

ISSN 1015-3918



**ANKARA ÜNİVERSİTESİ
ECZACILIK FAKÜLTESİ
DERGİSİ**

**JOURNAL OF FACULTY OF PHARMACY
OF
ANKARA UNIVERSITY**

**Cilt/Vol : 27
Sayı/No : 1
Yıl /Year: 1998**



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Ankara -1998

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Sahibi: Prof.Dr. Seçkin ÖZDEN

Editör: Prof.Dr. Feyyaz ONUR

Yayın Kurulu: Prof.Dr. Feyyaz ONUR (Başkan)
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Ankara Üniversitesi Eczacılık Fakültesi dergisi yılda 2 sayı yayınlanır.

Yayımlanana yazıların sorumluluğu yazara aittir.

Bu dergi **Chemical Abstracts (CA)**, **EMBASE / Excerpta Medica** ve **Medicinal Aromatic Plants Abstracts (MAPA)** da indekslenmektedir.

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JOURNAL OF FACULTY OF PHARMACY OF
ANKARA UNIVERSITY

Published by : Prof .Dr. Seçkin ÖZDEN

Editor : Prof.Dr. Feyyaz ONUR

Editorial Board: Prof.Dr. Feyyaz ONUR (Editor-in-Chief)
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Journal of Faculty of Pharmacy of Ankara University is published in semi-annual volumes.

All the articles appeared in this journal are published on the responsibility of the author.

This journal is indexed in **Chemical Abstracts (CA), EMBASE / Excerpta Medica and Medicinal Aromatic Plants Abstracts (MAPA).**

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Ankara Üniversitesi Basımevi,
1998

İÇİNDEKİLER / CONTENTS

	Sayfa
İlkay ÖREN - Bazı Antifungal Etkili Benzoksazol ve Benzimidazollerin V. albicans'a Karşı Kantitatif Yapı-Etki İlişkileri. QSAR Of Some Antifungal Active Benzoxazoles and Benzimidazoles Against C. albicans.	1
Nilüfer TARIMCI, Dilek ERMİŞ - Preparation and In vitro Evaluation of Sustained Release suppositories of Indomethacine. Indometazinin Sürekli Etkili Supozituvuarlarının Hazırlanması ve İn vitro Değerlendirilmesi.	11
Nurgün ERDEMOĞLU, Funda BİNGÖL, Bilge ŞENER - Linaria genistifolia (L.) Miller ssp. confertiflora (Boiss.) Davis Anatomik Bir Çalışma. An Investigation on the Anatomy of <i>Linaria genistifolia</i> (L.) Miller ssp. <i>confertiflora</i> (Boiss.) Davis.	23
Özge İNAL, Altan YÜKSEL - Kontakt Lensler ve Lens Çözeltileri. Contact Lenses and Lens Solutions.	31
Serap GÜR - Mesanein Non-Adrenerjik Non-Kolinerjik Kontrolü. Non-Adrenergic, Non-Cholinergic Control of the Urinary Bladder.	51

**QSAR OF SOME ANTIFUNGAL ACTIVE BENZOXAZOLES AND
BENZIMIDAZOLES AGAINST *C. albicans***

**BAZI ANTIFUNGAL ETKİLİ BENZOKSAZOL VE BENZİMİDAZOLLERİN
C. albicans'a. KARŞI KANTİTATİF YAPI-ETKİ İLİŞKİLERİ**

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SUMMARY

In this study, QSAR analysis of some isosteric heterocyclic compounds including benzoxazole and benzimidazole derivatives were studied for the antifungal activity against Candida albicans. The Free-Wilson approach is a valuable alternative tool in QSAR analysis for evaluating the lead generation and / or optimization in drug design. Predictions for the lead optimization in this series of compounds have been described by the results obtained from Free-Wilson analysis with the determination of the activity contributions of the performed structural descriptors.

The results of Free-Wilson analysis suggest that having 2-cyclohexylethyl moiety at the 2nd position on the heterocyclic system is significant for the antifungal activity and 5-Cl group is the most favorable substituent among the others.

Key words: Benzoxazole, Benzimidazole, Antifungal activity, QSAR, Free-Wilson analysis

ÖZET

Bu çalışmada, benzoksazol ve benzimidazol halkası içeren bazı izosterik heterosiklik bileşiklerin Kantitatif Yapı-Etki ilişkileri analizi Candida albicans'a karşı gösterdikleri antifungal aktiviteleri için araştırıldı, İlaç tasarımında kılavuz bileşik oluşumunun ve /veya optimizasyonunun sağlanmasında Free-Wilson metodu, Kantitatif Yapı-Etki İlişkileri analizinde yararlı alternatif bir yöntemdir. Bu seri bileşikler içinde kılavuz bileşiğin optimizasyonu, kullanılan yapısal parametrelerin Free-Wilson analizi sonucunda elde edilen aktiviteye katkıları ile belirlenmiştir.

Free-Wilson analizi sonucunda, antifungal aktivite için heterosiklik sistemin 2. konumda 2-sikloheksizetil yapısını içermesi ve 5-Cl grubunun diğerlerine göre daha önemli olduğu saptanmıştır.

Anahtar Kelimeler: Benzoksazol, Benzimidazol, Antifungal aktivite, Kantitatif yapı-etki ilişkileri. Free-Wilson analizi

INTRODUCTION

Most fungi are completely resistant to the action of the antimicrobial drugs. Only a few substances have been discovered which exert an inhibitory effect on the fungi pathogenic for man, and most of these are relatively toxic (1). The need for more and better antifungal agents is becoming more critical because of the increasing detection of systemic mycoses in patients suffering from debilitating diseases such as neoplasia and in persons on long-term total parenteral nutrition (2). A variety of useful antifungal drugs have been developed in the last three decades. But, very few compounds have as yet been found which combine the properties required for the treatment of systemic yeast infections (3).

Recent studies led to the discovery of certain highly substituted imidazole derivatives such as miconazole and clotrimazole (4, 5) which possess good clinical activity in dermatophytoses and nonsystemic candidiasis. Unfortunately, systemic use of miconazole has been accompanied by reversible thrombocytosis and anemia and of clotrimazole by severe gastrointestinal disturbances (2).

Recently, we reported the synthesis and their *in vitro* antifungal activities of different derivatives of 5-substituted-2-cyclohexyl, 2-cyclohexylmethyl, 2-(2-cyclohexylethyl), 2-cyclopentyl, 2-cyclopentylmethyl, 2-(2-cyclopentylethyl) benzoxazoles and benzimidazoles against the fungus *Candida albicans* (6,7).

The Free-Wilson approach is a satisfactory method applied for quantitative structure-activity relationships (8-10). The basic assumption of this procedure is that within a homologous series of drugs individual segments of molecules make additive and constant contributions to biological activity. If such contributions are known, biological activity can be estimated by simple addition for all the compounds obtainable by any new combination of the segments involved.

Free-Wilson analysis can be applied to homologous series where only substituents are varied in a constant parent molecule. The selection of these substituents, however, must not meet all of the requirements of the extrathermodynamic approach. In particular, series where only a few substituents are varied in many positions can be analysed by the Free-Wilson method.

One possible way to deal with such situations is to use parameters derived directly from two-dimensional chemical structures as molecular descriptors; such descriptors will be called "structural parameters". The structures to be considered are broken down into fragments and the structural parameters then indicate the occurrence of these fragments in these structures. Fragmentation and description of molecules in structural terms can be performed in many different ways, and different approaches can be used to related structural parameters to biological activity. In the most simple case it

is assumed that substituents at certain positions (fragments) make additive and constitutive contributions to biological activity which can be expressed in terms of a linear model called the Free-Wilson model.

In this present study, QSAR analysis of some antifungal active fused heterocycles **1-20** having cyclohexyl, cyclohexylmethyl, 2-cyclohexylethyl, cyclopentyl, cyclopentylmethyl or 2-cyclopentylethyl moieties at the position 2 and bearing H, Cl, NO₂ groups at the position 5 or/and 6 given in Figure 1 against *C. albicans* were determined using the Free-Wilson method.

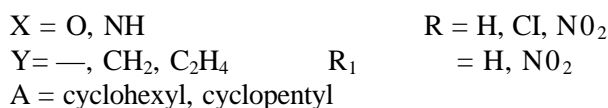
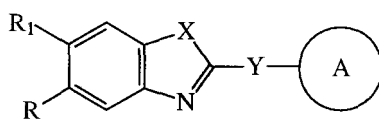


Figure 1

7

EXPERIMENTAL SECTION

The basic assumption of the Free-Wilson method is that within a set of congeners of compounds, individual segments of molecules called as molecular descriptors make additive and constant contributions to biological activity (9).

This expression is formulated as,

$$\text{Log } 1/C = \sum a_i x_i + \mu \quad \text{Eq 1}$$

Where a_i is the contribution of the i th substituent, and if the substituent is present in the molecule, x_i has a value of 1, otherwise a value of 0. The overall average activity calculated for the unsubstituted (constant) molecule is indicated as μ .

At the first step, the structure matrix given in Table 1 has been drawn up by listing the structural parameters x_i and used in Eq 1 that yielded the linear equation set of the compounds **1-20**.

At the next step, additional restrictive equations, so called symmetry conditions, have been formulated that the sum of the varying groups equaled to 0.

Symmetry equations for the sample are;

For position R,

$$9a_{11} + 8a_{12} + 3a_{13} = 0 \quad \text{Eq 2}$$

For position R₁,

$$16a_{21} + 4a_{22} = 0 \quad \text{Eq 3}$$

For position X,

$$9a_{31} + 11a_{32} = 0 \quad \text{Eq 4}$$

For position Y,

$$7a_{41} + 7a_{42} + 6a_{43} = 0 \quad \text{Eq 5}$$

For position A,

$$14a_{51} + 6a_{52} = 0 \quad \text{Eq 6}$$

a_{11} , a_{21} , a_{31} , a_{41} and a_{51} have been selected as a dependent variable at each position from Eqs 2-6.

$$a_{11} = -8/9a_{12} - 3/9a_{13} \quad \text{Eq 7}$$

$$a_{21} = -4/16a_{22} \quad \text{Eq 8}$$

$$a_{31} = -11/9a_{32} \quad \text{Eq 9}$$

$$a_{41} = -7/7a_{42} - 6/7a_{43} \quad \text{Eq 10}$$

$$a_{51} = -6/14a_{52} \quad \text{Eq 11}$$

Expression obtained from Eqs 7-11 are combined as substitutes of a_{11} , a_{21} , a_{31} , a_{41} and a_{51} in the equation set of the linear system and a correlation matrix derived. The correlation matrix given in Table 2.

Molecular descriptor values used in the Multiple Regression analysis were obtained from the correlation matrix and the observed log 1/C values have been used as dependent variable.

Besides the group contributions, the ranges of the activity contribution values of the substitution position sides were also calculated by the equations given below;

$$R = a_{12} - a_{13} \quad \text{Eq 12}$$

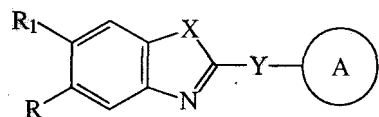
$$R_1 = a_{22} - a_{21} \quad \text{Eq 13}$$

$$X = a_{32} - a_{31} \quad \text{Eq 14}$$

$$Y = a_{43} - a_{41} \quad \text{Eq 15}$$

$$A = a_{51} - a_{52} \quad \text{Eq 16}$$

Correlation and regression analysis of this QSAR study performed by using IBM-computer working with Statgraff 2.6 Statistic Program package.

Table 1: Structure matrix of the compounds 1-20 for the Free-Wilson analysis

Comp No:	R	R ₁	X	Y	A	R			R ₁		X		Y			A		MIC mg/ml	log 1/C
						a ₁₁ -H	a ₁₂ -Cl	a ₁₃ -NO ₂	a ₂₁ -H	a ₂₂ -NO ₂	a ₃₁ -O-	a ₃₂ -NH-	a ₄₁ -	a ₄₂ -CH ₂ -	a ₄₃ -C ₂ H ₄ -	a ₅₁ cyclohexyl	a ₅₂ cyclopentyl		
1	Cl	H	O	—	cyclohexyl	1			1		1					1		50	3.6730
2	H	NO ₂	O	—	cyclohexyl					1	1					1		50	3.6702
3	Cl	NO ₂	O	—	cyclohexyl	1				1	1					1		50	3.7299
4	H	H	NH	—	cyclohexyl							1				1		25	3.9031
5	NO ₂	H	NH	—	cyclohexyl			1			1					1		50	3.6902
6	H	H	NH	—	cyclopentyl						1						1	50	3.5416
7	Cl	H	NH	—	cyclopentyl	1					1					1		50	3.6201
8	NO ₂	H	O	CH ₂	cyclohexyl			1		1					1		50	3.7160	
9	H	NO ₂	O	CH ₂	cyclohexyl				1	1	1				1		25	3.9965	
10	Cl	NO ₂	O	CH ₂	cyclohexyl	1				1	1				1		25	4.0531	
11	H	H	NH	CH ₂	cyclohexyl							1			1		25	3.9325	
12	Cl	H	NH	CH ₂	cyclohexyl	1					1				1		25	3.9974	
13	NO ₂	H	NH	CH ₂	cyclohexyl			1			1				1		50	3.7143	
14	H	H	NH	CH ₂	cyclopentyl						1					1	50	3.5752	
15	H	H	O	C ₂ H ₄	cyclohexyl					1					1		12.5	4.2629	
16	Cl	H	O	C ₂ H ₄	cyclohexyl	1				1					1		12.5	4.3239	
17	H	H	NH	C ₂ H ₄	cyclohexyl						1				1		12.5	4.2610	
18	Cl	H	O	C ₂ H ₄	cyclopentyl	1				1						1	25	3.9777	
19	H	H	NH	C ₂ H ₄	cyclopentyl						1					1	25	3.9074	
20	Cl	H	NH	C ₂ H ₄	cyclopentyl	1				1						1	25	3.9759	
Sum.						9	8	3	16	4	9	11	7	7	6	14	6		

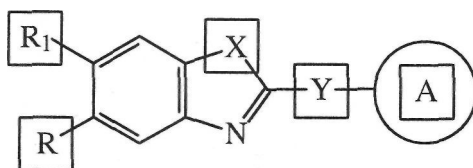
Table 2: Correlation matrix derived from symmetry equations

Comp. No:	R	R ₁	X	Y	A	R		R ₁	X	Y		A	MIC mg/ml	log 1/C
						a ₁₂ -Cl	-NO ₂			a ₂₂ -NO ₂	a ₃₂ -NH-			
1	Cl	H	0	—	cyclohexyl	1	0	-0.25	-1.22	-1	-0.857	-0.43	50	3.6730
2	H	NO ₂	0	—	cyclohexyl	-0.89	-0.33	1	-1.22	-1	-0.857	-0.43	50	3.6702
3	Cl	NO ₂	0	—	cyclohexyl	1	0	1	-1.22	-1	-0.857	-0.43	50	3.7299
4	H	H	NH	—	cyclohexyl	-0.89	-0.33	-0.25	1	-1	-0.857	-0.43	25	3.9031
5	NO ₂	H	NH	—	cyclohexyl	0	1	-0.25	1	-1	-0.857	-0.43	50	3.6902
6	H	H	NH	—	cyclopentyl	-0.89	-0.33	-0.25	1	-1	-0.857	1	50	3.5416
7	Cl	H	NH	—	cyclopentyl	1	0	-0.25	1	-1	-0.857	1	50	3.6201
8	NO ₂	H	0	CH ₂	cyclohexyl	0	1	-0.25	-1.22		0	-0.43	50	3.7160
9	H	NO ₂	0	CH ₂	cyclohexyl	-0.89	-0.33	1	-1.22		0	-0.43	25	3.9965
10	Cl	NO ₂	0	CH ₂	cyclohexyl	1	0	1	-1.22		0	-0.43	25	4.0531
11	H	H	NH	CH ₂	cyclohexyl	-0.89	-0.33	-0.25	1		0	-0.43	25	3.9325
12	Cl	H	NH	CH ₂	cyclohexyl	1	0	-0.25	1		0	-0.43	25	3.9974
13	NO ₂	H	NH	CH ₂	cyclohexyl	0	1	-0.25	1		0	-0.43	50	3.7143
14	H	H	NH	CH ₂	cyclopentyl	-0.89	-0.33	-0.25	1		0	1	50	3.5752
15	H	H	0	C ₂ H ₄	cyclohexyl	-0.89	-0.33	-0.25	-1.22	0		-0.43	12.5	4.2629
16	Cl	H	0	C ₂ H ₄	cyclohexyl	1	0	-0.25	-1.22	0		-0.43	12.5	4.3239
17	H	H	NH	C ₂ H ₄	cyclohexyl	-0.89	-0.33	-0.25	1	0		-0.43	12.5	4.2610
18	Cl	H	0	C ₂ H ₄	cyclopentyl	1	0	-0.25	-1.22	0		1	25	3.9777
19	H	H	NH	C ₂ H ₄	cyclopentyl	-0.89	-0.33	-0.25	1	0		1	25	3.9074
20	Cl	H	NH	C ₂ H ₄	cyclopentyl	1	0	-0.25	1	0		1	25	3.9759
Sum.						0	0	0	0	0	0	0		

RESULTS AND CONCLUSION

QSAR analysis for the lead optimization predictions obtained from Free-Wilson approach in these set of compounds **1-20** indicating group activity contributions, the ranges of position side values was summarized in Table 3. Observed, calculated and residual log $1/C$ values of the compounds **1-20** were also given in Table 4.

Table 3: Group activity contributions, ranges of position side values and the statistical data obtained from Free-Wilson analysis against *C. albicans*



Position	Group	Activity	contributions
		Position	Group
R		0.231	
	-H		0.008
	-Cl		0.056
R ₁	-NO ₂	0.016	-0.175
	-H		-0.003
	-NO ₂		0.013
X		0.064	
	-O-		-0.035
Y	-NH-	0.476	0.029
	—		-0.196
A	-CH ₂ -	0.320	-0.044
	-C ₂ H ₄ -		0.280
	cyclohexyl		0.096
	cyclopentyl		-0.224
Constant contribution (μ)		3.8761	
Number of compounds (n)		20	
Square of correlation coefficient (R^2)		0.9164	
Standart deviation (s)		0.0852	
F value (F)		18.80	
F table value (F) (%99)		4.64	

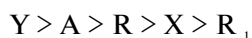
Table 4: Observed, calculated and residual log 1/C values of the compounds 1-20 obtained from Free-Wilson analysis

Comp No:	R	R ₁	X	Y	A	MIC µg/ml	Observed log 1/C	Calculated log 1/C	Residual
1	Cl	H	O	—	cyclohexyl	50	3.6730	3.7942	-0.1212
2	H	NO ₂	O	—	cyclohexyl	50	3.6702	3.7618	-0.0916
3	Cl	NO ₂	O	—	cyclohexyl	50	3.7299	3.8104	-0.0805
4	H	H	NH	—	cyclohexyl	25	3.9031	3.8090	0.0940
5	NO ₂	H	NH	—	cyclohexyl	50	3.6902	3.6262	0.0639
6	H	H	NH	—	cyclopentyl	50	3.5416	3.4888	0.0527
7	Cl	H	NH	—	cyclopentyl	50	3.6201	3.5374	0.0826
8	NO ₂	H	O	CH ₂	cyclohexyl	50	3.7160	3.7154	0.0005
9	H	NO ₂	O	CH ₂	cyclohexyl	25	3.9965	3.9144	0.0820
10	Cl	NO ₂	O	CH ₂	cyclohexyl	25	4.0531	3.9630	0.0900
11	H	H	NH	CH ₂	cyclohexyl	25	3.9325	3.9616	-0.0291
12	Cl	H	NH	CH ₂	cyclohexyl	25	3.9974	4.0102	-0.0128
13	NO ₂	H	NH	CH ₂	cyclohexyl	50	3.7143	3.7788	-0.0645
14	H	H	NH	CH ₂	cyclopentyl	50	3.5752	3.6414	-0.0662
15	H	H	O	C ₂ H ₄	cyclohexyl	12.5	4.2629	4.2222	0.0406
16	Cl	H	O	C ₂ H ₄	cyclohexyl	12.5	4.3239	4.2708	0.0530
17	H	H	NH	C ₂ H ₄	cyclohexyl	12.5	4.2610	4.2855	-0.0245
18	Cl	H	O	C ₂ H ₄	cyclopentyl	25	3.9777	3.9506	0.0270
19	H	H	NH	C ₂ H ₄	cyclopentyl	25	3.9074	3.9654	-0.0580
20	Cl	H	NH	C ₂ H ₄	cyclopentyl	25	3.9759	4.0140	-0.0381

Table 3 reveals that the lead optimization predictions obtained from Free-Wilson approach in these set of compounds 1-20 can be summarized as follows;

The constant contribution descriptor μ is decisive for the antifungal activity as a leading segment at the molecule.

Additive contribution range of the position side values of the groups are found significant as the following order of;



Nonpolar group like Cl at position R is causing an increase in the activity.

It is shown that the position 2 is important for the antifungal activity against *C. albicans* and substituting this position with 2-cyclohexylethyl, enhances the potency.

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Başvuru tarihi : 17.12.1997

Kabul tarihi : 05.03.1998