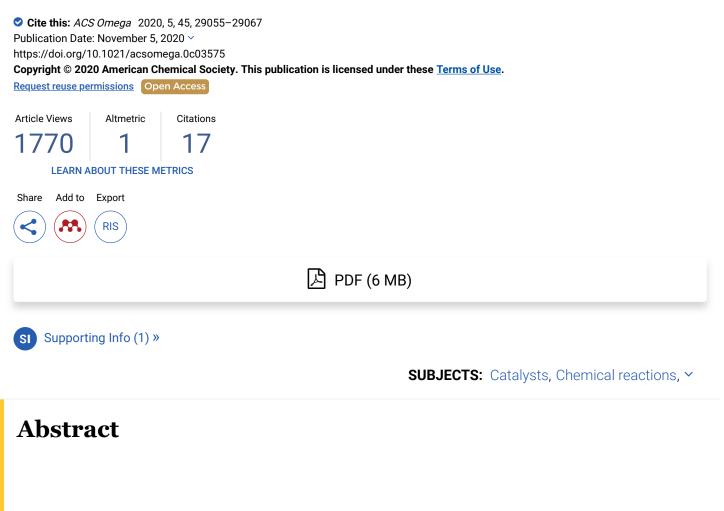


Rapid Construction of Substituted Dihydrothiophene Ureidoformamides at Room Temperature Using Diisopropyl Ethyl Ammonium Acetate: A Green Perspective

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An economic, sustainable, and straightforward environmentally friendly synthesis of highly diversified polyfunctional dihydrothiophenes is successfully achieved via diisopropyl ethyl ammonium acetate as a room-temperature ionic liquid. Multicomponent synthesis contains domino processes; the benefit of this present protocol is highlighted by its readily available starting materials, superior functional group tolerance, purity of synthesized compounds was checked by high-performance liquid chromatography results in up to 99.7% purity for the synthesized compounds, reaction mass efficiency, effective mass yield, and excellent atom economy. In addition, a series of 2-(N-carbamoyl acetamide)-substituted 2,3-dihydrothiophene analogs were synthesized, and selected samples were chosen for testing their in vitro antibacterial and antifungal activities. Furthermore, a molecular docking study against sterol 14α -demethylase could provide valuable insight into the mechanism of antifungal action providing an opportunity for structure-based lead optimization.

Introduction

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Over the last few decades, ionic liquids (ILs) have received an excessive deal of attention in an extensive range of different areas, attainment from material synthesis to an alternative reaction as well as separation science media. (1-6) ILs can be considered as green substitutes for organic solvents. They differ from molecular solvents by their "structure and organization" and their idiomatic ionic character, which can lead to precise effects, building them multipurpose and tunable materials. (7-11) Also, ILs have been successfully used in many multicomponent reactions. (12-17) Therefore, ILs may be an ideal medium of multicomponent domino-type reactions without pressure-tight equipment and organic solvents.

Multicomponent reactions (MCRs) offer a chance for the combination of three or more flexible and straightforward building blocks in a one-pot operation, constructing complex structures by the

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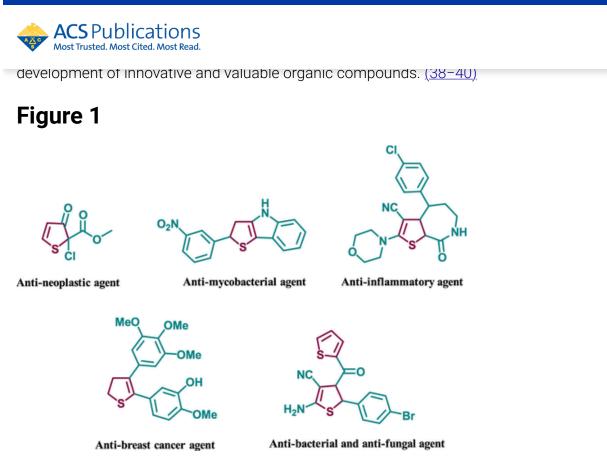
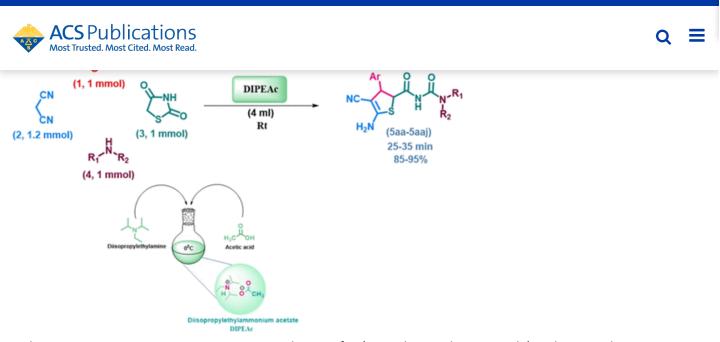


Figure 1. Chemical structures of some biologically active dihydrothiophenes.

In the past few decades, the careful study of the previously closely reported literature discloses pioneering work for the one-pot reaction of 1,3-thiazolidinedione, malononitrile, aldehyde, and aniline. In 2009, Sun et al. have reported the synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophenes by domino reactions of 1,3-thiazolidinedione in acetonitrile using organic amines as a catalyst. (<u>41</u>) In 2011, Shi et al. reported an improved synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene derivatives under ultrasound irradiation. (<u>42</u>) Again in 2011, Lu et al. reported an efficient synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene derivatives under ultrasound irradiation. (<u>42</u>) Again in 2011, Lu et al. reported an efficient synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene derivatives by domino reactions of 1,3-thiazolidinedione under catalyst-free conditions. (<u>43</u>) Whereas, in 2019, Kordnezhadian et al. carried out DBU-functionalized MCM-41-coated nanosized hematite (DBU-F-MCM-41-CNSH): a new magnetically separable basic nanocatalyst for the diastereoselective one-pot four-component synthesis of dihydrothiophene ureidoformamides. (<u>44</u>) Each of these reported procedures has its own merit, but all suffer from limitation of the synthesis to only a narrow range of dihydrothiophene ureidoformamides, a difficulty to isolate products or long reaction time, and harsh reaction conditions.

Thus we wished to explore a more "eco-friendly" and expeditious protocol for the construction of

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Scheme 1. One-Pot Four-Component Synthesis of 2-(*N*-Carbamoyl acetamide)-Substituted 2,3-Dihydrothiophenes (**5aa**–**5aaj**) in the Presence of DIPEAc as a New Room-Temperature Ionic Liquid

Results and Discussion

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Chemistry

Nowadays, the selection of a suitable reaction medium is of crucial importance for the successful organic synthesis. Ionic liquids (ILs) have become quite a popular solvent and accelerant in the chemical community due to their new generations, which contain catalytic functional groups, especially acidic (3,45) and basic groups. (46–49) Among the ILs, room-temperature ionic liquids (RTILs) have been the prominence of multiple current scientific surveys because of the properties being non-explosive, non-volatile, easy to handle, thermally robust, and nearly-zero vapor pressures. (50) Our group has elaborated on the applications and syntheses of the ILs, (51) and we have wished to utilize them as a reaction media and a homogeneous catalyst support for a high-yield, rapid, and recyclable process for the synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene.

Just following our goal to determine an innovative, sustainable, green, and synthetic path to choose the superior experimental conditions for the production of 2-(*N*-carbamoyl acetamide)-substituted 2,3dihydrothiophene, a model reaction was started with an equimolar (1:1:1:1) involving the fourcomponent domino reaction between 4-chlorobenzaldehyde (**1a**) (1.0 mmol), malononitrile (**2**) (1.0 mmol), 1,3-thiazolidinedione (**3**) (1.0 mmol), and aniline (**4a**) (1.0 mmol). <u>Scheme 2</u> is selected to elevate catalysts and conditions, reaction media, and catalyst dosage. As shown in <u>Table 1</u>, it was seen that when the reaction was performed in the absence of a catalyst in water acetonitrile and ethanol, the

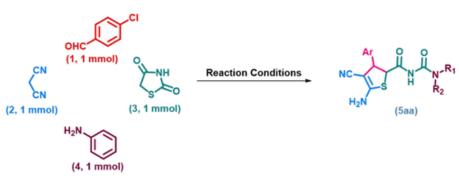
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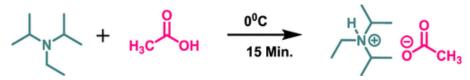
experiments and found the optimal reaction temperature (room temperature) (<u>Table 1</u>, entries 14–16, 18–22, and 28–38). In the presence of DIPEAc, this time, we have been seeing a new spot on the thinlayer chromatographic plate along with some unreacted 4-chlorobenzaldehyde (**1a**) and malononitrile (**2**) (<u>Table 1</u>, entries 13–22). Structural elucidation from ¹H NMR, ¹³C NMR, and liquid chromatographymass spectrometry (LC–MS) analysis of the isolated product gave an uncompleted Knoevenagel adduct because of the hygroscopic nature of malononitrile; then, we changed the equimolar ratio of malononitrile and optimized the model reaction, and we observed that the respective molar proportion for the model reaction between 4-chloro-benzaldehyde (**1a**) and malononitrile (**2**) changed from 1:1:1:1 to 1:1.2:1:1. This means that during the reaction, the molar proportion of malononitrile (**2**, 1.2 mmol) required was some more that of 4-chlorobenzaldehyde (**1**, 1 mmol), thiazolidinedione (**3**, 1 mmol), and aniline (**4**, 1 mmol). Employing this aspect, compound **5a** was isolated in 94% yield after only 30 min at room temperature. The model reaction in water using phase transfer catalysts was found to form the desired **5a** in less yields. Therefore, it can be thought that DIPEAc (<u>Scheme 3</u>) is a superior and green catalyst and solvent compared to the others shown in <u>Table 1</u>.

Scheme 2



Scheme 2. One-Pot Four-Component Reaction of 4-Chlorobenzaldehyde (**1a**, 1 mmol), Malononitrile (**2**, 1 mmol), 1,3-Thiazolidinedione (**3**, 1 mmol) and Aniline (**4a**, 1 mmol) under Different Conditions

Scheme 3



Scheme 3. Synthesis of Diisopropyl Ethyl Ammonium Acetate (DIPEAc)

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3		(1:1:1:1)	CHCN
4	Cs ₂ CO ₃	(1:1:1:1)	EtOH
5	p-TSA	(1:1:1:1)	H ₂ O
6	β-CD	(1:1:1)	H ₂ O
7	СТАВ	(1:1:1:1)	H ₂ O
8	Et ₃ N	(1:1:1:1)	EtOH
9	[Et ₃ NH][HSO ₄]	(1:1:1:1)	[Et ₃ NH]
10	[DBUH][OAc]	(1:1:1:1)	[DBUH][
11	ChCl:2urea	(1:1:1:1)	ChCl:2u
12	ChCl:2ZnCl ₂	(1:1:1:1)	ChCI:2Z
13	PEG-400	(1:1:1:1)	PEG-40
14	PEG-400	(1:1:1:1)	H ₂ O
15	pyrrolidine ammonium acetate	(1:1:1:1)	pyrrolid
16	piperidine ammonium acetate	(1:1:1:1)	piperidi
17	triethyl ethylammonium acetate	(1:1:1:1)	triethyl
18	[bmim]Br	(1:1:1:1)	[bmim][
19	DABCO	(1:1:1:1)	H ₂ O
20	dicationic ionic liquid	(1:1:1:1)	dicatior
21	DIPEAc	(1:1:1:1)	H ₂ 0

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25	DIPEAc	(1:1.2:1:1)	EtOH
26	DIPEAc	(1:1.2:1:1)	DIPEAc

^aReaction conditions: 4-chloro benzaldehyde (1.0 mmol), malononitrile (1.2 mmol), 1,3-thiazolidinedione (1.0 mmol), and aniline (1.0 mmol) in a medium (4 mL) stirred at room temperature. ^bIsolated yields b: no condensation. Bold values are for highlighting the good result.

From this optimization of catalysts, we can conclude that DIPEAc gives a better yield of the desired product. After the effective optimization of the catalyst, we also investigated the catalyst loading in the model reaction DIPEAc for the synthesis of **5a**. To assure the volume of the DIPEAc, the model reaction was investigated by a number of trials by changing the catalyst loading from 1 to 5 mL; as the amount of DIPEAc rises progressively, there is a steady growth noticed in the product yield. Then, 4 mL of DIPEAc furnished the **5a** in 94% yield at room temperature (<u>Table 2</u>, entry 1) and completed an excellent renovation of the reactants into the product in room-temperature DIPEAc at room temperature for 30 min (HPLC = 98.81%). Further, an increase in the amount of DIPEAc showed no significant difference in reaction time and product yield. The model reaction was performed without any solvent and catalyst. The very small amount of the product was obtained after a prolonged period (<u>Table 2</u>, entry 1).

Table 2. Solvent Impact on the Reaction for Synthesis of 2-(N-Carbamoyl Acetamide)-Substituted2,3dihydrothiophene Derivatives in Room Temperature DIPEAc^a

entry	DIPEAc	temp. (°C)	solvent	time (min)	yield (%) ^b
1	0	room temp.		24 h	Trace
2	1 mL	room temp.		30	60
3	2 mL	room temp.		30	75
4	3 mL	room temp.		30	80
5	4 mL	room temp.		30	94
6	5 mL	room temp.		45	95
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11	20 mol %	reflux	DMF	320	52
12	20 mol %	reflux	CH ₂ Cl ₂	320	50

^aReaction conditions: 4-chloro benzaldehyde (1.0 mmol), malononitrile (1.2 mmol), 1,3-thiazolidinedione (1.0 mmol), and aniline (1.0 mmol) in a medium (4 mL) stirred at room temperature. ^bIsolated yields b: no condensation. Bold values are for highlighting the good result.

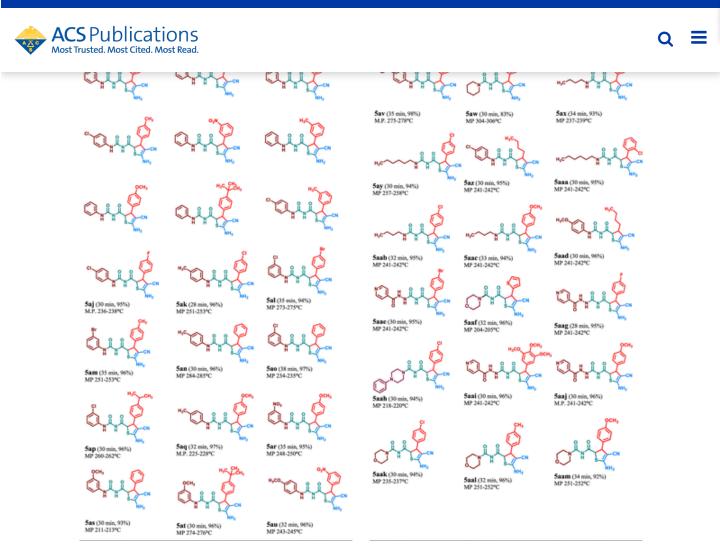
Moreover, the performance of DIPEAc was examined by using 20 mol % DIPEAc in different solvents (Table 2, entries 7–12). In ethanol, we detect a preferred reaction medium, and no sticky reaction mass was formed. Still, the isolated reaction yield (73%, E-factor = 0.62) was not satisfactory than that of DIPEAc (94%, E-factor = 0.25), the reaction conversion with a readily superior yield. Meanwhile, in acetonitrile, water, methanol, CH_2Cl_2 , DMF, and DCM reaction, output had lesser yields at reflux temperature. Not any of the solvents persist the superiority of yield and time over the non-solvent condition. Hence, the solventless condition was considered as excellent for the environmental suitability and cost.

As far as sustainable development is concerned, the fastest reaction time and highest efficiency for the formation of dihydrothiophene ureidoformamides were found at room temperature by using 4 mL of DIPEAc. Having excellent conditions in hand, the flexibility of the procedure was investigated for the synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophenes (**5aa–5aam**). We afterward examined the substrate scope by the reaction of different substituted aromatic/heteroaromatic aldehyde and aniline-subsumed methoxy, cyano, methyl, halogen (–Cl, –F, and –Br), hydroxyl, and nitro groups were used. The outcomes of all transformations carried out under these conditions are shown in <u>Table 3</u>. Aniline- and aldehyde-containing electron-withdrawing group like –NO₂ and electron-donating groups like –OMe and –Me on the aromatic ring was suited with this transformation, and comparable 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophenes (**5aa–5aam**) were achieved in good to high yields. To our enchantment, halogen-substituted 2-benzylidene malononitrile gave the products with high yields (**5ab**, **5ak**, and **5ao**). Furthermore, sterically crowded di- and trisubstituted benzaldehyde gave the desired outcomes in high yields (**5ah**, **5aai**, and **5aae**). The aliphatic aldehydes and heteroaryl aldehydes also remained under the current reaction medium without any trouble (**5ay**, **5aad**, and **5aah**).

Table 3. Substrate Scope for Synthesis of 2-(*N*-Carbamoyl acetamide)-Substituted 2,3-Dihydrothiophene Derivatives in Room-Temperature DIPEAc^a



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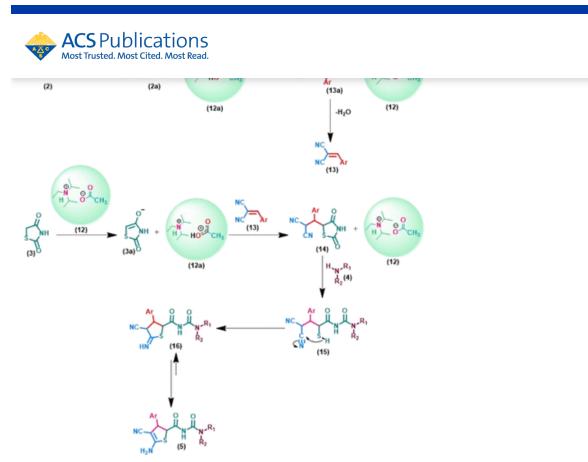
^aReaction conditions: 4-chloro benzaldehyde (1.0 mmol), malononitrile (1.2 mmol), 1,3-thiazolidinedione (1.0 mmol), and aniline (1.0 mmol) in a medium (4 mL) stirred at room temperature. Isolated yields in parentheses: no condensation.

To describe the mechanism of this one-pot multicomponent reaction of aryl aldehyde (1), malononitrile (2), 1,3-thiazolidinedione (3), amines (4), and formation of 2- (N-carbamoyl acetamide)-substituted 2,3dihydrothiophenes (**5aa–5aam**) in the presence of room-temperature DIPEAc (12), we suggest a plausible reaction mechanism, which is demonstrated in <u>Scheme 4</u>.

Scheme 4

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Scheme 4. Plausible Mechanism for the Formation of 2-(*N*-Carbamoyl acetamide)-Substituted 2,3-Dihydrothiophenes

The first step is the DIPEAc (12) catalyzing the synthesis of arylidene malononitrile (13) derived from the Knoevenagel condensation of the aromatic aldehyde (1) with malononitrile (2) after removing one molecule of water. In the next stage, 1,3-thiazolidinedione (3) in the existence of DIPEAc (12) transforms to its corresponding enolate (3a) and adds to the arylidene malononitrile (13) by Michael addition of the carbanion of 1,3-thiazolidinedione to arylidene cyanoacetamide (14). Then, the cyclic secondary amine (4) attacks the carbonyl group of 1,3-thiazolidinedione to open its ring and cause the formation of a sulfide anion (15) and the intramolecular addition of a sulfide anion to the cyano group in the intermediate (16). After this, at last thiophene (5) is produced by a dehydrogenation process in air. The DIPEAc medium was the best promoter for the preparation of dihydrothiophene derivatives; reasons for this could be explained as follows: (1) the use of DIPEAc raises the solubility of reagents, which results to a higher interfacial area and lower mass transfer resistance. (52). (2) The promoting impact of DIPEAc to the reaction could be credited to its hydrophobic, polarity, and hydrogen-bonding effects (53,54) (Scheme 4), making it easy to develop related products. It was revealed that only polar protic solvents could give the expected outcome, and the hydrogen-bonding effect is the main difference between other solvents and polar protic solvents, so the hydrogen-bonding effect may be the key factor to facilitate the

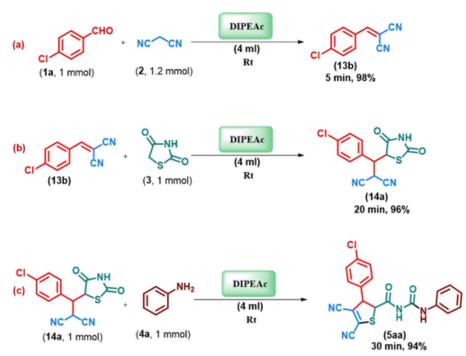
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chlorobenzylidene)malononitrile (13b) was achieved in 98% yield after 5 min (<u>Scheme 5</u>a). In the next stage, the possibility of the Michael addition of 1,3-thiazolidinedione (3) to the arylidene malononitrile (13b) in the presence of DIPEAc was verified. Therefore, the newly synthesized 2-(4-chlorobenzylidene) malononitrile (1 mmol) was treated with 1,3-thiazolidinedione (3, 1 mmol) in the presence of DIPEAc (4 mL) at room temperature and as expected, 2-((4-chlorophenyl) (2,4-dioxothiazolidin-5-yl) methyl) malononitrile (14a) was achieved in 95% yield after 20 min (<u>Scheme 5</u>b). Eventually, the synthesized 2-((4-chlorophenyl)-(2,4-dioxothiazolidin-5-yl) methyl) malononitrile (14a, 1 mmol) under the ideal reaction conditions, and the desired product (5aa) was achieved in 94% isolated yield after 30 min. (<u>Scheme 5</u>c)

Scheme 5



Scheme 5. Study of the Reliability of the Proposed Reaction Pathway for the Synthesis of 5-Amino-4cyano-3-phenyl-*N*-(phenyl carbamoyl)-2,3-dihydrothiophene-2-carboxamide in the Presence of DIPEAc at Room Temperature

As far as sustainability is concerned, it is required to confirm the existing procedure with wellestablished "green metrics" such as the atom economy, reaction mass efficiency, E-factor, optimum efficiency, and practical mass yield. (56-59)

The E-factor is an extensively used green metric for chemical reactions. The lesser the value of the E-

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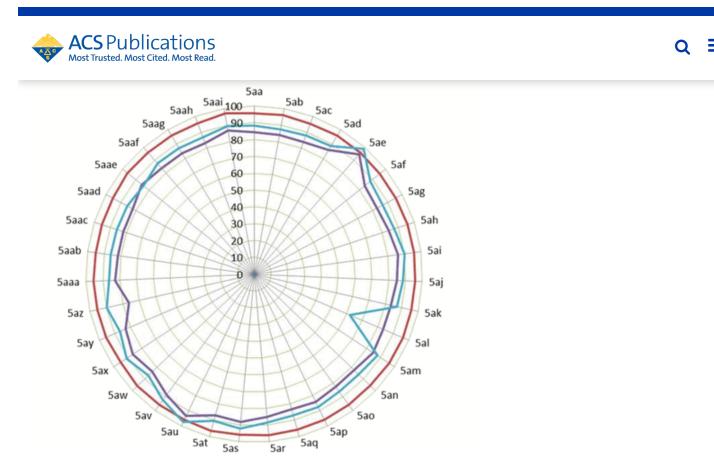


Figure 2. Radar chart of measured green metrics for the synthesis of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophenes (**5aa**–**5aam**).

Later, we measure the green chemistry metrics for both model reactions to afford **5aa** and **5ab** under optimized reaction conditions, as demonstrated in <u>Table 4</u>. The outcome revealed that the values of green chemistry metrics such as atom economy (AE), E-factor, optimum efficiency (OP), reaction mass efficiency (RME), and effective mass yield (EY) are nearly as close to their ideal values as demonstrated here (see the Supporting Information for detailed calculations).

sr. no.	green chemistry merits	ideal value	product (5aa)	product (5ab)
1	E-factor	0	0.20	0.18
2	atom economy (AE)	100%	95.67	96.02
3	reaction mass efficiency (RME)	100%	84.50	84.09

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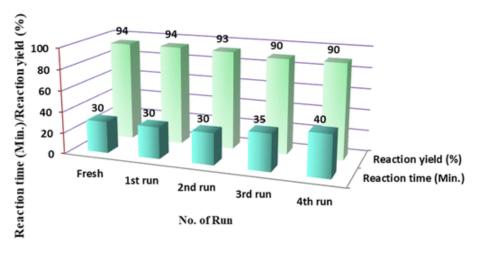




recovered catalyst were proven to be almost identical to the fresh one.

Figure 3

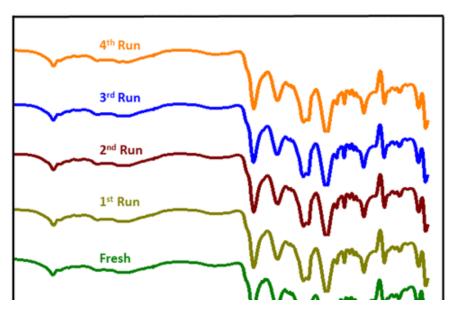
Catalyst Recycling



Reaction time (Min.) = Reaction yield (%)

Figure 3. Recyclability of the catalyst.

Figure 4



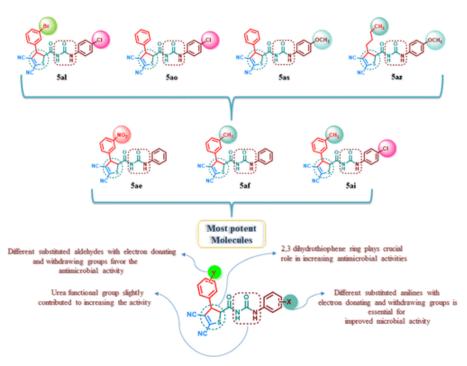
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In the route of identifying several novel antimicrobial agents, we are especially curious about the present work with novel 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene derivatives. In this paper, the antimicrobial screening of 39 newly synthesized compounds was screened against four bacteria and four fungi using the agar well diffusion method. Zone of inhibition (µg/mL) values are displayed in Table 5. (60,61) The outcomes showed that majority of the synthesized compounds demonstrated antimicrobial activities against four bacteria *Pseudomonas aeruginosa* (*P. aeruginosa*), *Escherichia coli* (*E. coli*), *Bacillus subtilis* (*B. subtilis*), and *Staphylococcus aureus* (*S. aureus*) and four fungal strains *Aspergillus niger* (*A. niger*), *Aspergillus flavus* (*A. flavus*), *Fusarium oxysporum* (*F. oxysporum*), and *Candida albicans* (*C. albicans*). The standard drugs ampicillin, ciprofloxacin, miconazole, and fluconazole were used as standards for antibacterial and antifungal activities, respectively. The achieved results revealed that most of the compounds have demonstrated satisfactory to superb inhibitory activity against the four tested bacteria and fungi. The electronic property of the compounds has an adjacent correlation with their biological activity as illustrated in Figure 5. A stepwise molecular optimization to the scaffold of most potent compounds **5al**, **5ao**, **5as**, **5az**, **5ae**, **5af**, and **5ai** and the molecular area highlighted in Figure 5 explored the SAR.

Figure 5



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1	5aa	76 ± 0.22	60 ± 0.34	31 ± 0.09	95 ± 0.75	100
2	5ab	39 ± 0.46	92.4 ± 0.28	98.3 ± 0.20	51.3 ± 0.81	55 ±
3	5ac	71 ± 0.38	96 ± 0.40	71.7 ± 0.61	66.1 ± 0.64	80 ±
4	5ad	95 ± 0.38	130.7 ± 0.87	100 ± 0.25	76 ± 0.39	70 ±
5	5ae	97.7 ± 0.65	65.7 ± 0.87	24.6 ± 0.95	97.2 ± 0.81	30 ±
6	5af	66.8 ± 0.34	68.6 ± 0.54	74.2 ± 0.19	88.9 ± 0.37	30 ±
7	5ag	92.1 ± 0.74	57.1 ± 0.34	75.2 ± 0.37	93.7 ± 0.64	* <u>b</u>
8	5ah	52.0 ± 0.63	88.1 ± 0.35	90.4 ± 0.76	55.8 ± 0.19	150
9	5ai	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	35 ±
10	5aj	77.38 ± 0.67	44.3 ± 0.34	64.2 ± 0.75	109.2 ± 0.44	60 ±
11	5ak	59.3 ± 0.33	82.3 ± 0.57	35.3 ± 0.63	84.5 ± 0.71	55 ±
12	5al	88.4 ± 0.31	92.3 ± 0.44	80 ± 0.17	122.3 ± 0.34	35 ±
13	5am	71.9 ± 0.30	102.5 ± 0.94	66.8 ± 0.90	48.3 ± 0.84	70 ±
14	5an	98.3. ± 0.74	104.6 ± 0 .09	83.0 ± 0.22	52.8 ± 0.10	2500
15	5ao	41.6 ± 0.90	79.3 ± 0.4	101.3 ± 0.62	88.1 ± 0.22	40 ±
16	5ap	168.1 ± 0.64	195.1 ± 0.37	108.5 ± 0.61	156.6 ± 0.78	60 ±
17	5aq	151.0 ± 0.64	188.7 ± 0.74	164.1 ± 0.31	139.7 ± 0.47	30 ±
18	5ar	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	50 ±
19	5as	32 ± 0.46	90.01 ± 0.28	98.3 ± 0.20	50.3 ± 0.81	50 ±

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22	5av	61 ± 0.38	86 ± 0.40	61.7 ± 0.61	56.1 ± 0.64	50 ±
23	5aw	95.3. ± 0.74	108.6 ± 0 .09	73.0 ± 0.22	62.8 ± 0.10	40 ±
24	5ax	53.0 ± 0.73	98.1 ± 0.35	91.4 ± 0.77	56.8 ± 0.19	45 ±
25	5ay	72.9 ± 0.40	103.5 ± 0.94	56.8 ± 0.90	58.3 ± 0.84	60 ±
26	5az	76.8 ± 0.34	58.6 ± 0.54	84.2 ± 0.19	*	25 ±
27	5aaa	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	100
28	5aab	52.0 ± 0.63	88.1 ± 0.35	90.4 ± 0.76	55.8 ± 0.19	50 ±
29	5aac	59.3 ± 0.33	82.3 ± 0.57	35.3 ± 0.63	84.5 ± 0.71	150
30	5aad	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	80 ±
31	5aae	71 ± 0.38	96 ± 0.40	71.7 ± 0.61	66.1 ± 0.64	60 ±
32	5aaf	98.7 ± 0.77	104.6 ± 0.06	84.0 ± 0.46	88.1 ± 0.71	50 ±
33	5aag	98.7 ± 0.77	104.6 ± 0.06	84.0 ± 0.46	88.1 ± 0.71	25. ±
34	5aah	167.1 ± 0.64	193.1 ± 0.37	105.5 ± 0.61	152.6 ± 0.78	150
35	5aai	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	70 ±
36	5aaj	71 ± 0.38	96 ± 0.40	71.7 ± 0.61	66.1 ± 0.64	50 ±
37	5aak	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	100
38	5aal	61 ± 0.38	86 ± 0.40	61.7 ± 0.61	56.1 ± 0.64	60 ±
39	5aam	165.7 ± 0.54	166.7 ± 0.34	204.7 ± 0.33	191.5 ± 0.82	100
std	ampicillin	100 ± 1.24	100 ± 2.14	250 ± 2.99	250 ± 0.88	

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5 ± 2



std fluconazole

^aValues are the average of three readings.

^bAll asterisks denote that no activity was observed up to 200 µg/mL.

The compounds **5ae**, **5af**, **5al**, **5ao**, and **5as** revealed good antibacterial activity against all four bacterial pathogens. The compounds **5ae**, **5af**, **5ai**, **5az**, **5aai**, and **5aag** showed excellent activity against all four fungal strains because of the presence of electron-withdrawing substituents ($-NO_2$, -Br, and -CI) and electron-donating substituents ($-CH_3$ and $-OCH_3$) in the molecule. Compounds **5ae**, **5af**, and **5ai** showed good antibacterial and antifungal activity against the Gram-positive strains *E. coli*,*P. aeruginosa*,*S. aureus*,*B. subtilis*,*C. albicans*,*F. oxysporum*,*A. flavus*, and *A. niger*. We imagine that the presence of a 2,3-dihydrothiophene ring, $-NO_2$, $-CH_3$, and $-OCH_3$ moieties in the molecule contributes significantly to the antibacterial and antifungal activity.

Molecular Docking

To clarify the possible mechanism by which the dihydrothiophene ureidoformamides investigated herein can induce antifungal activity, and to further guide the SAR, an in silico binding study through molecular docking was performed against an important fungal target-sterol 14a-demethylase (CYP51), inhibition of which could prevent the conversion of lanosterol to ergosterol, causing the accumulation of 14amethyl sterols in the cell and leading to impaired cell growth in fungi. This in silico molecular docking approach has now become an integral part of the drug discovery protocol, especially in the absence of available resources to perform the enzymatic assays, imparting knowledge on binding modes, affinities, and the associated thermodynamic interactions with the target enzyme that govern the inhibition of the causative pathogen. The Glide (grid-based ligand docking with energetics) module integrated into the Schrödinger molecular modeling package (Schrödinger, LLC, New York, NY, 2015) was used to predict the binding modes of dihydrothiophene ureidoformamides into the active site of the sterol 14ademethylase (CYP51) enzyme. (62) The 3D crystal structure of sterol 14a-demethylase (CYP51) complexed with its inhibitor fluconazole (PDB code: 3KHM) was retrieved from the RCSB Protein Data Bank (PDB) (https://www.rcsb.org/pdb) and preprocessed using the protein preparation wizard applying the OPLS-2005 force field, which includes the elimination of all crystallographically observed water molecules (as there is no conserved interaction with the enzyme), appending the missing hydrogen/ side-chain atoms corresponding to pH 7.0 considering the appropriate ionization states for the acidic and basic amino acid residues, identification of atom/residue overlaps, creating the disulfide bonds, assignment of reasonable charge and protonation state to the obtained structure, and finally energy minimization of the obtained structure until the average r m s.d. of non-hydrogen atoms converged to

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of sterol 14 α -demethylase using the extra precision (XP) Glide scoring function.

The enzyme-inhibitor complexation predicted by molecular docking showed that all these dihydrothiophene ureidoformamides could bind to CYP51 with a significant binding affinity adopting a similar orientation, and their complexes were stabilized by the formation of several bonded and non-bonded interactions. Their binding energies, which signify the binding relationship, were observed to be negative (-59.484 to -38.094 kcal/mol), while the average docking score was seen to be -7.841 kcal/mol. Furthermore, analysis of the per-residue interaction between these molecules and the active site residues of the enzyme was carried out to identify the most significantly interacting residues and their type of thermodynamic interactions, which is critical in lead optimization. This analysis is discussed for one of the most active analog **5aag** and is summarized in Table S1 (see in the Supporting Information) for the remaining molecules in the series.

The lowest energy-docked conformation of 5aag (Figure 6) showed that it is deeply embedded into the active pocket of CYP51 with significant binding affinity, producing a Glide docking score of -9.702 and Glide binding energy of -59.484 kcal/mol. It could occupy the same coordinates as the native ligand engaging in a close network of bonded and non-bonded interactions with the residues forming the active site. It was found to be stabilized into the active site through a series of significant van der Waals interactions observed with Val461 (-2.854 kcal/mol), Met460 (-2.482 kcal/mol), Thr459 (-1.137 kcal/ mol), Thr295 (-1.33 kcal/mol), His294 (-1.317 kcal/mol), Ala291 (-3.078 kcal/mol), Leu208 (-2.718 kcal/mol), Glu205 (-1.264 kcal/mol), Met106 (-3.171 kcal/mol), and Tyr103 (-3.504 kcal/mol) through the N-(5-amino-4-cyano-3-(4-flurophenyl)-2,3-dihydrothiophene core group, while the pyridine-linked carboxamide side chain showed a similar network of interactions with Cys422 (-1.111 kcal/mol), Leu356 (-2.936 kcal/mol), Ala287 (-1.553 kcal/mol), Leu127 (-1.254 kcal/mol), Tyr116 (-4.448 kcal/ mol), Phe110 (-1.011 kcal/mol), and Ile105 (-1.124 kcal/mol) residues lining the active site. Furthermore, the enhanced binding affinity of **5aag** is also attributed to significantly favorable electrostatic interactions observed with Ala291 (-1.297 kcal/mol), Phe290 (-1.312 kcal/mol), Tyr 116 (-1.361 kcal/mol), and Glu101 (-1.015 kcal/mol) residues. Being a metalloprotein, CYP51 was expected to engage in a significant interaction with the ligand through the prosthetic Hem moiety, which was observed for 5aag also wherein the compound showed significant van der Waals (-4.963 kcal/mol) and electrostatic (-3.696 kcal/mol) interactions with them contributing significantly to binding affinity. While the non-bonded (steric and electrostatic) interactions were observed to be the major driving force for the mechanical interlocking of **5aag**, the higher binding affinity is also supported by a prominent hydrogen-bonding interaction observed through Ala291 (2.419 Å), Phe290 (2.214 Å), and Tyr116 (2.679 Å).

Eiaura 6

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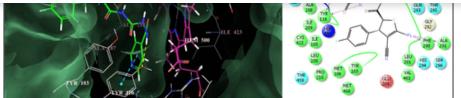


Figure 6. Binding mode of **5aag** into the active site of sterol 14 α -demethylase (CYP51) (on the right side: the pink lines represent the hydrogen bonding, while the green lines signify π - π stacking interactions).

Furthermore, a very close $\pi-\pi$ stacking interaction was also observed with Tyr116 (2.68 Å). Such hydrogen-bonding and π -stacking interactions serve as an "anchor" guiding the orientation of the ligand into the 3D space of the enzyme active site and facilitate the steric and electrostatic interactions therein. A similar network of bonded and non-bonded interactions was observed for other molecules in the series, guiding their binding to CYP51 (see the Supporting Information, S1–S25). The inference derived from this in silico binding studies is now fruitfully utilized for the structure-based lead optimization to arrive at potent molecules with this scaffold.

Conclusions

In conclusion, we have designed a very facile, simple, conveniently practical, and energy-efficient method for easy access of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene derivatives in the presence of DIPEAc (diisopropyl ethyl ammonium acetate) as a reusable catalyst and reaction medium via a one-pot four-component domino reaction at room temperature. Operational simplicity, clean reaction profiles, mild reaction conditions, absence of tedious separation procedures, high atom economy, energy efficiency, excellent yields, and the use of a low-cost and environmentally sustainable ionic liquid are the main advantages of the present method. Likewise, recycling of the reaction media is an added superiority to this protocol. Keeping in mind that the synthetic significance of such pharmacologically relevant 2,3-dihydrothiophene scaffolds directly relates to medicinal chemistry, the present methodology with mild operational simplicity and reaction conditions offers the possibility of its use with cost-effective and environmentally friendlier ways for Gram-scale industrial syntheses as well. Also, a series of 2-(*N*-carbamoyl acetamide)-substituted 2,3-dihydrothiophene analogs were screened for their in vitro antibacterial and antifungal activities. Molecular docking studies of all new derivatives showed a high binding affinity toward sterol 14α-demethylase (CYP51). They provided clues for further modification of the scaffold to improve the activity and selectivity toward the target.

Experimental Section

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Avance DRX-250 and Bruker Avance DRX-400, respectively, in pure DMSO- d_6 or CDCI₃ solvents. IR spectroscopy (Shimadzu FT-IR 8300), in cm⁻¹, was employed for the characterization of the compounds. The HPLC analysis of all compounds was performed on an MS-Agilent 6120 quadrupole system.

General Procedure for the Synthesis of Diisopropyl Ethyl Ammonium Acetate (DIPEAc)

A mixture of anhydrous acetic acid (1 mmol) and *N*,*N*-diisopropylethylamine (1 mmol) was stirred at 0-10 °C for 20 min. The viscous liquid, diisopropyl ethyl ammonium acetate, was achieved. (<u>16,26</u>)

General Procedure for the One-Pot Four-Component Synthesis of 2-(*N*-Carbamoyl acetamide)-Substituted 2,3-Dihydrothiophenes (5aa–5aaj) in the Presence of DIPEAc

A mixture of appropriate aldehyde (1.0 mmol) and malononitrile (1.2 mmol, 0.06 g) was added to a 25 mL round-bottom flask containing a suspension of DIPEAc (4 mL), and the resulting mixture was stirred at room temperature for 2 min. Then, 1,3-thiazolidinedione (1.0 mmol, 0.11 g) and the amine (1.0 mmol) were added, and the reaction mixture was stirred at the same temperature, and use of the TLC followed the progress of the reaction. At the end of the reaction (indicated by thin-layer chromatography), the solvent was recovered, and 5 mL of distilled water was added to the crude product and allowed to stir at room temperature to make the catalyst soluble and for complete solidification of the final product. The precipitates were washed thoroughly twice with 5 mL of distilled water to afford the desired products in pure form. All synthesized compounds were fully characterized based on analytical and spectral studies such as IR, ¹H NMR, ¹³C NMR, and HPLC analysis (for representative compounds **5aa** and **5ab**).

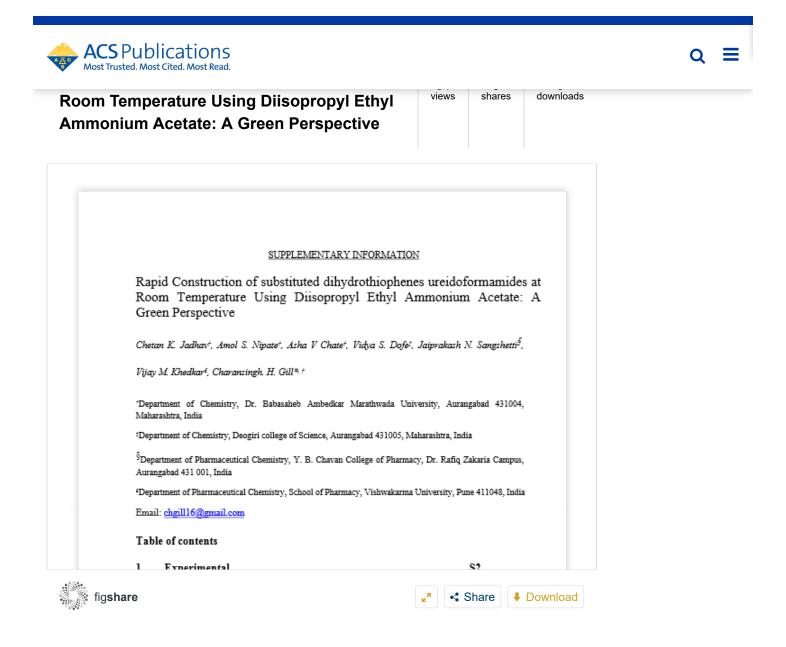
Supporting Information

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The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/ acsomega.0c03575.

 Analytical and spectroscopic data, a copy of ¹H NMR and ¹³C NMR spectra, HPLC reports (5aa-5ab), and calculation of green metrics for compound 5aa as a representative entry (PDF)

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Notes

The authors declare no competing financial interest. This work is dedicated to my beloved parents.

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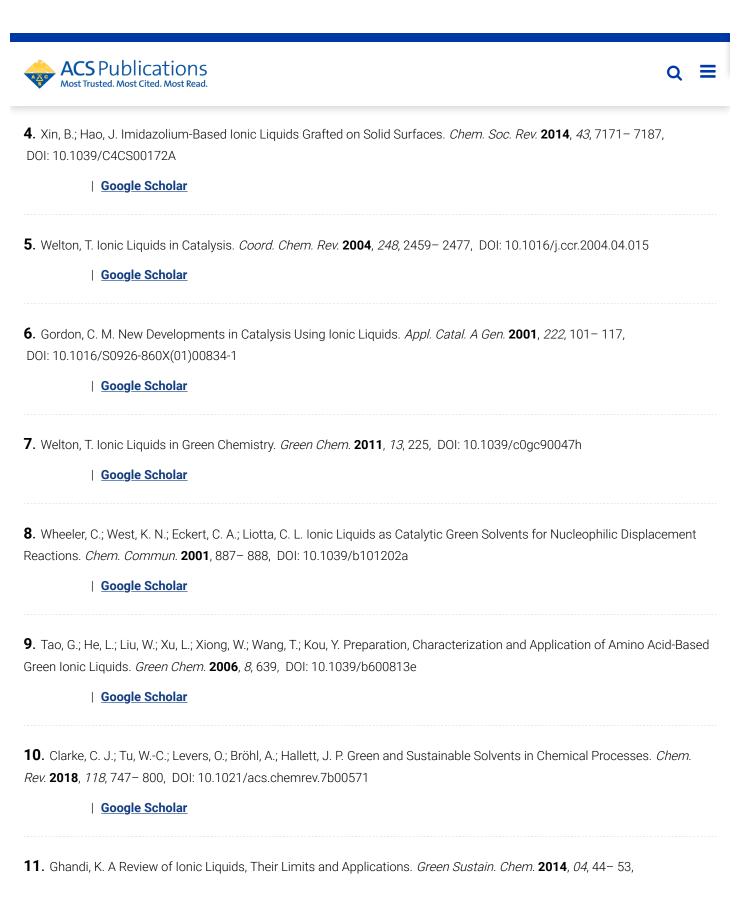
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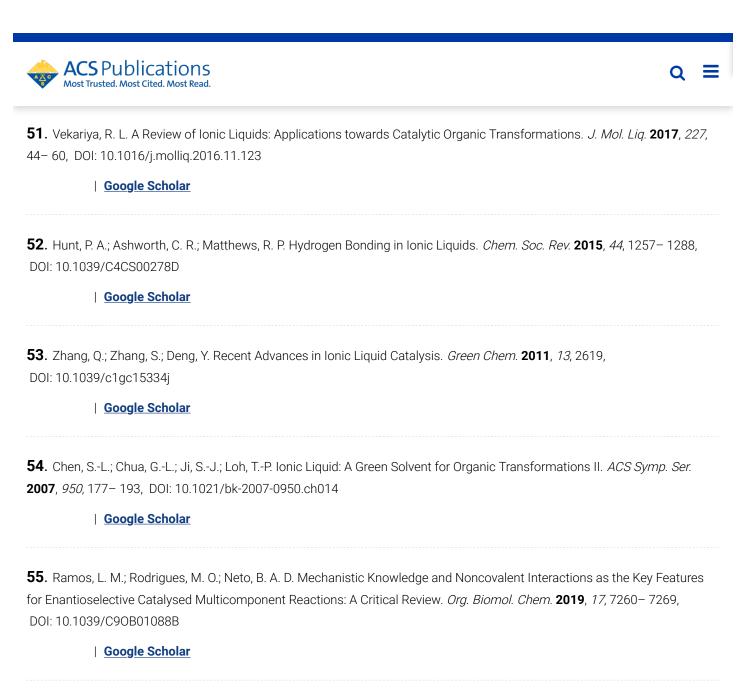
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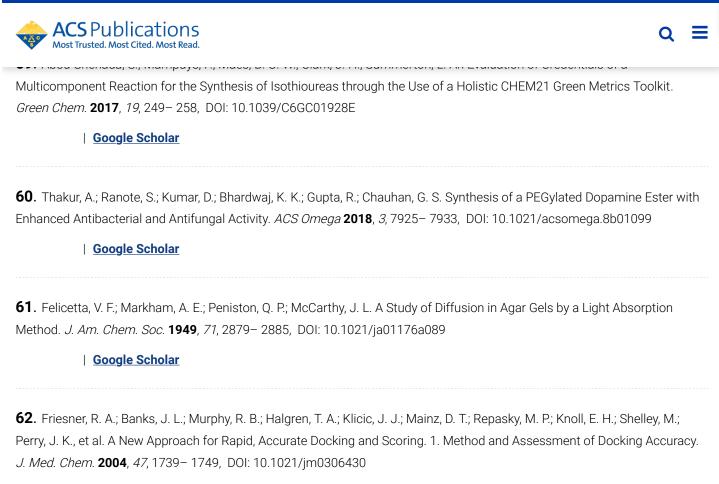
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