5–7.3 (m, 8*H*_{arom}), 6.84 (s, 1H, *H*_{pyrim}), 5.28 (hept, 1H, *J* = 6.2 Hz, CH(CH₃)₂), 1.34 (d, 6H, *J* = 6.2 Hz, CH(CH₃)₂). ¹³C NMR (CDCl₃): δ 171.9, 166.4, 165.0 (3s, 3*C*_{pyrim}), 153.2, 136.5 (2s, 2*C*_{arom}), 130.6, 129.1, 128.7, 127.0, 124.8, 121.9 (6d, 10CH_{arom}), 98.6 (d, CH_{pyrim}), 69.9 (d, CH), 21.8 (q, 2CH₃). MS (EI) *m/z*: 306 ([M]⁺, 24). Anal. Calcd. for C₁₉H₁₈N₂O₂: C, 74.49; H, 5.92; N, 9.14. Found: C, 74.36; H, 5.75; N, 9.40%.

Synthesis of 1-(4-chloro-phenyl)-2-[2-(2,4-dimethoxy-phenylamino)-pyrimidin-4-yloxy]-ethanone (11b). To a solution of the sulfone 10b (125 mg, 0.31 mmol) in dioxane (2 mL), the 2,4-dimethoxyaniline (100 mg, 0.62 mmol) was added. The reaction mixture with good stirring was heated at 100 °C until total consumption of the starting material (30 h., TLC monitoring). The solvent was removed under reduced pressure and the residue purified by flash-chromatography (n-hexane:EtOAc) to give 11b (25 mg, 20%) as an orange crystalline solid. m.p.: 122–123 °C. IR (KBr): v 3244 (br., NH), 1705 (m, C=O). ¹H NMR (CDCl₃): δ 8.20 (d, 1H, *J* = 5.4 Hz, *H*_{pyrim}), 7.96 (d, 2*H*_{arom}, *J* = 8.2 Hz), 7.77 (d, 1*H*_{arom}, *J* = 8.8 Hz), 7.54 (d, 2*H*_{arom}, *J* = 8.2 Hz, NH), 5.58 (s, 2H, CH₂O), 3.83 (s, 3H, CH₃O), 3.72 (s, 3H, CH₃O). ¹³C NMR (CDCl₃): δ 192.2 (s, C=O), 168.5, 159.6 (2s, 2C_{pyrim}), 158.6 (d, CH_{pyrim}), 155.4, 149.7, 140.2, 132.9 (4s, 4C_{arom}), 129.3, 129.2 (2d, 4CH_{arom}), 121.9 (s, C_{arom}), 120.3, 102.7 (2d, 2CH_{arom}), 98.6 (d, CH_{pyrim}), 98.5 (d, CH_{arom}), 67.3 (t, CH₂), 55.6, 55.4 (2q, 2CH₃). MS (EI) *m*/*z*: 401 ([M+2]⁻⁺, 17), 399 ([M]⁻⁺, 52). Anal. Calcd. for C₂₀H₁₈ClN₃O₄: C, 60.08; H, 4.54; N, 10.51. Found: C, 60.29; H, 4.48; N, 10.32%.

Removal of the 4-isopropoxy group. Synthesis of 2-phenoxy-6-phenyl-3*H*-pyrimid in-4-one (12). The 4-isopropoxypyrimidine 11a (87 mg, 0.28 mmol) was added to a mixture of AcOH (0.6 mL) and con. H₂SO₄ (0.6 mL). The reaction mixture was stirred at 90 °C for 15 min. After cooling, the mixture was neutralized with aq. 5 *N* NaOH and extracted with CH₂Cl₂ (3 × 5 mL). The combined organic layers were washed with brine and the separated organic layer was dried (MgSO₄), filtered and eliminated under reduced pressure to afford the pure pyrimidinone 12 (59 mg, 80%) as a colorless solid. m.p.: 263–264 °C. IR (KBr): ν 3060–2750 (br., NH), 1669 (s, C=O). ¹H NMR (DMSO-*d*₆): δ 12.75 (br., NH), 7.9–7.8 (m, 2*H*_{arom}), 7.6–7.5 (m, 8*H*_{arom}), 6.74 (s, 1H, *H*_{pyrim}). ¹³C NMR (DMSO-*d*₆): δ 166.0, 161.4, 158.7 (3s, 3*C*_{pyrim}), 151.8, 135.8 (2s, 2*C*_{arom}), 130.6, 129.5, 128.8, 126.6, 125.7, 121.7 (6d, 10CH_{arom}), 102.4 (d, CH_{pyrim}). MS (EI) *m*/*z*: 264 ([M]⁻⁺, 54). Anal. Calcd. for C₁₆H₁₂N₂O₂: C, 72.72; H, 4.58; N, 10.70. Found: C, 72.47; H, 4.81; N, 10.86%.

Synthesis of [1-benzyl-2-(2-benzylsulfanyl-pyrimidin-4-yloxy)-ethylamino]-phen yl-acetic acid (13): Petasis reaction.

To a stirred solution of glyoxylic acid monohydrate (74 mg, 0.78 mmol) in dichlorome thane (5 mL) was added the primary amine **9e**, previous deprotection of Boc using standard conditions, (275 mg, 0.78 mmol), followed by phenylboronic acid (99 mg, 0.78 mmol). After the flask was purged with nitrogen and sealed, the reaction mixture was stirred vigorously at room temperature for 3 days. The resulting precipitate was isolated by filtration and washed with to dichloromethane to give the pure pyrimidinone **13** (197 mg, 52%) as a colorless solid. m.p.: 134–135 °C. IR (KBr): v 3217 (br, NH), 1713 (s, C=O). ¹H NMR (DMSO-*d*₆): δ 8.42 (d, 1H, *J* = 5.8 Hz, *H*_{pyrim}), 7.9–7.8 (m, 2H, NH + COOH), 7.5–7.2 (m, 15*H*_{arom}), 6.69 (d, 1H, *J* = 5.8 Hz, *H*_{pyrim}), 4.63 (s, 1H, CHCOOH), 4.34 (s, 2H, PhCH₂S), 4.31 (dd, 1H, *J* = 4.0 Hz, *J*′ = 11.0 Hz, OCH₂), 4.17 (dd, 1H, *J* = 6.0 Hz, *J*′ = 11.0 Hz, OCH₂), 3.15 (br., 1H, PhCH₂CH), 3.02 (dd, 1H, *J* = 4.8 Hz, *J*′ = 13.6 Hz, PhCH₂CH), 2.85 (dd, 1H, *J* = 7.5 Hz, *J*′ = 13.6 Hz, PhCH₂CH). ¹³C NMR (DMSO-*d*₆): δ 172.6, 170.0, 168.0 (3s, 3C), 158.0 (d, CH_{pyrim}), 137.8, 137.7 (2s, 3C_{arom}), 134.1, 130.0, 129.2, 128.8, 128.4, 127.8, 127.3, 127.0, 126.4 (9d, 15CH_{arom}), 104.1 (d, CH_{pyrim}), 66.9 (t, CH₂), 62.4, 55.5 (2d, 2CH), 36.4, 34.2 (2t, 2CH₂).

MS (FAB⁺) m/z: 486 ([M+1]⁺⁺, 28). Anal. Calcd. for C₂₈H₂₇N₃O₃S: C, 69.25; H, 5.60; N, 8.65; S, 6.60. Found: C, 69.52; H, 5.84; N, 8.48; S, 6.31%.

Synthesis of 2-(2-benzylsulfanyl-6-phenyl-pyrimidin-4-yloxy)-1-phenyl-ethanol (14). To a stirred and cooled (0 °C) solution of pyrimidinone **9d** (500 mg, 1.21 mmols) in MeOH (6 mL) was added NaBH₄ (165 mg, 4.24 mmols) in small portions while stirring (vigorous evolution of gas observed). Stirring was continued for 2 h at 0 °C. The solution was evaporated to dryness, and the crude residue was partitioned between EtOAc (10 mL) and aq satd NH_4Cl solution (15 mL). The organic layer was separated, washed with H_2O (5 mL), dried (MgSO₄), and evaporated to give a residue which was purified by flash chromatography using hexanes/EtOAc to afford pure (14) as a colorless solid (403 mg, 80%). m.p.: 140–141 °C. IR (KBr): 3400 (br., OH). ¹H NMR (CDCl₃): δ 8.1–8.0 (m, 2H_{aron}), 7.5–7.3 (m, $13H_{arom}$), 6.90 (s, 1H, H_{pvrim}), 5.17 (dd, 1H, J = 8.4 Hz, J' = 3.2 Hz, PhCH), 4.53 (s, 2H, PhCH₂S), 4.47 (dd, 1H, J = 11.6 Hz, J' = 8.4 Hz, CH₂O), 4.03 (dd, 1H, J = 11.4 Hz, J' = 3.2 Hz, CH_2O), 2.90 (br., 1H, OH). ¹³C NMR ($CDCI_3$): δ 170.9, 169.6, 165.1 (3s, $3C_{\text{pyrim}}$), 139.8, 137.7, 136.4 (3s, 3C_{arom}), 130.8, 128.8, 128.7, 128.6, 128.5, 128.4, 128.2, 127.1, 126.2 (9d, 15CH_{arom}), 99.1 (d, CH_{pyrim}), 72.5 (d, CH), 71.7, 35.4 (2t, 2CH₂). MS (EI) *m*/*z*: 414 ([M]⁺, 33). Anal. Calcd. for C₂₅H₂₂N₂O₂S: C, 72.44; H, 5.35; N, 6.76; S, 7.74. Found: C, 72.62; H, 5.14; N, 6.52; S, 7.46%.

3.4. Microbiological Methods

Thirty-two isolates of *M. tuberculosis* from respiratory (n = 21) and non-respiratory (n = 11) clinical samples, identified by conventional methods [109] and by DNA hybridization probe (Accuprobe[®] Gen Probe Inc., San Diego, CA, USA) were selected from the laboratory collection of Dept. Microbiología, Hospital Clínico Universitario. Valencia, Spain. Susceptibility to first-line antituberculosis drugs (ethambutol, isoniazid, rifampin and streptomycin) was tested by a fluorometric method (Bactec[®] MGIT 960, Becton-Dickinson, Franklin Lakes, NJ, USA) and by a microdilution method [109].

The organisms were grown in modified Middlebrook 7H9 broth supplemented with 10% OADC enrichment (Difco Laboratories, Franklin Lakes, NJ, USA) for seven days at 37 °C. The inoculum size was obtained by dilution of *M. tuberculosis* isolates suspensions in 7H9 broth to yield an absorbance equivalent to that of a MacFarland n° 0.5 standard.

Antimicrobial susceptibility test was performed in 96-well microplates using serial twofold microdilution in 7H9 broth. Initial drug dilutions were prepared in deionized water or, if not soluble, dimethyl sulfoxide. Subsequent twofold dilutions were performed in 150 μ L of modified 7H9 broth in the microplates to provide a final test range of 128 to 0.125 mg/L. Ten μ L of a suspension of mycobacteria were added to the wells. Plates were covered with Parafilm "M"[®] (Laboratory Film, American National CanTM, Chicago, IL, USA), and incubated for 12 days at 37 °C. Starting at day 13 of incubation, 20 μ L of Resazurina[®] (Sigma 2127, St. Louis, MO, USA) with a concentration of 250 mg/L were added to the wells, and the microplates were reincubated at 37 °C for an additional period of 48 h. MIC₅₀ and MIC₉₀ were determined as the lowest concentrations of the compounds yielding no visible changes from blue to pink [110].

4. Conclusions

New chemical scaffolds have been identified that could render new lead drugs in this field, by using easy to calculate descriptors, such as structural invariants. The only substructure common to all the selected molecules is the pyrimidine ring, and the most frequent substituent is the benzylsulfanyl in position 2. The 1,2,4-triazolo[1,5-*a*]pyrimidine system is present in structures 7 and 6*a*, which have the same substituted groups in 2 and 5. Only one pyrimidone was selected (**12**). All these structures are proposed as new base structures in order to design new combinatorial synthesis projects.

Although several action mechanisms are represented in the training group, the usefulness of the method is appreciable. One possible explanation of this fact could be that the equation retains the structural features involved in all the mechanisms considered. In the opinion of the authors, this approach is not able to find new action mechanisms, but it is possible to obtain new unexpected molecular structures acting through the known mechanisms which combine the properties of the known compounds. This feature could be useful in order to avoid problems of resistance.

Supplementary Materials: The supporting information can be downloaded at: https://www.mdpi.com/article/10.3390/ijms232315057/s1.

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