



Thermal, Spectral, Antibacterial And Antioxidant Studies On Inclusion Complexes Of 2- (Thiazolidinone-2') Imino-3-Allyl 5-Arylidene-4-Thiazolidinone With β -Cyclodextrin)

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Abstract: Novel Arylidene derivatives of 2-(Thiazolidinone-2') imino-3-allyl-4-thiazolidinone were synthesized. To enhance solubility, β -cyclodextrin was utilized to form inclusion complexes. The presence of mixes and inclusion complexes was confirmed by analyzing the analytical data and spectrum properties. We used a cup-and-plate experiment to measure the antibacterial and antioxidant effects against *Bacillus subtilis*, *Escherichia coli*, and *Proteus vulgaris*. When compared to the parent chemicals, the inclusion complexes showed far better antioxidant and antibacterial capabilities.

Index Terms - β -cyclodextrin, *Escherichia*, antibacterial activity, inclusion complexes, cup plate method

1. INTRODUCTION

Yearly reports on synthetic heterocyclic structures encompass a variety of both well-established and newly discovered frameworks. These novel compounds, known for their diverse biological activities, have garnered significant attention from researchers. Chemists are delving into their complete synthesis and potential in chemotherapy. Utilizing advanced analytical and spectroscopic tools, scientists are able to analyze structural modifications and assignments based on synthetic efforts. Thiazolidinone serves as a pivotal pharmacophore, with its five-membered ring containing both sulfur and nitrogen. The research on thiazolidinones is expansive due to their unique biological activities, exemplified by antimicrobial properties and various others such as substituted thiazolidinones have been discovered.¹⁻², antioxidant³, anti-HIV⁴, antihistaminic⁵, anti-convulsant⁶⁻⁷, anti-inflammatory⁸⁻¹⁰ effects. In this current study, an assortment of arylidene derivatives of 2-(Thiazolidinone-2') imino-3-allyl-4-thiazolidinone have been synthesized from asymmetrical thiourea. However, the challenge of poor solubility in polar media may impede their bioavailability and efficacy as medicinal agents. To address this issue, a promising strategy involves enhancing the bioavailability of these compounds by forming inclusion complexes with cyclodextrin¹¹, a hydrophobic cavity. Cyclodextrins are commonly used as model hosts to improve the solubility of water-insoluble guests, making them more easily assimilated in water. Among cyclodextrins, β -cyclodextrin stands out as a cost-effective and safe option for creating inclusion complexes^[12-13]. Thus, this study aims to synthesize several derivatives of 2-(Thiazolidinone-2') imino-3-allyl-5-arylidene-4-thiazolidinone from asymmetrical thiourea¹⁴ and investigate their complexation with β -cyclodextrin to enhance their properties and potential applications. The analysis of the compound's logical and spectral data has led to the characterization of the compound and its inclusion complexes. In addition, studies have been conducted to determine if the drug's antioxidant and antibacterial effects are amplified during complex formation.

precipitate was filtered through a G-4 crucible after being rinsed with distilled water. We used Higuchi-Connors^[18] solubility assay to find out how the compounds dissolved in water. Using a rotary flask shaker, each chemical was subjected to room temperature shaking for 48 hours in a series of conical flasks. Filtration using Whatman-42 filter paper was followed by analysis of the solutions using a UV-visible spectrophotometer in the 200-400 nm range. The maximum absorbance (λ -max) was then compared to the β -cyclodextrin concentrations. Benesi Hilderbrand relation determines inclusion complexes' thermodynamic stability constant K_T . $1/\Delta A = 1/\Delta \epsilon + 1/K_T [Guest]_0 / \Delta \epsilon$. $[\beta\text{-CD}]_0$ "Where ΔA is change in absorbance, $\Delta \epsilon$ is change in molar extension coefficient, $[Guest]_0$ is concentration of compound in inclusion complex and $[\beta\text{-CD}]_0$ is molar concentration of β -CD.

2.4 Antibacterial evaluation

Cup-plate method examined compound antibacterial activity.^[19] The test chemicals were dissolved in 500 μ g/ml DMSO. Escherichia coli (MTCC 40), Proteus vulgaris (MTCC 87), and Bacillus subtilis (MTCC 321) were introduced to a 100 ml sterile nutritive stock and cultured at 37 $^\circ$ C for 24 hours. It was McFarland who standardized the thickness of bacterial suspensions. Agar plates were sterile-injected with test organisms prior to creating 6-mm-diameter wells. Micropipette-encapsulated drug and test synthetics (500 g/ml) were chilled to 8-10 $^\circ$ C in the chiller for diffusion into the medium. The Petri dishes were heated to 37 $^\circ$ C for 18 to 24 hours after an initial two-hour cold incubation phase. The venire scale was used to determine the size of inhibition zones in petri dishes. Analyzing the difference between the test compounds' and the reference medication's zone of inhibition allowed researchers to account for the results (Antibiotic medication). Tagashira and Ohtake^[20] suggested assessing cell reinforcement activity by DPPH (2, 2-diphenyl-1-picrylhydrazyl) radical seeking. Figure 4.22 depicts the DPPH's no-cost, no-obligation extreme search experiment of synthetic blends and composite structures. To form 1, 1-diphenyl-2-picryl hydrazine, cell reinforcement, with its speedy hydrogen gift, combines with DPPH, a nitrogen-focused extremist with a specific absorbance at 517 nm. Cellular reinforcement rummaging is suggested by staining. At concentrations of 100 and 500 g/mL, we tested the cell reinforcement capacity of the mixtures and candidate structures, respectively. Synthetic and incorporation structures' cell reinforcement or extreme rummaging action (RSA) is shown in Table 4.4.

3. RESULTS AND DISCUSSION

Limited solubility in polar solvents can restrict the pharmacological effectiveness of thiazolidinone derivatives. However, by creating inclusion complexes with cyclodextrins, these molecules can improve their solubility and therapeutic potential. Both Table-I and Table-II include analytical and spectral data for the compounds that were produced and their inclusion complexes, correspondingly. The compound forms are distinguished based on their spectral characteristics and the presence of elemental sulfur, with infrared and sulfur content measurements aligning with expectations. Their melting points provide evidence of the presence of chemical inclusion complexes with β -cyclodextrin. The compounds A, B, C, and D have melting points of 175 $^\circ$ C, 158 $^\circ$ C, 145 $^\circ$ C, and 92 $^\circ$ C, respectively. In contrast, the inclusion complexes of these compounds have melting temperatures of 180 $^\circ$ C, 162 $^\circ$ C, 152 $^\circ$ C, and 98 $^\circ$ C, respectively. The fact that the molecules need more heat to be released from the β -cyclodextrin cavity suggests that the complex has been successfully formed, which explains the discrepancy in melting points. The solubility of compounds in a β -cyclodextrin solution in the aqueous phase showed a linear dependence on concentration. The slopes of the graphs were all less than 1, suggesting a possible 1:1 stoichiometry for these compounds. For each inclusion complex, the Benesi-Hilderband relation was used to find its thermodynamic stability constant (K_T).^[21] Through the plot of $1/A$ vs $1/[\beta\text{-CD}]_0$ for the substances, strong linear relationships were observed, enabling the calculation of K_T values for each complex. In Table I, we can see that the K_T values for the inclusion complexes of compounds with β -cyclodextrin were 588.69, 725.5, 575.9, and 805.4 M^{-1} for A, B, C, and D, specifically. The inclusion complexes shown stability as a consequence of host-guest interactions such hydrophobic contacts and van der Waals forces, since these findings were within the usual range of 100-1000 M^{-1} .^[20,21] The calculation of the corresponding thermodynamic parameters, including the ΔG value at 298K, ($\Delta G = -2.303RT \log K$) was facilitated by determining the stability constant (K_T values) at various temperatures assuming a 1:1 stoichiometry.

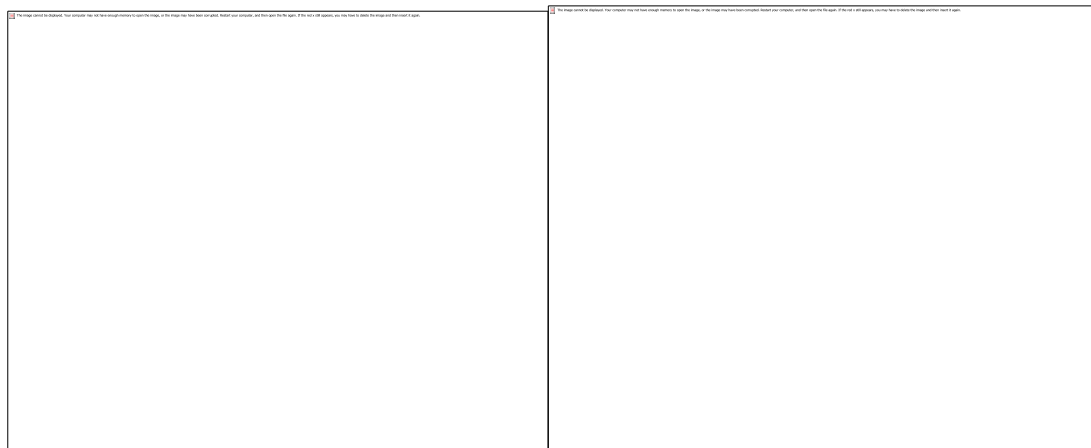


Figure: 1 A

Figure: 1B

The obtained activation energies for all inclusion complexes in Table-I are negative, indicating that the process of inclusion is thermodynamically favorable. “Analysis of the IR data in Table II of Compound A shows characteristic bonds such as C-S, C-C, C-N, C=O, C-H, and N-H at 680, 126, 159, 170, 296, and 3109 cm^{-1} , respectively. Similarly, the IR spectra of I.C. A display distinctive absorptions at 745, 124, 1597, 172, 2965, and 312 cm^{-1} , confirming the presence of C-S, C-C, C-N, C=O, C-H, and N-H bonds. The IR analyses of complexes B, C, and D along with their inclusion complexes, also exhibit characteristic absorption frequencies in accordance with the spectral data. In all cases, a reduction in the IR frequencies of the C=O bonds is observed, accompanied by broadening, weakening, and smoothing of the peaks. Conversely, upon formation of the inclusion complex, the N-H stretching vibration frequency shifts to a higher wavenumber. Weak interactions including hydrogen bonding, van der Waals forces, and hydrophobic contacts between the host and guest molecules are formed as a result of these alterations, which show that the compounds have been effectively contained within the β -cyclodextrin cavity.^[19-20]

3.1 Antibacterial Study

The results of the antibacterial tests indicate that the zones of inhibition against three bacterial strains (*Proteus vulgaris*, *Escherichia coli*, and *Bacillus subtilis*) increase in size from simple to complex structures for both the combinations and their corresponding inclusion complexes (Table III, Figs. 2, 3, and 4). The drugs are effective against sensitive bacteria, while showing no activity against resistant strains. Additionally, the compounds tend to exhibit greater stability and bioavailability compared to the pure medication. By encapsulating the drug molecules within the hydrophobic core of β -cyclodextrin, and allowing the hydrophilic surface to interact with the surrounding environment, there is an enhancement in bioavailability and sensitivity to bacteria.

3.2 Antioxidant results

In Table-IV, we can see the antioxidant activities, which are shown as radical scavenging activity (RSA), for the compounds and their inclusion complexes at concentrations of 100 µg/ml (Figure-6) and 500 µg/ml (Figure-5). Improvements in solubility and bioavailability are responsible for the chemicals' increased RSA when inclusion complexes are formed. An improvement in bio-accessibility enhances the antioxidant activity of the compounds. All compounds exhibit dose-dependent DPPH radical scavenging activity.

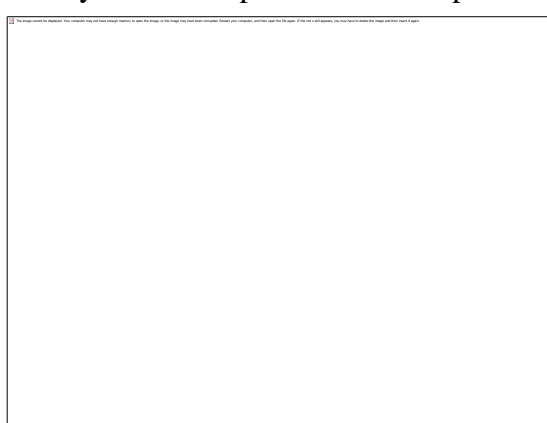


Figure: 2



Figure: 3

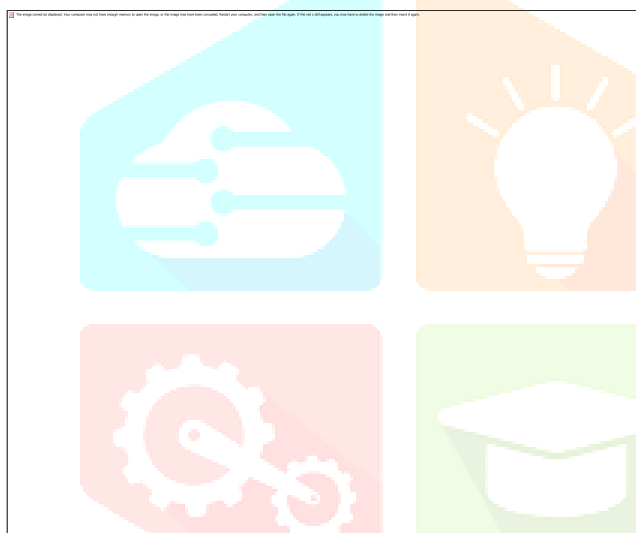


Figure: 4

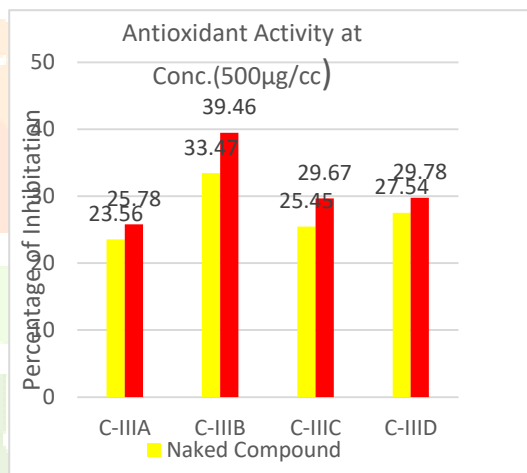


Figure: 5

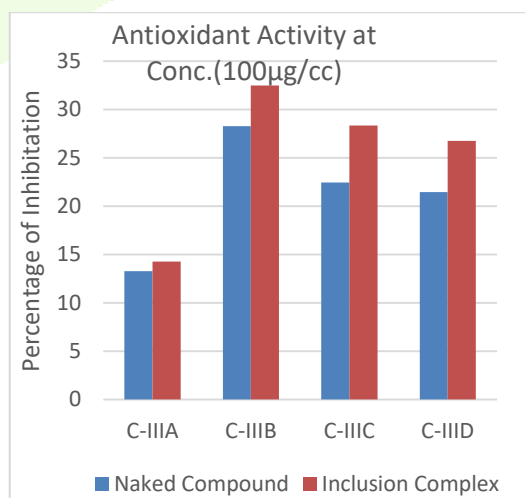


Figure: 6

Table: 1 Physical properties and thermodynamics parameter of the synthesized compounds and their complexes

Sl	Compound/Complex	Ar.	Color	M.P. in °C	% of yield	K	ΔG (kj/mol)
1	A	Phenyl	Brownish red	175	60	-	
2	I.C.A		Light yellowish	208	40	588.69	-15.804
3	B	p-Cl-C ₆ H ₄	Deep yellow	158	65		
4	I.C. B		Brownish yellow	162	41	725.5	-16.33
5	C	p-NO ₂ C ₆ H ₄	Dull brown	145	55		
6	I.C- C		Pale yellow	152	45	575.9	-15.737
7	D	p-OCH ₃ -C ₆ H ₄	Dull white	92	54		
8	I.C.- D		White	98	39	805.4	-16.58

Table 2: Spectral data of synthesized compounds and their inclusions

Sl no.	Compound/Complex	Ar.	IR(KBr) cm ⁻¹	¹ H NMR	UV _{λmax}
1	Compound A	Phenyl	742.59(C-S str), 1492.60(C=C Str), 1589.34(C=N str), 1645.28(C=O str), 3194.12(N-H str)	¹ H NMR (CDCl ₃) : δ 6.81-8.23 (d, 6H, Ar-H), 4.23(s,1H,C-NH), 7.58(s,1H,C-H), 7.34-7.61 (m,8H, Ar-H) ¹ H NMR (CDCl ₃)	275
2	I.C.A		746.45(C-Sstr), 1494.83(C=C str), 1581.83(C=N str), 1714.12(C=O str), 3224.34(N-H str)	δ 6.12-7.81 (d, 6H, Ar-H), 3.83 (s,1H,C-NH), 7.11(s,1H,CH), 6.82-7.24 (m,8H, Ar-H)	278
3	C.B	p-Cl-C ₆ H ₄	692.44(C-C str), 744.52(C-S str), 1487.12(C=C Str), 1583.56(C=N str), 2916.37(Ar-Hstr) 1701.22, 1645.28(C=O str), 3197.89(N-H str)	¹ H NMR (CDCl ₃) : δ 6.95-8.6 (d, 6H, Ar-H), 4.4(s,1H,CNH), 7.80(s,1H,C-H), 7.56-7.9 (t,8H, Ar-H)	265
4	I.C. B		692.44(C-Cstr), 746.45(C-S str), 1489.05(C=C Str), 1539.20(C=Nstr), 1714.72(C=O str), 3030.17(ArHstr), 3325.28, 3194.12(N-H str)	¹ H NMR (CDCl ₃): δ 6.3 -7.45 (d, 6H, Ar-H), 3.9 (s,1H,C-NH), 7.25(s,1H,CH), 6.9-7.3 (t,8H, Ar-H)	267

5	Compound C	P – NO ₂ C ₆ H ₄	742.59(C-Sstr), 850.61, 1338.60(N=Ostr), 1616.35(C=C Str), 1450.26(C=N str), 1696.36 (C=Ostr), 2916.37(Ar-HStr), 3196.50 (N-Hstr)	¹ H NMR (CDCl ₃) : δ 6.72-7.44 (d, 6H, Ar-H), 4.34(s,1H,C-NH),7.91(s,1H,C-H),7.51-7.92(m,8H, Ar-H)	296
6	I.C- C		690.52(C-Cl str), 748.38(C-S str), 1157.92(C-N str), 1494.83(C=C Str), 1597.60(C=N str), 2922.16(ArHstr)	¹ H NMR (CDCl ₃): δ 6.33-7.12 (d, 6H, Ar-H), 4.14 (s,1H,C-NH),7.33(s,1H,CH),7.11-7.54 (m,8H, Ar-H)	299
7	Compound D	p-OCH ₃ -C ₆ H ₄	748.38(C-Sstr), 1425.40(C=Nstr), 1494.83(C=C Str), 1593.20(C=N str), 1712.97 (C=o), 3062.96(Ar-HStr)	¹ H NMR (CDCl ₃) : δ 6.6-8.5 (d, 6H, Ar-H), 4.3(s,1H,CNH),7.65(s,1H,C-H),7.3-7.6 (t,8H, Ar-H),3.95 (s,3H,OCH ₃)	271
8	I.C.- D		748.38(C-Sstr), 1417.60(C-NStr), 1456.26(C=Cstr), 1508.33(C=N str), 1699.29, 1637.56(C=O str), 3331.07(N-H str)	¹ H NMR (CDCl ₃): δ 6.1-7.8 (d, 6H, Ar-H), 3.7 (s,1H,C-NH),7.5(s,1H,CH),6.6-7.1 (t,8H, Ar-H)3.65 (s,3H,OCH ₃)	275

Table-3: Antibacterial activity of the synthesized compounds and their complexes

Compound/Complex	Diameter of zone of inhibition(mm)		
	<i>Proteus vulgaris</i>	<i>Escherichia coli</i>	<i>Bacillus subtilis</i>
Comp- A	7	8	10
I.C.A.	10	11	13
Comp-B	15	10	12
I.C. B	18	17	17
Comp-C	14	12	16
I.C.C	18	16	18
Comp-D	12	8	10
I.C. D	16	13	14

Table 4: Antioxidant activity of synthesized compounds and their inclusions”

Compound	Percentage of Inhabitation			
	Conc.(100µg/cc)		Conc.(500µg/cc)	
	Naked Compound	Inclusion Complex	Naked Compound	Inclusion Complex
A	13.27	14.27	23.56	25.78
B	28.25	32.47	33.47	39.46
C	22.43	28.32	25.45	29.67

D	21.46	26.74	27.54	29.78
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4. CONCLUSION:

In summary, the findings indicate that the solubility, bioavailability, thermodynamic stability, antibacterial, and antioxidant activities of the synthesized compounds, specifically 2-(thiazolidinone-2) imino-3-allyl 5-arylidene-4-thiazolidinone, are significantly enhanced through the formation of inclusion complexes with β -cyclodextrin.

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