



Synthesis and characterization of new 4*H*-chromene-3-carboxylates ensuring potent elastase inhibition activity along with their molecular docking and chemoinformatics properties

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ABSTRACT

A new series of 4*H*-chromene-3-carboxylate derivatives were synthesized using multicomponent reaction of salicylaldehyde, ethyl acetoacetate and dimedone in ethanol with K₃PO₄ as a catalyst at 80 °C. The structures of all newly synthesized compounds were confirmed by spectral techniques viz. IR, ¹H NMR, ¹³C NMR, and LCMS analysis. The newly synthesized compounds **4a** to **4j** were screened against elastase enzyme. Interestingly, all these compounds found to be potent elastase inhibitors with much lower IC₅₀ value. The compound **4b** was found to be most potent elastase inhibitor (IC₅₀ = 0.41 ± 0.01 μM) amongst the synthesized series against standard Oleanolic Acid (IC₅₀ value = 13.45 ± 0.0 μM). The Kinetics mechanism for compound **4b** was analyzed by Lineweaver-Burk plots which revealed that compound inhibited elastase competitively by forming an enzyme-inhibitor complex. Along with this, all the synthesized compounds (**4a** – **4j**) exhibits excellent DPPH free radical scavenging ability. The inhibition constant *K_i* for compound **4b** was found to be 0.6 μM. The computational study was comprehensible with the experimental results with good docking energy values (Kcal/mol). Therefore, these molecules can be considered as promising medicinal scaffolds for the treatment of skin-related maladies.

1. Introduction

In the development of modern organic chemistry, heterocyclic compounds play a significant role. With its origins rooted in organic synthesis and medicinal chemistry, heterocyclic compounds present themselves as a fundamental division of organic chemistry [1]. Chromenes are one of the major classes of naturally occurring heterocyclic compounds. They found in a large number of natural products and possess a wide range of biological activities [2]. Out of them functionalized 4*H*-chromenes have received considerable attention in recent years for their excellent biological properties such as anticoagulant [3], anticancer [4–7], spasmolytic and anti-anaphylactic [8–10], antimicrobial [11–13], antioxidant [14], and antimalarial [15] activities. Along with this, they have a role in pigments [16] and photoactive materials and are utilized as potential biodegradable agrochemicals

[17]. This moiety also occurs in various natural products [18]. Several procedures for the preparation of 4*H*-chromenes with various starting materials have been reported in recent years [19].

Elastin is the protein widely distributed in the vertebrate tissues of human beings especially abundant in ligaments, lungs and skin. It is being hydrolyzed or cleaved by elastase enzyme belonging to the class of serine proteases [20]. It is also the key enzyme which attacks the all major matrix protein of connective tissue [21]. Skin aging is a naturally occurring process due to more exposure of skin by chronic ultra violet radiation. The continuous exposure of ultra violet radiations causes the formation of reactive oxygen species and lipid peroxides results into physical changes in the connective tissues and skin [22]. It leads to the loss of skin elasticity which can ultimately become the reason of wrinkle formation, brown spots, skin cancer, uneven pigmentation, leathery appearance, solar elastosis, laxity and melanoma [23–25].

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Elastase is the major responsible enzyme for wrinkle formation and dehydration on the skin [26]. It also causes delayed wound healings, increased inflammation progress and tissue permeability.

Recently, some medicinal plant's extracts have been reported as elastase inhibiting cosmeceuticals [27,28]. Along with this, some synthesized compounds, such as safranal [29], aryl and heteroaryl oxime ester derivatives [30], substituted perhalo-2-nitrobuta-1,3-dienes [31], and thiazol-2-(3H)-ones [32] have been reported as elastase inhibitors, but still the need for the development of superior and novel inhibitors is demanded. As per our continuation in synthesis of chromene compounds [33,34] and experience of previous efforts in enzyme inhibitory potential [35,36], we focus our attention towards design and synthesis of 4H chromene compounds which could be potent inhibitors against elastase activity.

2. Result and Discussion:

2.1. Chemistry:

In the beginning, we design and synthesize new 4H-chromene-3-carboxylates using multicomponent reaction of salicylaldehyde (**1**) (1 mmol), ethyl acetoacetate (**2**) (1 mmol) and dimedone (**3**) (1 mmol) in ethanol (Scheme 1). Favorably, the reaction proceeds well and resulting into the expected 4H-chromene-3-carboxylate product which was further confirmed by spectral techniques such as IR, ¹H & ¹³C NMR and LCMS. The reaction of salicylaldehyde, ethyl acetoacetate and dimedone was chosen as a model reaction.

Initially, we tried to carry the model reaction without catalyst in ethanol at reflux condition (Entry 1, Table 1). But the reaction did not proceed well and results into very low yield of product. After that screening of various catalysts (K₂CO₃, K₃PO₄, NH₄OAC, DABCO and NaHCO₃) was done with model reaction using 30 mol% of each catalyst (Entries 2 – 7, Table 1). We found that out of all catalysts screened, K₃PO₄ drag the desired reaction with higher yield in less reaction time (Entry 3, Table 1). Therefore, we perform the screening of catalytic amount of K₃PO₄ for model reaction (Entries 8 – 10, Table 1). From Table 1, we can conclude that the 20 mol % of K₃PO₄ (Entry 9, Table 1) was found to be the enough and good catalytic amount for the proposed transformation. Therefore, the final optimized reaction conditions for present protocol are 20 mol % of K₃PO₄ in ethanol at 80 °C.

With this optimized reaction conditions in hand, such as 20 mol % of K₃PO₄ in ethanol at 80 °C, the generality of protocol was assessed using variety of substituted salicylaldehydes with electron donating and withdrawing group in its structure (Table 2). All reactions proceeds well in result of expected products with good to excellent yield. The products containing electron withdrawing group (Entries 2 – 6, Table 2) in its structure shows higher practical yield and short reaction time than the compounds containing electron donating group (Entries 7 – 10, Table 2). All the synthesized compounds were characterized with spectral techniques viz., IR, NMR and LCMS. Pleasingly, the

Table 1
Screening of catalyst.

Entry	Catalyst	mol %	Time in min	Yield %
1	–	–	180	32
2	K ₂ CO ₃	30	140	74
3	K ₃ PO ₄	30	125	85
4	NH ₄ OAC	30	140	78
6	DABCO	30	150	77
7	NaHCO ₃	30	145	71
8	K ₃ PO ₄	10	130	82
9	K ₃ PO ₄	20	120	87
10	K ₃ PO ₄	40	125	84

Reaction conditions: salicylaldehyde (1 mmol), Ethyl Acetoacetate (1 mmol), Dimedone (1 mmol), Solvent: Ethanol (10 mL), Temperature: 80 °C

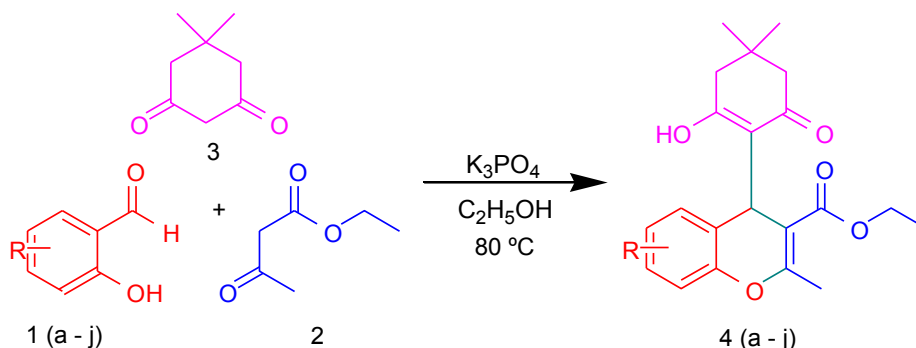
spectral data of synthesized compounds is in accordance with their expected structure. After successful synthesis and its characterization, all the synthesized compounds (**4a-4j**) were subjected to screen for their elastase activity. Satisfyingly, all synthesized compounds (**4a-4j**) exhibits good elastase inhibitory activity which was further supported by *in silico* analysis.

2.2. Biology:

2.2.1. Elastase inhibition and structure activity relationship:

All newly synthesized compounds (**4a-4j**) were scrutinized for their inhibitory potentials against elastase enzyme. The results were outlined in Table 3. Most of the compounds exhibits potent inhibitory activity against elastase as evident from their much lower IC₅₀ values (IC₅₀ value = 0.41 ± 0.01 to 8.71 ± 0.66) as compared to standard oleanolic acid (IC₅₀ value = 13.45 ± 0.01 μM). All compounds revealed much lower IC₅₀ values than the standard oleanolic acid which signifies their potency for elastase inhibition.

The structure activity relationship (SAR) was predicted by the presence of substituent group on aryl ring, as it is the only variable part and remaining all other parts are similar in all molecules. The presence of electron withdrawing group on aromatic ring shows better activity than the compound containing electron donating group. Compound **4b** having Bromo substituent in its structure shows lowest IC₅₀ value (IC₅₀ = 0.41 ± 0.01 μM) as compared to standard oleanolic acid (IC₅₀ = 13.45 ± 0.01 μM). It means compound **4b** shows better elastase activity than the standard oleanolic acid. Along with this compounds **4c**, **4d** **4e** and **4f** (IC₅₀ = 7.64 ± 0.42, 7.66 ± 0.77, 3.66 ± 0.55 and 7.23 ± 0.64 μM) also contains electron withdrawing group in its structure possesses lower IC₅₀ value than the standard. Compound **4g** – **4j** contains electron donating group in its structure. Here compound **4g** exceptionally shows lower IC₅₀ value (IC₅₀ = 0.99 ± 0.08 μM) than electron withdrawing group containing compounds except **4b** (IC₅₀ = 0.41 ± 0.01 μM). Compounds **4h**, **4i** and **4j** (IC₅₀ = 8.71 ± 0.66, 11.38 ± 0.99 and 7.83 ± 0.77 μM)



Scheme 1. Multicomponent Synthesis of 4H-chromene-3-carboxylates.

Table 2
Library of newly synthesized 4*H*-chromene-3-carboxylates.

Entry	Code	Structure of compound	Time in min	Yield %
1	4a		120	87
2	4b		100	87
3	4c		110	88
4	4d		95	88
5	4e		100	89
6	4f		115	81
7	4g		125	78
8	4h		130	76
9	4i		120	77

Table 2 (continued)

Entry	Code	Structure of compound	Time in min	Yield %
10	4j		125	79

Reaction conditions: Respective salicylaldehyde (1 mmol), Ethyl Acetoacetate (1 mmol), Dimedone (1 mmol), Catalyst: 20 mol % K_3PO_4 , Solvent: Ethanol (10 mL), Temperature: 80 °C

Table 3
Elastase (Elastase from porcine pancreas) inhibitory activity of 4a-4j derivatives.

Compound	Elastase $IC_{50} \pm SEM$ (μM)
4a	9.86 \pm 0.89
4b	0.41 \pm 0.01
4c	7.64 \pm 0.42
4d	7.66 \pm 0.77
4e	3.66 \pm 0.55
4f	7.23 \pm 0.64
4g	0.99 \pm 0.08
4h	8.71 \pm 0.66
4i	11.38 \pm 0.99
4j	7.83 \pm 0.77
Oleanolic Acid	13.45 \pm 0.01

SEM = Standard error of the mean; values are expressed in mean \pm SEM.

shows comparatively higher IC_{50} value than compound 4c, 4e and 4f but lower than the standard oleanolic acid ($IC_{50} = 13.45 \pm 0.01 \mu M$). Amongst the all synthesized and screened compounds, 4b found to be more potent against the elastase inhibition. Therefore, it can be concluded that all 4*H*-chromene-3-carboxylate moiety bearing electron withdrawing groups on its structure possibly interacts more with the enzyme.

2.2.2. Kinetic Mechanism:

The inhibition kinetic study was performed to understand the inhibitory mode of synthetic compounds against elastase. Based upon IC_{50} results, the most potent compound 4b was selected to determine their inhibition type and inhibition constant. The kinetic study of the enzyme involves the examination of Lineweaver-Burk plot of $1/V$ versus substrate *N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide $1/[S]$ in the presence of different inhibitor concentrations resulted into a series of straight lines (Fig. 1A). The Lineweaver-Burk plot of compound 4b showed that V_{max} remains same without significantly affecting the slopes. Whereas, K_m increases with increasing concentration and V_{max} remains the same with insignificant difference. This behavior indicates that compound 4b inhibits the enzymes in competitive manner (Fig. 1A). The enzyme Inhibition dissociation constant (K_i) was determined by plotting slope against inhibitor concentration of 4b (Fig. 1B), which was found to be 0.6 μM .

2.2.3. Free radical scavenging:

All the synthesized 4*H*-chromene-3-carboxylates (4a-4j) were assessed for their DPPH free radical scavenging ability at the concentration of (100 $\mu g/mL$). The results were presented in Fig. 2. In comparison with standard vitamin C, all synthesized 4*H*-chromene-3-carboxylate compounds (4a-4j) shows excellent free radical scavenging activity.

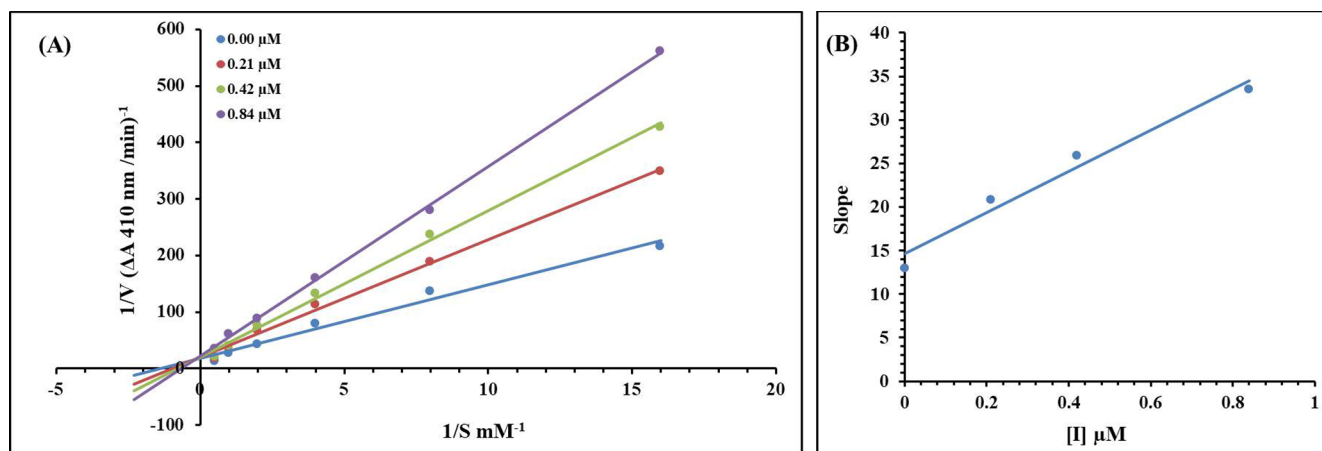


Fig. 1. Lineweaver-Burk plot for inhibition of elastase from porcine pancreas in the presence of Compound **4b**. (A) Concentrations of **4b** were 0.00, 0.21, 0.42 and 0.84 μM , Substrate *N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide concentrations were 2, 1, 0.5, 0.25, 0.125 and 0.0625 mM, respectively. (B) The insets represent the plot of the slope versus inhibitor **4b** concentrations to determine inhibition constant.

2.2.4. Chemoinformatics properties of newly synthesized 4*H*-chromene-3-carboxylates (**4a** – **4j**):

The basic Chemoinformatics properties of all synthesized compounds (**4a**–**4j**) were predicted by using computational tools. The predicted properties like molecular weight (Mol. Wt. g/mol), number of hydrogen bond acceptor and donors (No. HBA/HBD), LogP, polar surface area (PSA, A^2) and molecular volume (Mol. Vol, A^3) were shown in Table 4. The Mol. Wt (g/mol) and number of HBA and HBD in all synthesized compounds were found to be comparable with the standard values < 500 (g/mol) Mol. Wt. and < 10 HBA with < 5 HBD, respectively. Though **4f** exhibited greater Mol. Wt. (609 g/mol) compared to standard results, it follows Lipinski's rule. It is well known that the exceed values of HBA and HBD results in poor permeation [37,38]. The hydrogen-bonding capacity has been considered as significant parameter for drug permeability. Our results justified that the all synthesized compounds (**4a**–**4j**) possess < 10 HBA and < 5 HBD values which were comparable with standard values. Additionally, the analyzed data for synthesized compounds showed that PSA is very helpful parameter for drug absorption prediction in drug discovery [37,38]. The reported study showed that molecules with poor absorption are more likely to be observed to violate the Lipinski's rule. Multiple

Table 4

Predicted Chemoinformatics properties of synthesized compounds **4a** – **4j**.

Compounds	Mol. Wt. (g/mol)	No. HBA	No. HBD	Mol. LogP	Mol PSA (A^2)	Mol. Vol (A^3)
4a	358	5	1	2.9	57	452
4b	434	5	1	5.1	56	440
4c	390	5	1	5.0	56	435
4d	511	5	1	5.8	56	460
4e	424	5	1	5.6	56	450
4f	609	5	1	5.7	56	477
4g	372	6	2	4.0	74	428
4h	386	6	1	4.4	64	450
4i	370	5	1	4.7	56	439
4j	400	6	1	4.7	63	467

examples are available for RO5 violation amongst the existing drugs [39–40].

2.2.5. Docking Analysis:

2.2.5.1. Structural evaluation of target protein. Porcine pancreatic elastase consists of 240 amino acids having calcium atom embedded

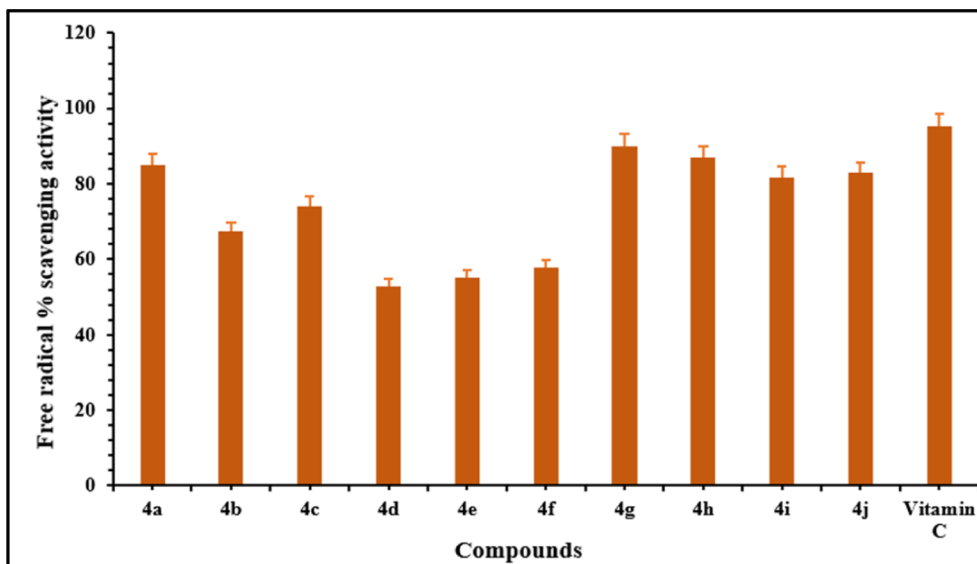


Fig. 2. Free radical scavenging activity of synthesized compounds. Values were represented as mean \pm SEM. All compounds concentrations were 100 $\mu\text{g}/\text{mL}$.

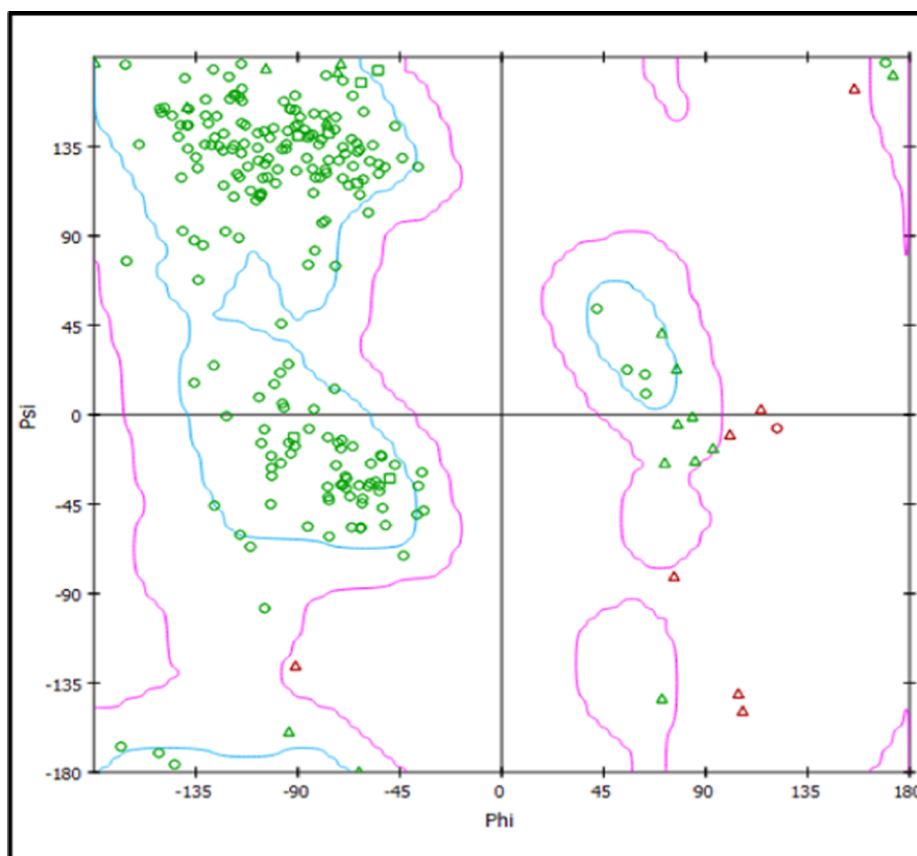


Fig. 3. Ramachandran plot of porcine pancreatic elastase.

in the protein structure. The x-ray diffraction study showed its resolution: 1.8 Å having unit cell coordinates $a = 52.530$, $b = 57.470$ and $c = 75.260$ with angles $\alpha = 90.00$, $\beta = 90.00$ and $\gamma = 90.00$, respectively. The Ramachandran graph values also showed 88.70% of residues were present in favored region while 98.70% residues were lie in allowed region (Fig. 3).

2.2.5.2. Docking energy analyses of synthesized compounds: To predict the best fitted conformational position of synthesized compounds (4a-4j) within the active region of targeted protein. The predicted docked complexes were evaluated on the basis of minimum energy values (kcal/mol) and interaction pattern. Docking results illustrates that all compounds showed good energy values ranging from -7.5 to -8.5 Kcal/mol. The docking energy values of all the docking complexes were calculated by using equation (1).

$\Delta G_{\text{binding}}$

$$= \Delta G_{\text{Gauss}} + \Delta G_{\text{repulsion}} + \Delta G_{\text{Hbond}} + \Delta G_{\text{hydrophobic}} + \Delta G_{\text{tors}}$$

(1)

where, ΔG_{Gauss} : attractive term for dispersion of two gaussian functions, $\Delta G_{\text{repulsion}}$: square of the distance if closer than a threshold value, ΔG_{Hbond} : ramp function - also used for interactions with metal ions, $\Delta G_{\text{hydrophobic}}$: ramp function, ΔG_{tors} : proportional to the number of rotatable bonds.

Present docking results for synthesized compounds (4a-4j) indicates the energy value difference among all docking complexes were lower than standard error value (Fig. 4). Therefore, based on the basis of both *in vitro* and *in silico* docking energy results, 4b is ranked as best ligand which shows good inhibitory potential against targeted enzyme as compared to all other derivatives.

2.2.5.3. Binding pocket and ligand conformation. The binding pocket

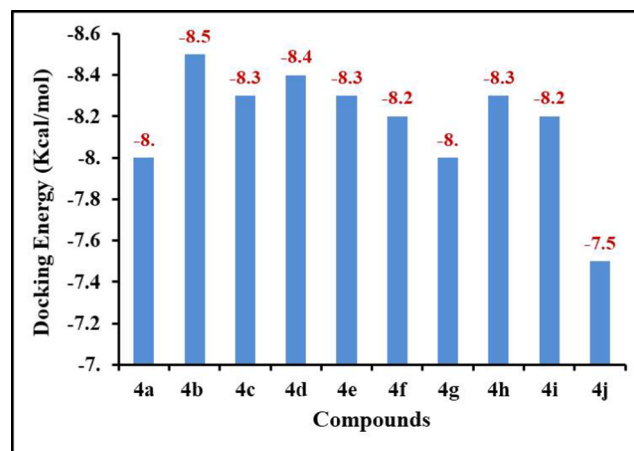


Fig. 4. Docking energy values of synthesized compounds.

analysis showed that compounds were confined in the active region of target protein (Fig. 5A). Results showed that compound 4b binds in good conformation in the binding pocket of target protein. The methyl part (CH_3) showed their entrance inside the binding pocket, whereas halogen moiety was present at the opening region of target protein. This incorporation may results in suitable configuration and conformation to ligand to be fitted in the binding pocket of elastase (Fig. 5B).

2.2.5.4. Hydrogen binding analysis. The docked complexes were analyzed on the basis of hydrogen and hydrophobic bonding interactions. Based on *in-vitro* result compound 4b was selected to check best conformational position inside active region of target protein. The compound 4b forms two hydrogen bonds at two

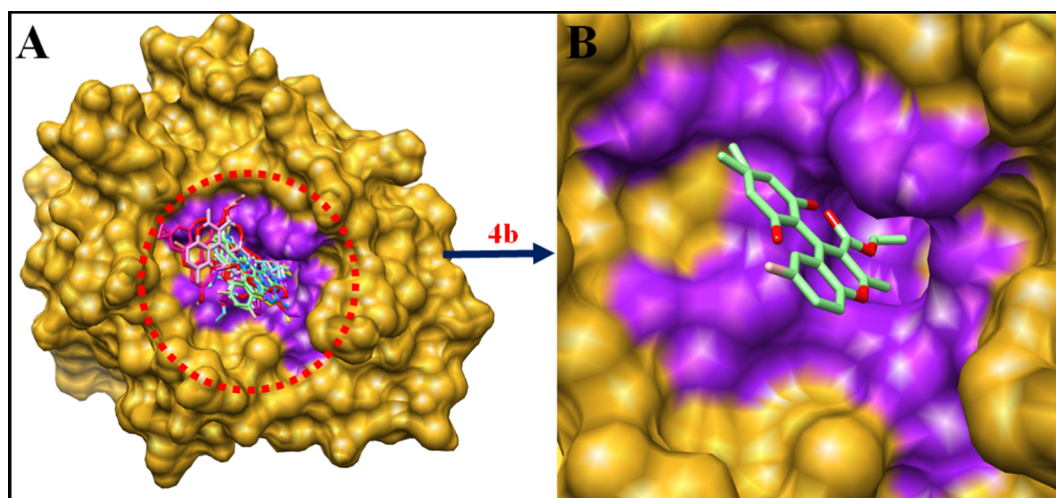


Fig. 5. Binding pocket of elastase (A) with embedded compound 4b (B).

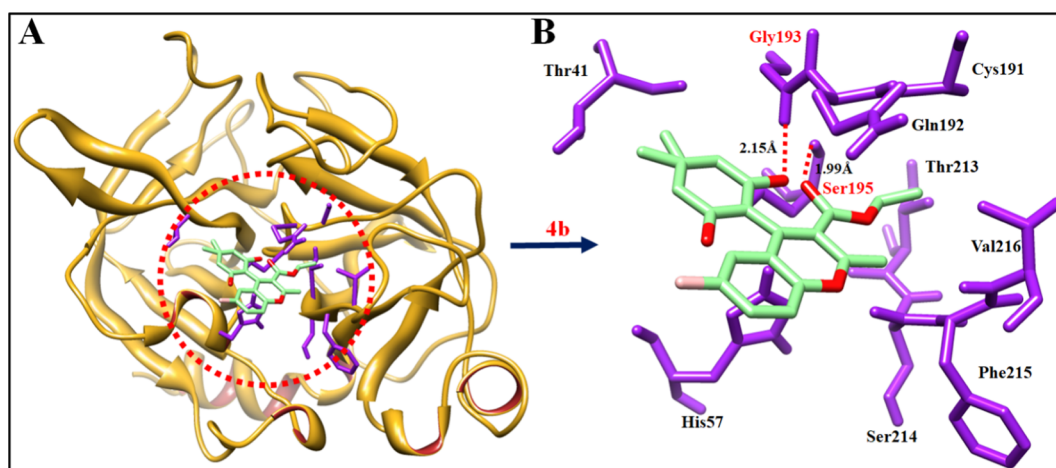


Fig. 6. Molecular docking interaction of 4b against receptor molecule. (A) The general overview of docking depiction. The protein structure is represented in dark brown color in ribbon format while ligand is highlighted in cyan color. (B) The closer view of binding pocket interaction with best conformation position of ligand against target protein. The interacted amino acids are highlighted in purple color and red dotted lines with distance mentioned in angstrom (\AA) are justified for hydrogen and bond distances.

different residues against target protein. The oxygen atoms of benzene ring form hydrogen bonds with Gly193 and Ser195 with bond length 2.15 and 1.99 \AA , respectively. Literature data also ensured the importance of these residues in bonding with other elastase inhibitors which strengthen our docking results [40,41]. The docking depiction is mentioned in Fig. 6A and Fig. 6B.

3. Conclusion:

The successful multicomponent synthesis of 4*H*-chromene-3-carboxylates was performed using K_3PO_4 as catalyst in ethanol at 80 $^\circ\text{C}$. The noteworthy features of this protocol are easy and simple workup procedure, high atom economy, excellent substrate scope, less reaction time and high yield of 4*H*-chromene-3-carboxylates. It is noteworthy that all synthesized compounds (4a – 4j) possessing superior elastase inhibitor activity. Particularly, compound 4b ($\text{IC}_{50} = 0.41 \pm 0.01 \mu\text{M}$) exhibits excellent activity as compared to standard Oleanolic Acid ($\text{IC}_{50} = 13.45 \pm 0.01 \mu\text{M}$). In addition, DPPH assay signposted excellent antioxidant properties for all the synthesized carboxylates. The *in silico* molecular docking investigation is in support of *in vitro* findings. Overall, from the results on biological activity and molecular docking studies of synthesized compounds 4a – 4j, it can be concluded that these compounds can be utilized as leading medicinal templates in

molecular drug design for elastase inhibition and skin disorder.

4. Experimental:

4.1. Chemistry

4.1.1. Materials and Method

Various substituted salicylaldehydes, Ethyl Acetoacetate, Dimezone and Potassium phosphate (K_3PO_4) were purchased from Sigma-Aldrich, Korea and used as received without further purification. Digimelt (SRS, USA) melting point apparatus was used to measure melting point of synthesized carboxylates. The formation of carboxylates (4a–4j) was confirmed by spectral techniques such as IR, NMR and LCMS Analysis. IR spectra were recorded on a Frontier IR Perkin–Elmer spectrophotometer. NMR spectra were recorded on a Bruker Avance III 600 MHz FT-NMR spectrometer in $\text{DMSO-}d_6$ using tetramethylsilane as an internal standard. The LCMS spectra were recorded on a Bruker MicroTof-Q spectrometer (Germany) coupled with Dionex Ultimate 3000/LC 09,115,047 (USA).

4.1.2. General synthetic procedure for the synthesis of 4*H*-chromene-3-carboxylates (4a – 4j)

The 50 mL of round bottom flask equipped with respective

substituted salicylaldehyde (1 mmol), ethyl acetoacetate (1 mmol) and K_3PO_4 (20 mol%) in ethanol (10 mL). The reaction mixture was refluxed at 80 °C till formation of Knoevenagel adduct. Then an equivalent amount of dimedone (1 mmol) was added in it and the resultant reaction mixture was further refluxed at 80 °C till the completion of reaction. The progress of reaction was monitored using thin layer chromatography. After completion of reaction, the reaction mixture was cooled and ice water was added in it. The expected product was precipitated out which was separated just by simple filtration. The crude product was again recrystallized using hot methanol in order to receive the product in its pure form. The formation of expected product was confirmed by spectral techniques such as IR, 1H & ^{13}C NMR and LCMS analysis.

4.1.2.1. Ethyl 4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4a). Off White powder; M.P.: 208 °C; IR: 3175, 2968, 1726, 1641, 1592, 1520, 1362, 1260, 1232, 1189, 1082, 1008, 850, 756, 680 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.36 (s, 1H, -OH), 7.08 – 7.11 (m, 1H), 6.92 – 6.99 (m, 3H), 5.05 (s, 1H), 2.52–2.55 (d, 2H, $J = 18$ Hz), 2.22 – 2.35 (q, 2H), 2.03 – 2.05 (d, 2H, $J = 18$), 1.05 (S, 3H), 0.98 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR (150 MHz, DMSO- d_6): δ 169.20, 195.55, 165.16, 150.13, 128.86, 127.30, 126.11, 124.60, 115.78, 11.28, 50.93, 43.47, 41.17, 32.06, 29.63, 28.20, 26.73 ppm; LCMS (ESI): 355.1765 (M – 1) m/z .

4.1.2.2. Ethyl 6-bromo-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4b). Cream color powder; M.P.: 225 °C; IR: 3118, 2961, 1733, 1614, 1474, 1364, 1250, 1180, 1136, 1079, 1039, 1011, 982, 969, 834, 820, 755, 658 cm^{-1} ; 1H NMR (600 MHz, DMSO- d_6): δ 10.54 (s, 1H, -OH), 7.26 – 7.28 (m, 1H), 6.93 – 7.04 (m, 2H), 5.03 (s, 1H), 2.58 (s, 2H), 2.23 – 2.34 (q, 2H), 2.02 – 2.05 (d, 2H, $J = 18$), 1.04 (S, 3H), 0.97 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR (150 MHz, DMSO- d_6): δ 196.08, 195.40, 164.92, 158.43, 154.10, 149.45, 146.06, 137.04, 133.00, 131.04, 130.15, 128.78, 125.93, 120.54, 118.87, 118.20, 116.81, 115.90, 110.94, 50.85, 41.00, 32.06, 30.43, 29.56, 28.08, 26.64 ppm; LCMS (ESI): 435.0804 M^+ and 347.0784 ($M + 2$) m/z

4.1.2.3. Ethyl 6-chloro-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4c). Light yellow powder; M.P.: 212 °C; IR: 2960, 1736, 1716, 1621, 1575, 1479, 1373, 1282, 1182, 1074, 1039, 1012, 933, 877, 836, 822 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.56 (s, 1H, -OH), 7.14 – 7.16 (m, 1H), 7.99 – 7.00 (d, 1H, $J = 6$ MHz), 6.90 – 6.91 (d, 1H, $J = 6$ MHz), 5.04 (s, 1H), 2.53–2.59 (m, 2H), 2.23 – 2.35 (q, 2H), 2.03 – 2.05 (d, 2H, $J = 12$ MHz), 1.05 (S, 3H), 0.97 (S, 3H), 0.90 (S, 6H) ppm; ^{13}C NMR (150 MHz, DMSO- d_6): δ 196.09, 164.90, 153.69, 149.00, 146.13, 134.28, 129.98, 128.32, 128.08, 128.00, 127.27, 118.61, 117.76, 110.86, 50.86, 41.01, 32.04, 29.60, 28.11, 26.64 ppm; LCMS (ESI): 391.1307($M + 1$) m/z

4.1.2.4. Ethyl 6,8-dibromo-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4d). Beige powder; M.P.: 231 °C; IR: 3175, 2960, 1747, 1667, 1646, 1598, 1563, 1449, 1373, 1313, 1246, 1206, 1181, 1149, 1038, 1011, 852, 764, 699 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.71 (s, 1H, -OH), 8.22 – 8.25 (m, 1H), 7.62 – 7.63 (d, 1H, $J = 6$ MHz), 7.01 – 7.02 (d, 1H, $J = 6$ MHz), 5.06 (s, 1H), 2.55–2.60 (m, 2H), 2.25 – 2.27 (d, 2H, $J = 12$ MHz), 2.03 – 2.06 (d, 2H, $J = 18$ MHz), 1.06 (S, 3H), 0.97 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.05, 195.09, 164.53, 157.76, 150.92, 146.54, 145.77, 138.84, 132.77, 132.65, 130.56, 130.14, 126.49, 121.53, 116.85, 115.93, 111.31, 110.73, 50.82, 40.80, 32.13, 30.37, 29.53, 28.07, 26.54 ppm; LCMS (ESI): 514.12 M^+ m/z

4.1.2.5. Ethyl 6,8-dichloro-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4e). Off White powder;

M.P.: 207 °C; IR: 3174, 2961, 1746, 1667, 1646, 1600, 1450, 1378, 1313, 1257, 1207, 1190, 1150, 1040, 1026, 1012, 886, 854, 764 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.72 (s, 1H, -OH), 7.41 (s, 1H), 6.86 (s, 1H), 5.06 (s, 1H), 2.54–2.60 (m, 2H), 2.25 – 2.28 (d, 2H, $J = 18$ MHz), 2.04 – 2.07 (d, 2H, $J = 18$ MHz), 1.06 (S, 3H), 0.97 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 197.26, 196.06, 164.63, 164.41, 145.13, 129.78, 127.91, 127.40, 127.01, 121.21, 111.18, 50.82, 40.79, 32.11, 32.04, 29.52, 29.03, 28.04, 26.54 ppm; LCMS (ESI): 425.0918 M^+ m/z

4.1.2.6. Ethyl 4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-6,8-diiodo-2-methyl-4H-chromene-3-carboxylate (4f). Beige powder; M.P.: 223 °C; IR: 3181, 2959, 2872, 1738, 1645, 1590, 1551, 1443, 1373, 1312, 1254, 1242, 1206, 1177, 1147, 1010, 980, 872, 854, 764, 694, 677 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.64 (s, 1H, -OH), 7.85 (s, 1H), 7.18 (s, 1H), 5.01 (s, 1H), 2.56 – 2.59 (t, 2H, $J = 6$ & 12 MHz), 2.35–2.38 (d, 2H, $J = 18$ MHz), 2.24 – 2.27 (d, 2H, $J = 18$ MHz), 2.02 – 2.05 (d, 2H, $J = 18$ MHz), 1.06 (S, 3H), 0.97 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.00, 164.79, 159.35, 154.14, 152.06, 149.60, 143.77, 137.26, 129.55, 126.10, 121.17, 111.56, 88.86, 86.82, 50.85, 40.87, 32.14, 30.43, 29.59, 28.06, 26.56 ppm; LCMS (ESI): 608.0570 M^+ m/z

4.1.2.7. Ethyl 7-hydroxy-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4g). Light Brown powder; M.P.: 234 °C; IR: 3174, 2953, 2885, 1744, 1619, 1588, 1515, 1451, 1368, 1308, 1263, 1231, 1172, 1143, 1097, 1032, 1012, 973, 844, 814, 769, 662 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.20 (s, 1H, -OH), 9.34 (s, 1H, -OH), 6.73 – 6.74 (d, 1H, $J = 6$ MHz), 6.40 – 6.41 (d, 1H, $J = 6$ MHz), 6.31 (s, 1H), 4.92 (s, 1H), 2.29 – 2.31 (d, 2H, $J = 12$ MHz), 2.20–2.22 (d, 2H, $J = 12$ MHz), 2.00 – 2.02 (d, 2H, $J = 12$ MHz), 1.03 (S, 3H), 0.97 (S, 3H), 0.89 (S, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.27, 195.62, 165.07, 156.57, 150.54, 129.23, 116.40, 112.24, 111.66, 102.35, 50.97, 43.55, 41.22, 32.02, 29.62, 28.21, 26.78 ppm; LCMS (ESI): 373.1644($M + 1$) m/z

4.1.2.8. Ethyl 4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-6-methoxy-2-methyl-4H-chromene-3-carboxylate (4h). Light Beige powder; M.P.: 222 °C; IR: 3192, 2942, 1728, 1634, 1591, 1497, 1468, 1422, 1373, 1318, 1265, 1223, 1199, 1152, 1130, 1027, 1012, 855, 814, 746, 662 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.32 (s, 1H, -OH), 6.88 – 6.89 (d, 1H, $J = 6$ MHz), 6.67 – 6.69 (q, 1H), 6.46 (s, 1H), 5.02 (s, 1H), 3.63 (s, 3H), 2.50 – 2.53 (d, 2H, $J = 18$ MHz), 2.21–2.32 (q, 2H), 2.01 – 2.03 (d, 2H, $J = 12$ MHz), 1.04 (S, 3H), 0.98 (S, 3H), 0.90 (S, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.15, 195.65, 165.33, 159.05, 156.28, 156.08, 149.62, 147.30, 144.30, 127.06, 125.12, 122.82, 119.06, 117.70, 116.54, 113.24, 112.79, 112.70, 110.51, 56.30, 55.66, 50.93, 41.22, 32.03, 30.48, 29.65, 28.13, 26.73 ppm;

4.1.2.9. Ethyl 4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2,6-dimethyl-4H-chromene-3-carboxylate (4i). White powder; M.P.: 213 °C; IR: 3104, 2962, 2942, 1735, 1626, 1613, 1585, 1494, 1372, 1307, 1245, 1223, 1202, 1172, 1153, 1076, 1041, 1011, 934, 886, 818, 781, 657 cm^{-1} ; 1H NMR(600 MHz, DMSO- d_6): δ 10.29 (s, 1H, -OH), 6.89 – 6.91 (q, 1H), 6.82 – 6.83 (d, 1H, $J = 6$ MHz), 6.75 (s, 1H), 5.01 (s, 1H), 2.31 – 2.33 (d, 2H, $J = 12$ MHz), 2.22–2.25 (d, 2H, $J = 18$ MHz), 2.16 (s, 3H), 2.01 – 2.04 (d, 2H, $J = 18$ MHz), 1.05 (S, 3H), 0.98 (S, 3H), 0.90 (S, 3H), 0.89 (s, 3H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.18, 165.25, 148.15, 133.38, 128.98, 127.82, 125.78, 115.54, 111.15, 50.96, 43.50, 41.21, 32.04, 29.65, 28.13, 26.71, 20.81 ppm; LCMS (ESI): 369.2051($M - 1$) m/z

4.1.2.10. Ethyl 8-ethoxy-4-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-2-methyl-4H-chromene-3-carboxylate (4j). Light yellow powder; M.P.: 142 °C; IR: 2960, 2871, 2565, 1729, 1634, 1612, 1582, 1566,

1471, 1371, 1332, 1316, 1257, 1227, 1207, 1195, 1132, 1089, 1074, 1027, 886, 788, 760, 727 cm^{-1} ; ^1H NMR(600 MHz, DMSO- d_6): δ 10.31 (s, 1H, -OH), 6.88 (t, 1H, $J = 12$ MHz), 6.77–6.78 (d, 1H, $J = 6$ MHz), 6.51–6.53 (d, 1H, $J = 12$ MHz), 5.03 (s, 1H), 4.03–4.05 (q, 2H), 2.56–2.59 (d, 1H, $J = 18$ MHz), 2.21–2.37 (q, 2H), 2.02–2.04 (d, 2H, $J = 12$ MHz), 1.35 (t, 3H, $J = 6$ & 12 MHz), 1.05 (s, 3H), 0.99 (s, 3H), 0.89 (s, 6H) ppm; ^{13}C NMR(150 MHz, DMSO- d_6): δ 196.23, 158.68, 147.73, 146.39, 139.86, 126.83, 125.32, 124.17, 122.12, 120.19, 117.88, 111.25, 65.06, 64.32, 56.50, 32.10, 30.43, 29.60, 28.12, 26.70, 19.01, 15.20 ppm; LCMS (ESI): 400.2220 M^+ m/z

4.2. Biology:

4.2.1. In vitro Methodology:

4.2.1.1. Elastase inhibition assay: The elastase (Elastase from porcine pancreas) inhibition activity was performed by following already reported method [42–45] with few modifications. In order to perform the inhibition of elastase activity, the amount of released *p*-nitroaniline, which was hydrolyzed from the substrate (*N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide) by elastase, was determined by measuring the absorbance at 410 nm. In detail, 0.8 mM solution of *N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide was prepared in a 0.2 M Tris-HCl buffer (pH 8.0) and this buffer (130 μL) was added to the test sample (10 μL) in a 96 well microplate. The microplate was pre-incubated for 10 min at 25 $^\circ\text{C}$ before an elastase (0.0375 Unit/mL) stock solution (10 μL) was added. After enzyme addition, the microplate was kept at 25 $^\circ\text{C}$ for 30 min, and the absorbance was measured at 410 nm using microplate reader (SpectraMax ABS, USA). IC_{50} values were calculated by nonlinear regression using GraphPad Prism 5.0 (GraphPad, San Diego, CA USA). All experiments were carried out in triplicate. The elastase inhibition activities were calculated according to the following formula:

$$\text{Elastaseinhibitionactivity(\%)} = \frac{\text{OD}_{\text{Control}} - \text{OD}_{\text{Sample}}}{\text{OD}_{\text{Control}}} \times 100 \quad (2)$$

where, $\text{OD}_{\text{control}}$ and $\text{OD}_{\text{sample}}$ represents the optical densities in the absence and presence of sample, respectively. Oleanolic acid was used as the standard inhibitor for elastase.

4.2.1.2. Kinetic analysis of inhibition of elastase: By following our reported method [46] Kinetic analysis was carried out to determine the mode of inhibition. The Compound **4b** was selected on the basis of most potent IC_{50} values. Kinetic study was carried out by varying the concentration of *N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide in the presence of different concentrations of compound **4b** (0.00, 0.21, 0.42 and 0.84 μM). Briefly the *N*-succinyl-Ala-Ala-Ala-*p*-nitroanilide concentration was changed from 2, 1, 0.5, 0.25, 0.125 and 0.0625 mM for its kinetics studies and remaining procedure was same for all kinetic studies as describes in elastase inhibition assay protocol. Maximal initial velocities were determined from initial linear portion of absorbance's up to 10 min after addition of enzyme at per minute interval. The inhibition type on the enzyme was evaluated by Lineweaver-Burk plot of inverse of velocities ($1/V$) versus inverse of substrate concentration $1/[S]$ mM^{-1} . The Enzyme Inhibition dissociation constant K_i was determined by secondary plot of $1/V$ versus inhibitor concentration. The results were processed by using SoftMaxPro.

4.2.1.3. Free radical scavenging assay. Radical scavenging activity was determined by modifying already reported method [47,48] by 2, 2-diphenyl-1 picrylhydrazyl (DPPH) assay. The assay solution consisted of 100 μL of DPPH (150 μM), 20 μL of increasing concentration of test compounds and the volume was adjusted to 200 μL in each well with methanol. The reaction mixture was then incubated for 30 min at room temperature. Ascorbic acid (Vitamin C) was used as a reference inhibitor. The assay measurements were carried out by using a microplate reader (OPTI MAX , Tunable) at 517 nm. The reaction rates

were compared and the percent inhibition caused by the presence of tested inhibitors was calculated. Each concentration was analyzed in three independent experiments.

4.2.2. In silico analysis: Computational Methodology:

4.2.2.1. Retrieval of porcine pancreatic elastase structure. The crystal structure of porcine pancreatic elastase was accessed from Protein Data Bank (PDB) (www.rcsb.org) with PDBID 7EST. The retrieved target protein structure was minimized by using UCSF Chimera 1.6rc [49]. The stereo-chemical properties of elastase and Ramachandran plot were also accessed from PDB [51].

4.2.2.2. Chemo-informatics properties of synthesized compounds and molecular docking. The synthesized compounds were evaluated on the basis of chemo-informatics properties [37–41,49–52]. Molinspiration (<http://www.molinspiration.com/>), and Molsoft (<http://www.molsoft.com/>) online servers were used to predict the chemo-informatics and biological properties of ligands, respectively. Lipinski's rule of five was analyzed using Molsoft. Before docking experiment all the synthesized chemical structures (**4a-4j**) were sketched in ACD/ChemSketch tool and accessed in mol format. Furthermore, UCSF Chimera 1.10.1 tool was employed to energy minimization of each ligand separately having default parameters such as steepest descent steps 100 with step size 0.02 (\AA), conjugate gradient steps 100 with step size 0.02 (\AA) and update interval was fixed at 10. Finally, Gasteiger charges were added using Dock Prep in ligand structure to obtain the good structure conformation. Molecular docking experiment was performed on all the compounds, against elastase by using virtual screening tool PyRx with VINA Wizard approach [53]. The grid box parameters values in VINA search space ($X = 4.6898$, $Y = 57.6015$ and $Z = -5.7255$) were adjusted with default exhaustiveness value = 8 to maximize the binding conformational analysis. We have adjusted sufficient grid box size on binding pocket residues to allow the ligand to move freely in the search space. All the synthesized compounds were docked separately against target protein. In all docked complexes, the ligands conformational poses were keenly observed to obtain the best docking results. The generated docked complexes were evaluated on the basis of lowest binding energy (Kcal/mol) values and binding interaction pattern between ligands and receptor. The graphical depictions of all the docked complexes were accomplished by UCSF Chimera 1.10.1 and Discovery Studio (2.1.0), respectively.

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Conflict of interest

Authors declare no conflict of interest.

Appendix A. Supplementary data

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