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## QSRT STUDIES OF 2-(5-ARYL-1, 3, 4-OXADIAZOL-2-YL)-N-(5-METHYLISOXAZOL-3-YL)-ACETAMIDE DERIVATIVES AS AN ANTIMICROBIAL AGENTS

Srinivas Marri<sup>1,2</sup>, Ramu Kakkerla<sup>\*3</sup> and M. P. S. Murali Krishna<sup>4</sup>

Department of Chemistry<sup>1</sup>, JNTU, Kakinada - 533003, Andhra Pradesh, India.

Department of Chemistry<sup>2</sup>, Siddhartha Degree & P. G. College, Narsampet, Warangal - 506132, Telangana, India.

Department of Chemistry<sup>3</sup>, Satavahana University, Karimnagar - 505001, Telangana, India.

Department of Chemistry<sup>4</sup>, Andhra Polytechnic College, Kakinada - 533003, Andhra Pradesh, India.

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### Correspondence to Author:

**Dr. Ramu Kakkerla**

Assistant Professor,  
Department of Chemistry, Satavahana  
University, Karimnagar - 505001,  
Telangana, India.

**E-mail:** kakkerla2001@yahoo.co.in

**ABSTRACT:** Quantitative structure relationship techniques (QSRT) is among the most widely used computational technology for analog-based drug design. A molecular modeling approach using theoretical, computational analysis through chem sketch ACD lab online software was done for determination of probable antimicrobial activity. To develop a pharmacophoric model for inhibition, QSRT parameters PCP, ADME, and toxicity has been used. BCF (Bioconcentration factor), adsorption coefficient, log P, log D values, plasma protein binding, P-gp inhibition, AMES test, hERG inhibition, estrogen receptor, and Lipinski's type properties have been calculated. From a data set of 11 analogs, it has been concluded that all compounds are non-bio-accumulative, non-endocrine disruptors, non hERG inhibitors, genotoxic and within Lipinski's criteria. None of the compounds are practically ionizable at various body pH values, and all are zwitterionic compounds. From these data and *in-vitro* antimicrobial data compounds, 5c and 5f can be exploited for the formulation of bactericide and fungicide with a slight modification in their structure.

**INTRODUCTION:** Besides biological reactivity against a target, which is the primary requirement, there are so many essential properties and characteristics that are mandatory to be possessed by a molecule to be considered as a drug. The use of computational methods for designing molecules with a desired activity, reactivity or property has been a growing area in chemistry and medicine.

In this direction QSRT analysis has been performed considering some important activity properties like BCF (Bioconcentration factor), adsorption coefficient, partition coefficients, probable plasma binding property, p-gp inhibiting characteristics, non genotoxicity, AMES test possibilities, hERG inhibiting probabilities, endocrinal disruptor tendency, and drug's pharmacokinetics like adsorption, distribution, metabolism and elimination in the human body, and can be inferred from Lipinski's rule molecular properties.

1, 3, 4-Oxadiazoles are of great practical significance, which is primarily due to their large number of uses, in the most diverse areas, such as in drug synthesis, scintillation materials, in the

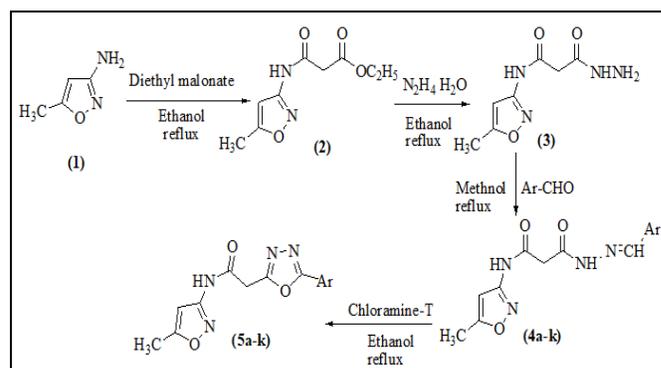
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production of polymers and dyes, and uses in photography as light screening agents. 1, 3, 4-Oxadiazole derivatives are reported to exhibit antibacterial<sup>1</sup>, antifungal<sup>2</sup>, analgesic and anti-inflammatory<sup>3</sup>, anticonvulsant<sup>4</sup>, anticoagulant<sup>5</sup>, and anticancer activity<sup>6, 7</sup>. A large number of synthetic derivatives of this molecule are well documented in the literature. Similarly, isoxazole derivatives are found to possess a wide variety of biological activities<sup>8, 9, 10, 11, 12</sup>.

Literature survey indicated that the compounds bearing the isoxazole moiety are endowed with various types of biological activities. The 1, 3, 4-oxadiazoles are heterocycles of current interest due to their broad-spectrum pharmacological activity. Therefore, in the current investigation, the QSRT studies of isoxazolyl 1, 3, 4-oxadiazoles have been undertaken to correlating antimicrobial activity. In this paper, we present QSRT studies of 2-(5-aryl-1, 3, 4-oxadiazol-2-yl)-N-(5-methyl isoxazol-3-yl)-acetamides.

**MATERIALS AND METHODS:** Several isoxazolyl 1,3,4-oxadiazoles have been synthesized and characterized by spectral analysis **Scheme I** and evaluated for antibacterial **Table 1** and antifungal activity<sup>13</sup> **Table 2**. The structures are drawn and submitted for theoretical, computational

analysis using through chem sketch ACD-lab online software<sup>14</sup> analyzed QSRT parameters viz., BCF, adsorption coefficient<sup>15</sup>, log P, log D values, Plasma protein binding<sup>16</sup>, P-gp inhibition<sup>17</sup>, AMES test<sup>18, 19</sup>, hERG inhibition<sup>20</sup>, estrogen receptor<sup>21, 22</sup> genotoxicity<sup>23, 24</sup> and Lipinski's type properties<sup>25</sup>. The data is tabulated and examined for analysis between the structural aspects of molecular activity. Under the light of changes of structural aspects of the molecule, the trends in the molecules are evaluated. This will help in designing new molecules with expected behaviors.



**SCHEME I: SYNTHESIS OF 2-(5-aryl-1, 3, 4-oxadiazol-2-yl)-N-(5-methylisoxazol-3-yl)-acetamides.** 4 & 5 Ar = a: C<sub>6</sub>H<sub>5</sub>; b: 4-ClC<sub>6</sub>H<sub>4</sub>; c: 4-OCH<sub>3</sub>C<sub>6</sub>H<sub>4</sub>; d: 4-NO<sub>2</sub>C<sub>6</sub>H<sub>4</sub>; e: 3-OCH<sub>3</sub>C<sub>6</sub>H<sub>4</sub>; f: 3-BrC<sub>6</sub>H<sub>4</sub>; g: 2-OH C<sub>6</sub>H<sub>4</sub>; h: 2-OH,3-OCH<sub>3</sub>C<sub>6</sub>H<sub>3</sub>; i: 3,4,5-(OCH<sub>3</sub>)<sub>3</sub>C<sub>6</sub>H<sub>2</sub>; j: 3,4-(OCH<sub>3</sub>)<sub>2</sub>C<sub>6</sub>H<sub>3</sub>; k: 2-CH<sub>3</sub>C<sub>6</sub>H<sub>4</sub>

**TABLE 1: ANTIBACTERIAL ACTIVITY OF N-(5-methylisoxazol-3-yl)-2-(5-phenyl-1,3,4-oxadiazol-2-yl) acetamides<sup>13</sup> (5a-k)**

Compound	Minimum Inhibitory Concentration (MIC) <sup>a,b</sup>					
	Bacterial strains					
	<i>P. aeruginosa</i>	<i>K. aerogenes</i>	<i>C. violaceum</i>	<i>B. subtilis</i>	<i>B. sphaericus</i>	<i>S. aureus</i>
5a	20	18	18	19	17	15
5b	11	11	8	9	8	7
5c	17	15	13	16	18	14
5d	10	8	7	6	8	6
5e	10	16	15	15	13	14
5f	13	11	14	11	12	10
5g	18	14	16	16	15	15
5h	19	16	20	17	18	21
5i	21	18	21	19	17	18
5j	23	20	22	18	16	15
5k	22	23	20	18	17	16
Ciprofloxacin	30	25	25	20	20	25

<sup>a</sup>Negative control (acetone)- no activity, <sup>b</sup>Concentration in µg/mL

**TABLE 2: ANTIFUNGAL ACTIVITY OF N-(5-methylisoxazol-3-yl)-2-(5-phenyl-1,3,4-oxadiazol-2-yl) acetamides<sup>13</sup> (5a-k)**

Compound	Zone of Inhibition mm <sup>a,b</sup>				
	Fungal strains				
	<i>A. niger</i>	<i>C. tropicalis</i>	<i>R. oryzae</i>	<i>F. moniliforme</i>	<i>C. lunata</i>
5a	52.5	46.5	51.0	41.2	50.5
5b	69.1	70.1	72.5	69.8	65.5
5c	56.0	57.0	58.0	61.0	59.5
5d	75.0	77.2	80.5	73.2	69.5
5e	48.0	51.0	48.5	53.2	57.2

5f	63.2	65.0	72.5	60.5	73.5
5g	55.8	58.3	59.1	60.0	61.1
5h	47.2	42.5	40.2	38.5	31.5
5i	39.0	38.3	31.2	37.0	51.0
5j	51.0	55.2	49.8	48.1	59.0
5k	45.5	60.1	55.3	48.1	39.5
Clotrimazole	26.5	30.6	33.5	25.5	35.8

<sup>a</sup>Negative control (acetone)- no activity, <sup>b</sup>Concentration 100 µg/mL

## RESULTS AND DISCUSSION:

**QSRT Studies:** In this paper, an attempt has been made to analyze 2-(5-aryl-1,3,4-oxadiazol-2-yl)-N-(5-methylisoxazol-3-yl)-acetamide derivatives (5a-5k) through the application of the quantitative structure relationship techniques (QSRT).

**QSPR Studies:** Table 3 shows the properties computed through quantitative structure-property relationship algorithm of ACD software. Very marginal variance is observed among all physical parameters.

**QSAR Study:** BCF of 5a-k structures is found to have a constant unity value concerning the change

in pH. In the case of the adsorption coefficient, a consistent value unity has been maintained between pH range 1-9, besides that, it tends to approach a constant value, as shown in Table 4. Since, BCF is <1000, 5a -5k compounds are non bioaccumulative.

Therefore, there is very less chance to be incorporated into biological food chains from the environment. Since the bioconcentration factor is greater than 1; it is indicative of a hydrophobic or lipophilic chemical. In the case of 5a-5k derivatives as they have unity value indicating very low hydrophilic nature, but not having of lipophobic or hydrophilic nature.

TABLE 3: GENERAL PHYSICAL PROPERTIES OF COMPOUNDS 5a-5k

Compound	Molar Refractivity	Molar Volume	Parachor	Index of Refraction	Surface Tension	Density	Polarizability
5a	73.30	210.0	584.5	1.615	59.9	1.353	29.05
5b	78.19	221.9	620.4	1.622	61.0	1.435	30.99
5c	79.98	234.0	641.2	1.599	56.3	1.342	31.70
5d	79.84	221.8	640.0	1.639	69.2	1.484	31.65
5e	79.98	234.0	641.2	1.599	56.3	1.342	31.70
5f	80.99	226.2	635.0	1.634	62.1	1.605	32.10
5g	75.18	208.4	599.5	1.640	68.4	1.440	29.80
5h	81.86	232.4	656.2	1.621	63.5	1.420	32.45
5i	93.33	282.0	754.5	1.576	51.2	1.327	37.00
5j	86.65	258.0	697.9	1.586	53.5	1.334	34.35
5k	78.12	226.3	622.2	1.606	57.1	1.318	30.97

TABLE 4: BCF, ADSORPTION COEFFICIENT, log P AND log D VALUES OF COMPOUNDS 5a-5k

Compound	BCF BIO		log P		log D				
	BCF	K <sup>OC</sup>	log P	Reliability	pH 1.6	pH 4.6	pH 6.5	pH 7.4	pH 8.2
5a	1	20.8C	1.71	0.65M	1.71	1.71	1.71	1.71	1.71
5b	1	54.3C	2.12	0.52M	2.12	2.12	2.12	2.12	2.12
5c	1	25.7C	1.73	0.48B	1.73	1.73	1.73	1.73	1.73
5d	1	19.8C	1.49	0.59M	1.48	1.49	1.49	1.49	1.49
5e	1	25.6C	1.73	0.48B	1.73	1.73	1.73	1.73	1.73
5f	1	70.5C	2.35	0.5M	2.35	2.35	2.35	2.35	2.35
5g	1	29.2C	1.3	0.55M	1.3	1.3	1.3	1.3	1.3
5h	1	27.6	0.81	0.51M	0.81	0.81	0.81	0.81	0.81
5i	1	26.4C	1.38	0.56M	1.38	1.38	1.38	1.38	1.38
5j	1	28.1C	0.96	0.47B	0.96	0.96	0.96	0.96	0.96
5k	1	37.1C	1.73	0.52M	1.73	1.73	1.73	1.73	1.73

The log P and log D computed values are found to be moderately reliable with reliability parameter between 0.56-0.65. For all compounds except for

5c, 5e, and 5j, whose reliabilities are found to be borderline. log D values for all of the derivatives at pH 1.7, 4.6, 6.5, 7.4 and 8 appears to be same for

all the compounds (5a–5k) indicating that at all those pH s no practical ionization of molecule occurs **Table 4**. Partition coefficient values are found to be in-between 0.86 to 2.83. log P, log D, adsorption coefficient, and BCF increased when halogen groups are present. By halogen substitution molecule's log P, log D, and aqueous solubility

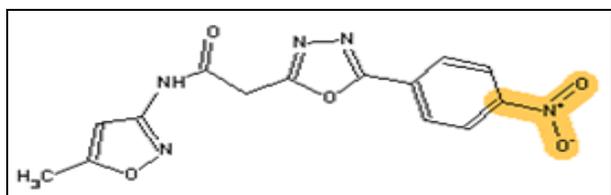
decreased. Presence of two methoxyl on aryl groups shown a decrease in log P value in 5j derivative on comparison with the un-substituted molecule, whereas the presence of either single methoxyl group or three methoxyl groups on aryl structural unit is showing a decreasing tendency.

**TABLE 5: PLASMA PROTEIN BINDING AND P-gp INHIBITION POSSIBILITIES WITH RELIABILITIES OF COMPOUNDS 5a-5k**

Compound	DBP			P-gp inhibition				
	PPB (%)	Reliability I	log K <sup>HSA</sup>	Reliability II	Probability	Reliability	K <sub>i</sub>	Reliability
5a	96.54	0.42B	4.32	0.4B	0.08	0.31B	0.01	0.56M
5b	97.45	0.52M	4.63	0.38B	0.1	0.2NR	0.01	0.46B
5c	95.28	0.49B	4.32	0.4B	0.11	0.2NR	0.02	0.54M
5d	95.73	0.41B	4.16	0.38B	0.1	0.2NR	0.02	0.49B
5e	95.28	0.49B	4.32	0.4B	0.11	0.2NR	0.02	0.54M
5f	96.44	0.43B	4.46	0.39B	0.1	0.21NR	0.02	0.48B
5g	93.77	0.49B	4.14	0.39B	0.05	0.13NR	0.01	0.53M
5h	90.78	0.4B	4.06	0.23NR	0.06	0.14NR	0	0.49B
5i	93.6	0.45B	4.34	0.34B	0.13	0.25NR	0.03	0.53M
5j	93.65	0.49B	4.25	0.35B	0.13	0.22NR	0.03	0.55M
5k	95.28	0.48B	4.33	0.4B	0.11	0.21NR	0.02	0.51M

% PPB of 5b is found to be moderately reliable in respect of plasma protein binding possibility and borderline for all other 5 series compounds. 5b has the highest value of 97.45, and 5h has the least value. The variation is very narrow for all as 94±4. No molecule is found to have a high probability of 0.8 or above reliability in respect to plasma protein binding parameter. log K<sup>HSA</sup> values of 5a-5k compounds are very narrowly distributed so synergetic effect on the availability of molecule concerning others is very less significant. Presence of 5b can enhance the percentage of 5h in blood as the difference in their K<sub>i</sub> value is significant to consider. All (5a-5k) are Zwitterionic compounds. These drugs are likely to bind to the majority of plasma proteins.

Further one may work except that all may bind to human serum albumin, alpha1-acid glycoprotein, and albumin although in most cases the binding is not extensive. In compound 5d nitroarenes exhibit genotoxic activity after metabolic transformation to hydroxylamines **Fig. 1**.



**FIG. 1: GENOTOXICITY OF COMPOUND 5d**

In this series, a constant value of 0.1 has been observing for most of them. From this one can conclude that there will be a least binding tendency concerning P-glycoprotein, 5a, 5h and 5g are having 0.05 order of binding aptitude that is 10 fold less than other members of the same series.

Since probability of potent P-gp inhibition (K<sub>i</sub><1 micro-M): 0.01, Reliability: Moderate (RI = 0.56 to 0.49) and McGowan's volume is found to be less than 200 cm<sup>3</sup>/mol with high reliability for 5a, so it can classify as non-inhibitor, 5g, and 5h will be Probably non-inhibitor as they are weak hydrophilic acid or amphiprotic compound, acid pK<sub>a</sub> > 5, MW < 360, AB/logP < 2.5, with high reliability. In cases of 5b, 5c, 5d, 5e, 5f, 5i, 5j and 5k no prediction can be made unless data for similar compounds be available to predict through the use for probabilistic estimation<sup>23, 26</sup> **Table 5**.

All other molecules of this series have AMES test probabilities in the range of 0.12 to 0.16 with non-reliable values of 0.17 or 0.18, except 5d with a value of 0.68 with borderline reliability 0.38. So it may show carcinogen activity. hERG inhibiting probabilities are found to be reliable, which is borderline except 5c. Still, one may except non inhibiting character as their values estimated appear to be very low, (the range is 0.1-0.2). Only 5g has 0.09 value.

They are expected to have log RBA less than -3 and  $K^i$  less than 10  $\mu$ M. Relative binding affinity values are within the criteria of expectation, so all the molecules of consideration do not tend to bind to alpha estrogens receptor **Table 6**. All

compounds are found to be within the Lipinski's criteria. It has found to be a topological surface area well below 145 and 142 (borderline value) in case of 5d; it may be associated with a reduction of genotoxicity by oxadiazole ring formation **Table 7**.

**TABLE 6: AMES TEST PROBABILITIES AND ESTROGEN RECEPTOR PROBABILITIES WITH RELIABILITIES OF COMPOUNDS 5a-5k**

Compound	AMES test		hERg Inhibitors		Endocrine Disruptions	
	Probability	Reliability	Probability	Reliability	Log RBA>-3	Log RBA>0
5a	0.13	0.2NR	0.12	0.22NR	0.2	0
5b	0.14	0.17NR	0.28	0.27NR	0.33	0
5c	0.16	0.17NR	0.13	0.23NR	0.21	0
5d	0.68	0.38B	0.1	0.19NR	0.18	0
5e	0.16	0.17NR	0.13	0.23NR	0.24	0
5f	0.14	0.18NR	0.15	0.18NR	0.36	0
5g	0.14	0.17NR	0.09	0.18NR	0.23	0
5h	0.17	0.17NR	0.13	0.17NR	0.24	0
5i	0.1	0.19NR	0.14	0.17NR	0.11	0
5j	0.12	0.18NR	0.13	0.19NR	0.19	0
5k	0.13	0.18NR	0.12	0.23NR	0.31	0

**TABLE 7: LIPINSKI'S TYPE PROPERTIES OF COMPOUNDS 5a-5k**

Compound	Lipinski's type properties				
	M. Wt.	HBD	HBA	TPSA	RB
5a	284.27	1	7	94.05	4
5b	318.71	1	7	94.05	4
5c	314.3	1	8	103.28	5
5d	329.27	1	10	142.88	5
5e	314.3	1	8	103.28	5
5f	363.17	1	7	94.05	4
5g	300.27	2	8	114.28	4
5h	330.29	2	9	123.51	5
5i	374.35	1	10	121.74	7
5j	344.32	1	9	112.51	6
5k	298.3	1	7	94.05	4

**CONCLUSION:** QSAR and QSPR studies for 5a-5k were presented. All the Compounds (5a-5l) are found to be promising non bio-accumulative, nonendocrine disruptors, non hERg inhibitors, and within Lipinski criteria. None of the compounds are practically ionizable at various body pH values. All the compounds (5a-5k) are zwitterionic compounds. It can be concluded that compounds 5a-5k may exhibit potential antimicrobial activity based on QSRT studies. QSRT studies may help in designing the pharmacophore modes to identify more potent isoxazolyl oxadiazole as an antimicrobial agent, which will be useful for the formulation of bacteriocide and fungicide.

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