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Article

Ethylene-di-amine Modified β-Cyclodextrin Catalyzed Green Synthesis of Pyrimidones and Its *In Silico* Study against ESBL *E. coli* Receptor

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Abstract: Modern organic synthesis is majorly focused on developing environmentally benign synthetic protocol's by employing green chemistry principles. Accordingly, in our recent research work, we herein report the use of modified supramolecular host cyclodextrin as an effective solid based green catalyst for accessing structurally diverse and medicinally relevant pyrimidone architectures. The catalyst and the synthesized compounds 4 (a-r) were characterized using FT-IR, NMR and GC-Mass spectroscopy. Major highlights of the reported work include: atom economical process, extremely milder reaction conditions, operational simplicity, high isolated yields, and excellent catalyst turnover number. The molecular docking studies suggest that the compound 4n has the hydrogen bonding, hydrophobic and π -pair interactions with the active site of active sites of CXT M 15 receptor.

Keywords: Ethylene-di-amine modified β -CD; Multicomponent reaction; 3,4-dihydropyrimidin-2(1H)-ones DHPM derivatives; Solvent-free conditions; Reusability

1. Introduction

After the initial discovery of Cyclodextrins (CD) in 1891 by Villiers, the use of this oligosaccharides for various catalytic applications was disclosed by various researchers across the discovery globe [1]. In the recent years CD have been extensively serving as host molecules in supramolecular chemistry. They offer several advantages over other host molecules, including wide availability from renewable sources, good water solubility, biocompatibility and simplicity in chemical modification. Their molecular recognition of guest molecules is also well recognized [2]. These "molecules of holes" have lately gained favour as building blocks for the self-assembly of supramolecular structures with varied porosity [3]. The key feature of CD is its cone-shape structure which provides a hydrophobic interior cavity capable of hosting a variety of guest molecules with appropriate polarity and sizes. Furthermore, the CD's hydrophilic outer surface, can act as a molecular shuttle, allowing hydrophobic molecules to be transferred to the aqueous phase [4].

CD are classified as α –CD with six D-glucose units, β –CD with seven D-glucose units, and γ –CD with eight D-glucose units are the most prevalent [5]. The hydroxy and acetyl functional groups on CD dictate the majority of their chemical characteristics, say solubility in specific. Particularly, 18.5 g of β –CD in 1 L of water at room temperature, the primary and secondary hydroxy groups on β –CD

have the potential to form a high intramolecular H-bonding network, which results in the reduction of water solubility. Thus, the hydroxy group on β –CD is replaced by the methoxy group and it reduces the H-bonding strength eventually and emphasizes the water solubility [6].

In modern drug discovery approaches, introducing multiple diversity points in a molecule is of immense potential for generating hit molecules for various therapeutic areas. Among the various synthetic approaches, MCRs play a vital role in accessing this structural diversity at multiple points. Such reactions explore the synthesis of bioactive molecules and also facilitate to complete the difficult task of pharmaceutical and therapeutic chemists. By this fact, the Biginelli reaction has been employed to synthesis various bioactive molecules of 3,4-dihydropyrimidin-2(1*H*)-ones (DHPMs) [7–9].

The importance of heterocyclic frameworks is well documented in literature as depicted by its importance in various fields like material science, agrochemicals and drug discovery [10–15]. 3,4-dihydropyrimidin-2(1*H*)-ones (DHPMs) are found to possess diverse range of pharmacological properties, including anti-tubercular, [16] antimicrobial, [17] anti-consultant, [18] anti-cancer, [19] anti-viral, [20] and anti-tubercular properties.

Our current research was mainly focused on employing cyclodextrins as catalyst for Biginelli condensation to access the pyrimidone analogues. It has been found that assorted catalyst and several methods were employed for the same. Many catalytic systems such as the use of strong protic acids H₂SO₄ [21], HCl [22], various Lewis and Bronsted acids like Bi(NO₃)₃ [23], Sulfated silica tungstic acid [24], sulfated tungstate [25], SiO₂-polyphosphoric acid (SiO₂-PPA) [26], Silica sulfuric acid [27], Bismuth Subnitrate [28], cellulose sulfuric acid [29], sulfated polyborate [30], H₂SO₄-Silica, [27], Al₂O₃-MeSO₃H [31], Al(HSO₄)₃ [32], Al₂O₃-SO₃H [33], sulfated zirconia [34], zeolites [35], and metal trifles [36] were successfully employed. Further methods involved are ultrasonic [37], microwave-assisted [38] and ionic liquids [39] employing different green catalysts [40–43]. However, most of these reported methodologies have several limitations including low yields, prolonged reaction times and use of metal catalysts.

While native cyclodextrins can operate as catalyst for a variety of organic reactions, the accession of acidic or basic functional groups allows to precise their functions. Per-6-amino-cyclodextrin, a successful modified form of cyclodextrin contains basic amino groups function as a catalyst and supramolecular ligand for reactions such as asymmetric Michael addition [44], Mizoroki-Heck coupling [45], N-arylation [46], cyanation [47], enantioselective Henry reaction [48] and also for the multicomponent synthesis of pyranopyrazoles and 2-amino-4H-benzo[*b*]pyrans [49,50]. The enhanced solubility in the reaction medium is a major hallmark of homogeneous catalytic systems thereby increasing its catalytic activity and substrate accessibility to the catalytic site when compared to heterogeneous catalysts [51–57].

The synthetic methodologies developed have several limitations such as the use of costly catalysts, hazardous chemicals, higher temperatures, longer reaction time, time-consuming procedures and lower yields. To fulfill the need for developing more environmentally benign protocols, our research was focused on the use of modified cyclodextrins as catalysts for its operational simplicity, enhanced yields, cost effective, faster reaction rates etc., for the synthesis of 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) derivatives. Subsequently, a series of 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) was synthesized using aromatic aldehydes, heterocyclic aldehyde, acetoacetate (Ethyl, Methyl & Ethyl-4-chloro), Urea with employing modified β -Cyclodextrins-Ethylene Diamine (E@ β -CD) as a carrier.

The synthesized compounds were further evaluated [58,59] for, the Molecular Docking study was performed using docking tool AutoDock Tool version 1.5.6, against the ESBL-producing *E. coli* receptors (CTX-M-15).

2. Materials and Methods

Aerobic conditions were used for all reactions. All the chemicals purchased from Sigma Aldrich and were used without additional purification, unless otherwise stated. On a Bruker spectrometer, NMR spectra were recorded at 400 MHz using different solvents, such as DMSO d_6 and CDCl₃, with

TMS as the internal standard. A BRUKER ALPHA II ECO-ATR instrument was used to measure the FT-IR spectrum from 4000–550 cm⁻¹ range. Thermo Fisher Instruments Limited's (US) LCQ Fleet recorded Electrospray ionization mass spectrometry (ESI-MS) was used, under negative ion mode.

2.1. General Procedure for the Synthesis of 3,4-dihydropyrimidin-2(1H)-ones DHPMs (4 a-r)

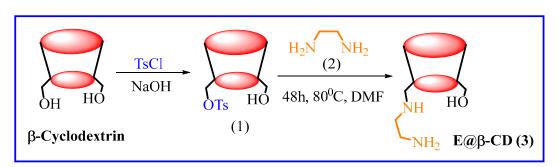
 $E@\beta$ -CD (0.001 mmol) was taken without solvent in a round-bottom flask, aldehydes (1 mmol), β -Keto-Ester (1 mmol) and urea (1 mmol) were added. The mixture was then stirred for 5 min at room temperature. TLC was used to ensure the completion of the reaction with ethyl acetate and n-hexane (1:4) as the eluent system. After this process, the mixture was extracted twice with ethyl acetate (2 x 20 ml). The organic layer of the extract was concentrated, washed with distilled water and dried over anhydrous Na₂SO₄. Without performing any additional purifications, the resulting solid was recrystallized with ethyl acetate.

2.2. Molecular Docking Studies

Molecular docking studies with the AutoDock Tool docking programme, version 1.5.6, the binding affinities of the synthesised compounds 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) (4b, 4d, and 4b) and the crystal structure of CTX-M-15 (PDB: 4HBU) were assessed. Using CHEMSKETCH, the heterocyclic ligand's structure was sketched in mol format, which was later translated to pdb format by OPENBABEL. The Protein Data Bank, located at https://www.rcsb.org/pdb, was used to download the CTX-M-15 protein in pdb format. The heterocyclic compounds (4b, 4d, and 4b) and the receptor (CTX-M-15) files were created using AutoDock tools. After removing all heteroatoms and water molecules, the receptor molecule was then assigned with polar hydrogen atoms and Kollman charges. Then, all other bonds were permitted to rotate while the rotatable bonds were assigned to ligand molecules. A box with a rigid spacing of 0.372 Å and 60 points each of the three dimensions (60 x 60 x 60 in x x y x z dimensions) was used to wrap the CTX-M-15 molecule. The Lamarckian evolutionary algorithm, as implemented in AutoDock, was used for the docking computations, and all other parameters were left at their default values. Using the DISCOVERY studio client, docked positions were shown in three dimensions, while LIG-PLOT plus was utilized to see hydrogen bonds and hydrophobic interactions in 2D.

3. Result and Discussion

In the current work, we have employed $E@\beta$ -CD (3) as an ace supramolecular host for the synthesis of three-component pyrimidine derivatives under solvent free condition at room temperature. The reusability of the catalyst for multiple times highlights the significance of the developed protocol in terms of green chemistry perspective. In Scheme 1, the procedure for the synthesis of $E@\beta$ -CD (3) is illustrated. Tosylation was assisted to develop Mono-6-tosyl- β -cyclodextrin (1) from commercially available β -CD and the intended product, $E@\beta$ -CD (3) was produced as a yellow solid after treating with ethylene-di-amine (2). $E@\beta$ -CD (3) was successfully synthesized in a single step with a 78% yield. NMR and ESI- MS spectra validated the structure of $E@\beta$ -CD (3).



ESI-MS spectrum (Figure S3) shows an m/z peak at 1177.89 which corresponds to [M+1 adduct]. The chemical shift values of the 1 H NMR and 13 C NMR spectrum are found to be in good agreement with the synthesized E@ β -CD (3) (Figure S1 and Figure S2).

Insight II molecular modelling studies were used to conduct energy minimization analyses to ascertain the prospective inclusion of ethylene-di-amine in $E@\beta$ -CD (3) of the β -CD cavity. Δ^a Ea (Kcal.M-1) values of Mode A and Mode B shown in Table 1, reveals that the existence of ethylene-di-amine group outside the $E@\beta$ -CD (3) cavity is more favored than inside. (Figure S4)

Table 1. Molecular modeling studies of ethylene-di-amine modified β -cyclodextrin.

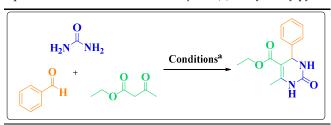
Mode of Inclusion	ΔE^{a}	(Kcal.M ⁻¹)
Di-amine group outside the pyr:β-CD cavity(Mode A) -	-64.8800
Di-amine group inside the pyr:β-CD cavity. (Mode B)) -	50.8492

 $\mbox{\sc a}Binding$ energy calculation of modified $\beta\mbox{-cyclodextrin}.$

3.1. E@β-CD Catalyzed for Multi-Component Reactions

The reaction was optimized using Urea, ethyl acetoacetate and benzaldehyde as substrates and the results are shown in Table 2.

Table 2. Optimized reaction conditions in E@β-CD (3) catalyzed by pyrimidones^{a,b.}



Entry	Catalyst	Medium	Time (h)	Yield (%)b
1	β-CD	Water	24	28
2	Ethylene-di-amine (2)	-	8	26
3	Methanol	-	8	28
4	Ethanol	-	8	32
5	Methylamine	-	8	35
6	Diethylamine	-	8	33
7	Triethylamine	-	8	34
8	Pyridine	-	8	36
9	E@β-CD (3)	DMF	24	58
10	E@β-CD (3)	DMSO	24	60
11	E@β-CD (3)	-	5 min	96
c12	E@β-CD (3)	-	5 min	96 ^b
^d 13	E@β-CD (3)	-	5 min	96°

^aReaction conditions: benzaldehyde (1 mmol, 1 equiv), urea (1.2 equiv) and ester (1.2 equiv), E@β-CD (3) (0.2 mol%) RT, 5 min. ^bIsolated Yield, ^c0.01 mmol of catalyst and ^d0.001mmol of catalyst.

The investigation of catalyst loading on the reaction time and yield was carried out. The minimal yield was observed when the plain β -CD was administrated as a catalyst in an aqueous medium at room temperature (entry 1). Subsequently, the use of ethylene-di-amine (2) as catalytic system resulted in poor yields of the product (entry 2). Besides a wide range of organic bases were employed as catalyst under solvent-free condition to see the outcome of the reaction, such as methanol (entry 3), ethanol (entry 4), methylamine (entry 5), diethylamine (entry 6), triethylamine (entry 7) and pyridine (entry 8) as catalysts (entries 3-8). Surprisingly, lower yields were observed in these cases.

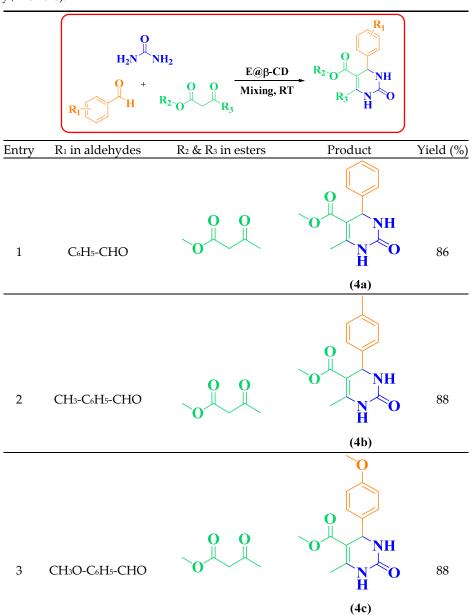
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Similar suboptimal yields were observed when $E@\beta$ -CD (3) was used for the first time as a catalyst in this reaction in DMF and DMSO (entries 9 and 10).

To our surprise, an excellent yield of 96% was observed (entry 11) at a shorter reaction time of 5 min when the catalyst is used with the substrates in solvent-free conditions. Similarly (entry 12), the addition of successive amounts of urea, ethyl acetoacetate, and aromatic aldehydes led to a quantifiable yield of pyrimidones even with 0.01 mmol amount of E@ β -CD (3) and the reaction was completed in 5 min. No traceable change in the yield after admitting with 0.001 mmol of catalyst reaction (entry 13). The protocol developed was highly atom economical and excellent isolated yields were observed. The impact of this efficient supramolecular host E@ β -CD (3) has been demonstrated in this study with high quantitative yield beside with the facile feature of reusability.

To further expand the substrate scope of the developed methodology, electronically biased aldehydes, β-ketoesters and urea were employed. The observed results are listed in Table 3.

Table 3. $E@\beta$ -CD (3) catalyst multicomponent reactions of aldehydes, amine and acetoacetate (Methyl, Ethyl/ 4-chloro).



^aReaction conditions: Aldehyde (1 mmol, 1 equiv), urea (1.2 equiv) and ester (1.2 equiv), E@ β -CD (3) (0.2 mol%) RT, 5 min. ^bIsolated yield in solvent-free conditions for 5 min at room temperature.

The substrates chosen were furnished the corresponding 3,4-dihydropyrimidin-2(1*H*)-ones (DHPMs) with good to excellent yield with a shorter reaction time shown in Table 3. Electron-withdrawing substituents on aromatic aldehydes like nitro groups gave excellent yield in longer reaction time (Table 3, entries 8 & 15), whereas the halogen substituents resulted in excellent yield in shorter reaction time (Table 3, entries 4, 6, 9, 10, 12, 13 and 16). Electron donating substituents like 4-methylbenzaldehyde showed a moderate yield in the shorter reaction time, whereas the methoxy and

di-methoxy substituents requires longest time and their yields are also moderate (Table 3, entries 3, 5, 11 & 14). These, may be due to their electron releasing nature. The bulkier aldehyde like α -Napthaldehyde and heterocyclic aldehyde like thiophene-2-carboxaldehyde (Table 3, entries 17 & 18) resulted in good yield. The optimized protocol tolerates a wide range of functional groups and gave broad scope to the Biginelli reactions. The present systems offer various advantages: (i) recyclability of the catalyst without significant loss of its catalytic activity, (ii) readily available, (iii) excellent yield in short reaction times, (iv) Simple and easy separation.

The reaction mechanism, which is important for revealing the specific process, was investigated systematically based on the previous reports [60]. A presumptive mechanistic pathway for the assemblage of 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) derivatives catalyzed by E@ β -CD (3) is presented in Scheme 2.

$$R_2$$
 R_1 R_2 R_2 R_3 R_4 R_4 R_5 R_5 R_5 R_5 R_6 R_6 R_6 R_7 R_8 R_8 R_8 R_9 R_9

Scheme 2. Proposed Mechanism for the synthesis of 3,4-dihydropyrimidin-2(1*H*)-ones (DHPMs) derivatives.

According to the proposed mechanism, $E@\beta$ -CD (3) primarily supports to activate the carbonyl group of aldehydes to give an intermediate (I), the nucleophilic addition of urea followed by cage elimination under acid conditions, then dehydration takes place to form the intermediate (II), wherein the ethyl acetoacetate was added. The final product 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) is be obtained immediately and we speculated that intermediate (III) can be obtained through the reaction between intermediate (II) and the enolized ethyl acetoacetate. Finally, intermediate (III) underwent cyclisation and followed by dehydration quickly and afforded the target product.

Once the reaction is completed, the catalyst was removed from the substrates or products and washed with EtOH, EtOAc and n-hexane, respectively. In Figure 1, the acquired white powder was subsequently dried in an oven and utilised in additional runs. Another crucial feature of this eco-friendly, effective and active heterogeneous catalyst is its recyclable nature.

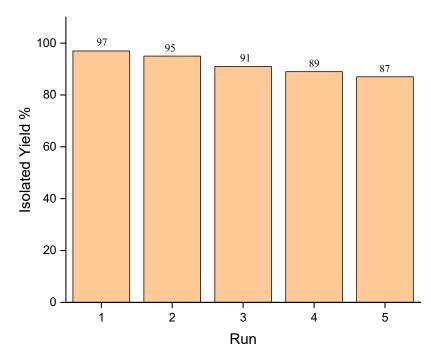


Figure 1. Reusability of the E@ β -CD (3) for the synthesis of DHPM.

Our catalyst serves better over other catalysts that have been reported for the Biginelli reactions between aromatic benzaldehydes, β -ketoesters and urea, details are presented in Table 4. The highlights of the developed protocol include good to excellent yield, involves mild reaction condition and easy work up procedure, recyclable and reusable nature of the catalyst.

Table 4. Comparison table for the catalyst performance of different supported catalysis in the synthesis of pyrimidines.

Entry	Catalyst	Reaction condition	n Time	Yield (%)) Ref
1.	E@β-CD / SF	Rt	30 min	97	This work
2.	KSF/SF	130 °C	48 h	74–88	[61]
3.	Lanthanide triflate/ SF	100 °C	1-1.5 min	81–91	[62]
4.	strontium(II) triflate/ SF	70 °C	4 h	85–97	[36]
5.	Mg-Al-CO ₃ hydrotalcite, SF	80 °C	30–60 min	71–74	[63]
6.	DBSA (5 mol %) / SF	80 °C	2.5-3h	81-94	[64]
7.	β-CD SF	100 °C	3 h	85	[65]
8	β-CD-SO₃H SF	100 °C	2 h	83	[66]
9	β-CD-HCl	EtOH/ reflux	8 h	92	[67]
10	nano-γ-Fe ₂ O ₃ -SO ₃ H SF	60 °C	3 h	90	[68]
11	PS-PEG-SO ₃ H	Dioxane/80°C	10 h	86	[69]
12	Fe ₃ O ₄ /PAA-SO ₃ H SF	RT	120 min	90	[70]
13	Bentonite/PS-SO3H SF	120 °C	30 min	89	[71]
14	Tartaric acid	EtOH/Reflux	4 h	92	[72]
15	Citric acid	EtOH/Reflux	4 h	96	[73]
16	Lactic acids	EtOH/Reflux	2.5 h	92	[74]
17	Ascorbic acid	Solvent free	6 h	85	[75]
18	Imidazole-1yl-acetic acid	l Water/reflux	30 min	94	[76]
19	Sulfanilic acid	Water	3 h	98	[77]
20	Phenyl Phosphonic acid	ACN/reflux	4 h	97	[78]

SF – Solvent Free; DBSA – p-Deodecylbenzene sulfonic acid; β -CD- SO_3H – Cyclodextrin modified Propyl Sulfonic acid; PS-PEG- SO_3H – Polysterene-poly(ethylene glycol) sulfonic acid; PAA- SO_3H – Pheylaceticacid sulfonic acid.

3.2. In-Silico Analysis

The Enterobacteriaceae that produces ESBLs, which includes *Escherichia coli* and *Klebsiella pneumoniae* are the rifeness of pathogenic bacteria. The Indian patients are frequently infected with urinary tract, pneumonia and septicemia infections by this ESBLs. These organisms are virtually resistant to all third generation cephalosporins because of their plethoric variety of ESBL genes. As a result, the hunt for alternative antimicrobial compounds to address this deteriorating clinical situation has been triggered to treat these infections. Resistance to third generation cephalosporins like ceftazidime and cefotaxime (by ESBL generating bacteria) are one of the most significant developing resistance challenges. These antibiotics are typically recommended to treat serious bacterial infections. Since beta lactamases play a significant part in gram-positive bacteria's negative resistance mechanisms, which are linked to easily transposable plasmidic resistant determinants. The centre of the antibacterial activity of beta lactam antibiotics is the amide bond in the beta lactam ring. These beta lactamases cleave the ring, rendering the antibiotics inactive against bacteria. The periplasm of the resistant bacterium secretes beta lactamases. According to reports, CTX-M-15 hydrolyzes cefotaxime with a high catalytic efficiency causes bacterial resistance to cefotaxime.

The examined ligands (4a, 4d and 4n) strongly interacted with the active sites of CTX-M-15 receptor via hydrogen bonds and hydrophobic interactions (Figure 2, and Table 5).

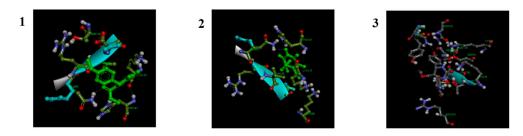


Figure 2. Docking poses of the compounds 4b (1), 4d (2) and 4n (3) with CXT M 15 receptor showing 3D interactions.

The ligand and the receptor interaction of the compound 4n shows five hydrogen bonds. The interaction of three hydrogen bonds are between the oxygen atom of the ligand and hydrogen atom of the receptor ASN10, SER 130 & THR 235 (O---H = 2.12 A° , dihedral angle = 142.45° ; O----H = 2.10 A° , dihedral angle = 140.82° ; O----H = 1.8 A° , dihedral angle = 169.9°) respectively. The rest two hydrogen bond are between the hydrogen atom of the ligand and oxygen atom of the receptors THR 235 and SER 130 (H---O = 2.19 A° , dihedral angle = 131.59° ; H----O = 2.2 A° , dihedral angle = 129.3°). Results of the 4d show that there are three hydrogen bonds in which all the interactions are between oxygen atom of the ligand and hydrogen atoms of the receptors GLN 93, LYS 137 and HIS 141 (O---H = 2.4 A° , dihedral angle = 97.40° ; O----H = 1.78 A° , dihedral angle = 154.74° ; O----H = 1.77 A° , dihedral angle = 156.08°). Hydrogen bond interaction for the compound 4b possess the same residues noted for the compound 4d with different bond distance and dihedral angle. The residues involved in the interactions are GLN 93, LYS 137 and HIS 141 (O---H = 2.01 A° , dihedral angle = 141.72° ; O----H = 1.8 A° , dihedral angle = 142.49° ; O----H = 2.04 A° , dihedral angle = 153.3°).

In addition to those hydrogen bonds, the binding models were stabilized with the hydrophobic interactions between the ligands (4b, 4d and 4n) and the residues SER 70, SER 237, TYR 105, GLY 236, ASN 170, ARG 94, GLU 96, VAL 95, THR 116, ARG 94, VAL 195(A) and GLU 96(A) of the CXT M 15 receptor molecule.

The validation of the best docking poses and binding affinity of the compounds 4b, 4d and 4n (Figure 2), are found within the active sites of CXT M 15 receptor. 2D interactions and binding affinity

are shown in (Figure 3) and Table 5 respectively. Results show that, there exists a similarity in hydrogen bonding found in CXT M 15 & ceftoxamine [79] and between 4n and CXT M 15.

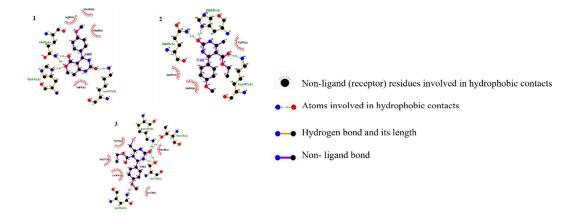


Figure 3. Docking poses of the compounds 4b (1), 4d (2) and 4n (3) with CXT M 15 receptor showing 2D interactions.

Table 5. Molecular docking parameters of the compounds 4 (a, d & m) with receptor (CXT-M-15).

Entry	vdW + H bond + electrostatic + dissolving energy (1) (kcal/mol)	Final total internal Energy (2) (kcal/mol)	Torsional free energy (3) (kcal/ mol)	Unbound system's energy (4) (kcal/mol)	Estimated free energy of binding [1 + 2 + 3-4] (kcal/mol)
4a	-7.53	-2.83	1.49	-2.83	-6.04
4d	-7.21	-2.28	1.49	-2.28	-5.72
4n	-6.21	-2.88	1.69	-2.25	-4.96

4. Conclusions

In summary, the synthesis of modified Ethylene-di-amine modified β -CD (E@ β -CD (3)) using mono-tosy- β -CD and ethylene diamine used as a promoter for the first time in Biginelli reaction. The catalyst and the synthesized compounds 4 (a-r) were characterized using FT-IR, NMR and GC-Mass spectroscopy. It's been evident and demonstrated that E@ β -CD (3) is a very effective and recyclable transition metal-free catalyst for the one-pot three-component synthesis of 3,4-dihydropyrimidin-2(1H)-ones (DHPMs) derivatives under solvent-free conditions. The amount of catalyst, the temperature and the solvent all had a substantial impact on the reaction system. Short reaction periods, high to exceptional yields, elimination of toxic transition metals or organic solvents, ease of workup, reusability of the catalyst and ease of product purification are the key benefits of this procedure. Further, the docking results reveal that the binding modes of compound 4n are in close approximation with the active sites of cefotaxamine hydrolysis sites with four common amino acids involved in the hydrogen bonding with CXT M 15 receptor.

Supplementary information: The data that supports the findings of this study are available in the supplementary material of this article. FT-IR, ¹H NMR, ¹³C NMR spectra and ESI-MS data are provided in Supplementary Materials.

Author Contributions: RMK and MSP contributed to writing - the original draft, methodology, conceptualization, validation and investigation. MMA, SA, ARB, RI and MAK contributed to validation, data curation and formal analysis.

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Conflicts of Interest: The authors declare that they have no conflict of interest.

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