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4 **Mechanistic evaluation of a novel cyclohexanone derivative's**
5 **functionality against nociception and inflammation:**
6 **an *in-vitro*, *in-vivo* and *in-silico* approach.**

7
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39 **Highlights**

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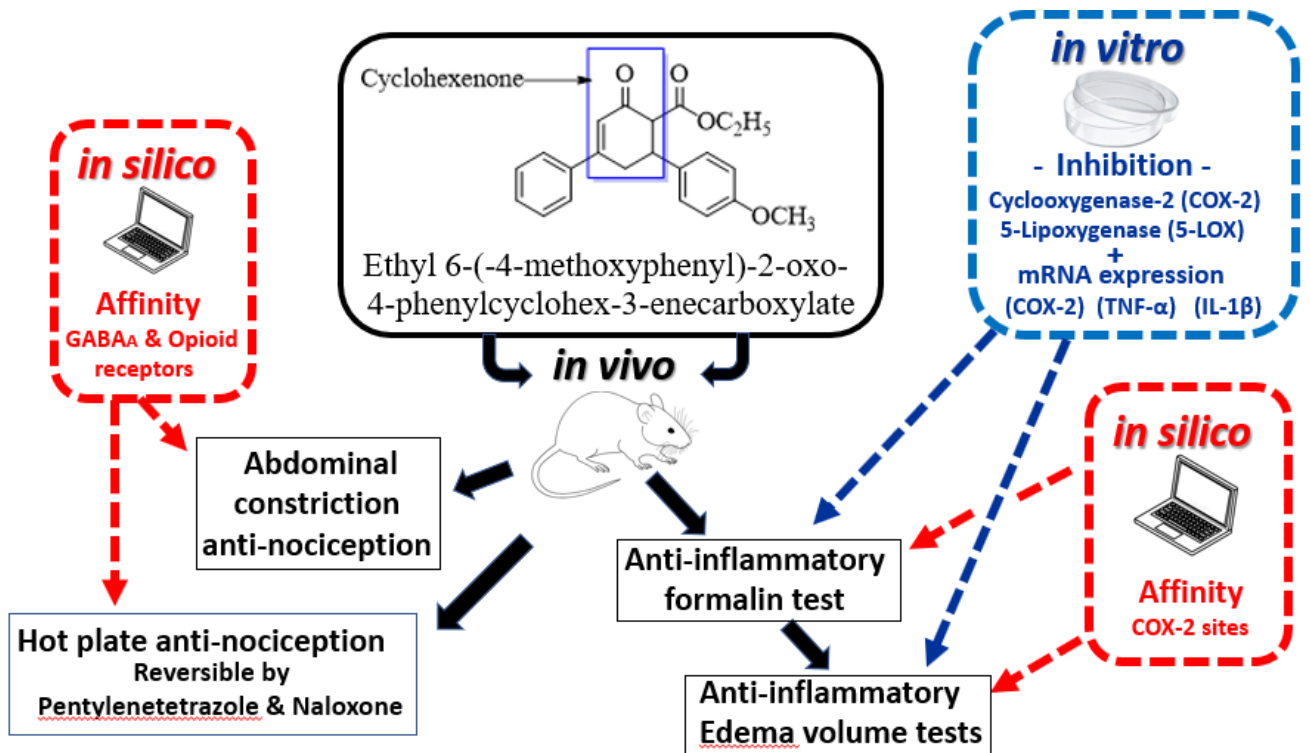
- 41 -A synthesised anti-inflammatory cyclohexanone (CHD) was tested *in vivo* and *in vitro*
- 42 -CHD inhibited COX-2 and 5-LOX enzymes plus COX-2, TNF- α and IL-1 β mRNA expression
- 43 -CHD also produced GABA_A and opioid mediated inhibitory activity in nociceptive tests
- 44 -In silico CHD had preferential affinity for GABA_A, opioid and COX-2 target sites
- 45 -CHD may possess therapeutic effectiveness in the management of inflammation and pain

46

47 **Graphic Abstract**

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52

53 **Abstract**

54 The synthesis of a novel cyclohexanone derivative (CHD; Ethyl 6-(4-metohxyphenyl)-2-oxo-4-
55 phenylcyclohexe-3-enecarboxylate) was described and the subsequent aim was to perform an *in*
56 *vitro*, *in vivo* and *in silico* pharmacological evaluation as a putative anti-nociceptive and anti-
57 inflammatory agent in mice. Initial *in vitro* studies revealed that CHD inhibited both
58 cyclooxygenase-2 (COX-2) and 5-lipoxygenase (5-LOX) enzymes and it also reduced mRNA
59 expression of COX-2 and the pro-inflammatory cytokines TNF- α and IL-1 β . It was then shown
60 that CHD dose dependently inhibited chemically induced tonic nociception in the abdominal
61 constriction assay and also phasic thermal nociception (i.e. anti-nociception) in the hot plate and
62 tail immersion tests in comparison with aspirin and tramadol respectively. The thermal test
63 outcomes indicated a possible moderate centrally mediated anti-nociception which, in the case of
64 the hot plate test, was pentylenetetrazole (PTZ) and naloxone reversible, implicating GABAergic
65 and opioidergic mechanisms. CHD was also effective against both the neurogenic and
66 inflammatory mediator phases induced in the formalin test and it also disclosed anti-inflammatory
67 activity against the phlogistic agents, carrageenan, serotonin, histamine and xylene compared with
68 standard drugs in edema volume tests. *In silico* studies indicated that CHD possessed preferential
69 affinity for GABA_A, opioid and COX-2 target sites and this was supported by molecular dynamic
70 simulations where computation of free energy of binding also favored the formation of stable
71 complexes with these sites. These findings suggest that CHD has prospective anti-nociceptive and
72 anti-inflammatory properties, probably mediated through GABAergic and opioidergic interactions
73 supplemented by COX-2 and 5-LOX enzyme inhibition in addition to reducing pro-inflammatory
74 cytokine expression. CHD may therefore possess potentially beneficial therapeutic effectiveness
75 in the management of inflammation and pain.

76

77 **Keywords**

78 Cyclohexenone, Anti-nociceptive, Anti-inflammatory, Cyclooxygenase-2, 5-lipoxygenase,
79 GABA_A/opioid receptors

80

81

82 **1. Introduction**

83 The process of drug discovery and development incorporating a novel chemical moiety with a
84 desirable therapeutic profile is a challenging task nowadays (DiMasi et al., 2010). Extensive
85 research has been carried out on pain and inflammation over a number of years, particularly
86 because these pathological conditions can greatly influence patient quality of life (Ali et al., 2015;
87 Chapman and Gavrin, 1999; Shahid et al., 2017a; Shahid et al., 2017b). Pathologically, pain may
88 be categorized as nociceptive, neuropathic or inflammatory and if protracted, it may progress into
89 a chronic pain syndrome involving additional symptoms such as anxiety and depression.
90 Nociceptive pain is typically initiated by stimulation of somatic sensory receptors designated as
91 nociceptors, which then transmit pain impulses to the central nervous system (CNS). Alternatively,
92 neuropathic pain arises from damage or lesions to the nervous system (Van Hecke et al., 2014).
93 Active inflammation is the hallmark of inflammatory pain and is characterized by the presence of
94 inflammatory mediators such as interleukin, TNF- α , prostaglandins (PGE₂, PGI₂, TXA₂),
95 histamine, serotonin, bradykinin and leukotrienes (LTs) (Fernandes et al., 2015). These
96 biochemical substances produce changes in neuronal sensitivity and invoke the onset of tissue
97 hypersensitivity associated with inflammation (Kidd and Urban, 2001). Currently, opioids and
98 non-steroidal anti-inflammatory drugs (NSAIDs) are the analgesic agents of choice often utilized
99 in the management of inflammatory pain. However, it is well documented that persistent use of
100 NSAIDs may well cause deleterious effects such as ulceration, hemorrhage or even perforation in
101 the gastrointestinal tract, cardiovascular system disorders and kidney damage (Gutthann et al.,
102 1996; Jones et al., 2008). Similarly, opioid analgesics are considered highly effective as analgesics,
103 but they are associated with dependence liability and other side effects which may limit their

104 usefulness (Laxmaiah Manchikanti et al., 2010; Mayer et al., 1995; Shahid et al., 2016).
105 Consequently, there is a genuine need for substitute drugs that retain the analgesic and anti-
106 inflammatory effectiveness of conventional analgesic agents without their untoward effects
107 (Fawad et al., 2018; Islam et al., 2017; Islam et al., 2019).

108 The key role of the cyclohexenone ring is well established in the field of biomedical research. It
109 has been documented that this functionality is an integral part of several interesting compounds
110 and is of considerable significance for the development of potentially valuable drugs (Das and
111 Manna, 2015; Fang et al., 2012). Chemically, the cyclohexenone nucleus, serves as a convenient
112 intermediate for synthesizing various heterocyclic compounds including fused pyrazoles,
113 isoxazoles, quinazolines (Senguttuvan and Nagarajan, 2010) and 2H-indazole (Gopalakrishnan et
114 al., 2008). Cyclohexenones are cyclohexane derivatives with a carbonyl group at position-1 and a
115 carbon-carbon double bond at position-2 (Fig. 1). The enone functional group and substitution at
116 a carbon atom in the six membered ring have been used to synthesize other substituted
117 cyclohexenones (Johnson et al., 2016). The pharmacological properties of cyclohexenone
118 derivatives include anti-inflammatory and anti-nociceptive effects (Ahmadi et al., 2012; Lednicer
119 et al., 1981a; Lednicer et al., 1981b; Liu et al., 2013; Ming-Tatt et al., 2013; Ming-Tatt et al., 2012;
120 Sheorey et al., 2016; Wang et al., 2011) as well as anti-neuropathic and antioxidant activity (Khan
121 et al., 2019). The present study was undertaken to evaluate a novel cyclohexenone derivative
122 (CHD; Ethyl 6-(4-metohxyphenyl)-2-oxo-4-phenylcyclohexe-3-enecarboxylate) as a possible
123 inhibitor of cyclooxygenase-2 (COX-2) and 5-LOX pro-inflammatory enzymes and subsequently
124 examine its effects against nociception using *in vivo* mouse models of pain and inflammation.
125 Additionally, the anti-nociceptive activity of CHD was also investigated in the presence of

126 pentylenetetrazole (PTZ) and naloxone in order to probe any possible underlying mechanisms,
127 which might have been corroborated by *in silico* and *in vitro* studies.

128

129 **2. Material and methods**

130 *2.1. Chemicals and drugs*

131 Naloxone, serotonin, histamine, PTZ, xylene, indomethacin, lambda carrageenan and aspirin were
132 purchased from (Sigma-Aldrich, USA). Formaldehyde was procured from Merck (Germany),
133 glacial acetic acid was obtained from Pancreac (Spain), tramadol (Tramal[®] 50mg/ml) was
134 acquired from Searle Ltd (Pakistan). Fresh preparation of chalcone was carried out in the
135 laboratory of ICS (University of Peshawar, Pakistan). Ethyl acetoacetate, ethyl acetate and
136 potassium carbonate were purchased from Merck (Pakistan). N-hexane and ethanol were procured
137 from Scharlau (Lahore, Pakistan). The cDNA synthesis kit, TRIzol reagent, master mix and
138 primers were acquired from Thermofishcer Scientific (USA).

139 *2.2. Chemistry*

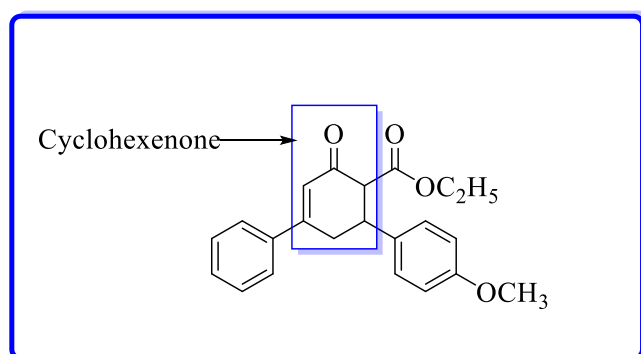
140 *2.2.1. General*

141 A Gallenkamp melting point apparatus was used to determine melting points. Purity was checked
142 by thin layer chromatography (TLC). A Shimadzu IR Prestige-21 FT-IR Spectrometer Instrument
143 (Tokyo, Japan) was utilized to record the Infrared spectra. ¹³C and ¹H NMR analyses (Agilent AV-
144 300,400 and 500 Tokyo, Japan) were accomplished with D₂O and DMSO-d₆ as solvents. Mass
145 spectra (ESI-MS) were obtained on (Qp 2010 plus, Shimadzu, Tokyo, Japan). Perkin Elmer 2400
146 CHN/O Analyzer was operated to determine Elemental analysis.

147

148 2.2.2. Synthesis of Ethyl 6-(4-methoxyphenyl)-2-oxo-4-phenylcyclohex-3-enecarboxylate

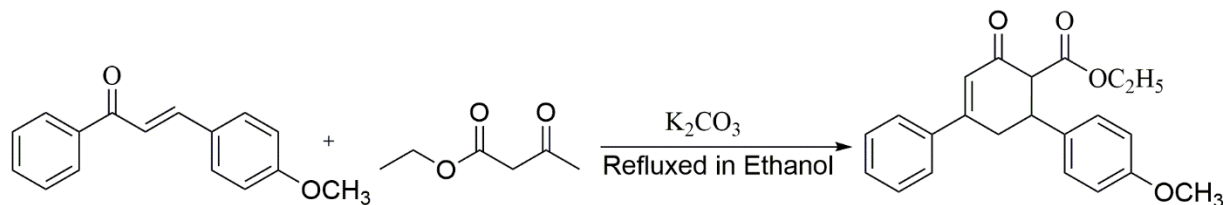
149 The synthesis was conducted according to the synthetic protocol as shown in Scheme 1. (E)-3-(
150 4-methoxyphenyl)-1-phenylprop-2-en-1-one (10 mmol) was refluxed with ethyl acetoacetate (20
151 mmol) in the presence of K_2CO_3 catalyst in 20 ml of ethanol for 3 hr. The product obtained was
152 recrystallized from ethanol; a brownish yellow powder was obtained having a yield of 85 %. M.p
153 = 92-95 $^{\circ}C$; R_f = 0.51 *n*-Hexane/ethyl acetate (7:3); IR (KBr) ν_{max} cm^{-1} : 3077 (Ar-H), 1689
154 (ketone C=O), 1735 (Ester C=O) 2870 (Aliphatic C-H); 1H -NMR ($CDCl_3$, 400MHz) δ :6.9-7.5 (m,
155 Ar-H),3.05 (d, 2H, J= 2.3), 2.9 (t,1H J=5.0, C-3), 2.6-2.8(q, 5H, CH_2CH_3 , J=7.0); ^{13}C -NMR (100
156 MHz, $CDCl_3$) δ : 199.0 (C=O), 125-130 (Ar-CH), 112 (C-6), 40.2 (OCH_3), 159.0 (C-19), and 44.39
157 (C-3). EI-MS; m/z (rel. int. %) 351 (M+), CHN Anal. Calcd for: C, 75.41; H, 6.33; O, 18.26.
158 Found: C, 74.81; H, 6.38. Formula: $C_{22}H_{22}O_4$, C=22, H=22, and O=4.



160 **Fig. 1.** Chemical structure of Ethyl 6-(4-methoxyphenyl)-2-oxo-4-phenylcyclohex-3-
161 enecarboxylate

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164
 165 **Scheme 1.** Synthetic scheme of Ethyl 6-(4-methoxyphenyl)-2-oxo-4-phenylcyclohexe-3-
 166 enecarboxylate

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168 2.3. *In vitro* activities

169 2.3.1. 5-LOX inhibition assay

170 The inhibitory potential of CHD was examined by utilizing human recombinant 5-LOX. In this
 171 assessment, the enzyme inhibition was determined through residual enzyme potential following
 172 10 to 15 min incubation at 25 °C in an incubator (Jan et al., 2020; Wisastra et al., 2013). The
 173 activity was estimated through linoleic acid (lipoxygenase substrate) conversion into hydroperoxy-
 174 octadecadienoate (HPOD). The alteration rate was calculated in the form of absorbance at 234 nm
 175 with UV- visible spectrophotometer. Ethylene diamine tetra acetic acid (EDTA 2 mM) and $CaCl_2$
 176 (2 mM) containing Tris buffer (50 mM) of PH 7.5 was used as an assay buffer for this assay. The
 177 enzyme 5-LOX (20,000 U/ml) was diluted with buffer in a ratio of 1:4000. The assay buffer was
 178 then diluted with 100 mM of inhibitor formerly blended with DMSO. Linoleic acid was then
 179 diluted with ethyl alcohol to 20 mM. Subsequently, various concentrations of CHD ranging from
 180 31.25 to 1000 μ g/ml and 1 ml of enzyme solution (1:4000) was mixed with 100 μ l adenosine
 181 triphosphate (2 mM), 790 μ l of Tris buffer plus 100 μ l inhibitor (1mM) and then incubated for ten
 182 min duration. Then to this mixture was added 10 μ l of substrate solution (20mM) and after ten
 183 seconds mixing of the enzyme with substrate, the substrate conversion rate was monitored. The

184 reaction rate in the absence of inhibitor was employed as positive control. The standard inhibitor
185 agent used in this assay was zileuton.

186 2.3.2. COX-2 Inhibition assay

187 The COX-2 inhibitory activity of the test compound was evaluated according to a previously
188 validated procedure (Burnett et al., 2007; Jan et al., 2020). A COX-2 enzyme solution (300 U/ml)
189 was prepared. For activation, 10 µl of enzyme solution was kept for 5 to 6 min on ice (4 °C) and
190 then mixed with 50 µl of co-factor solution comprising 0.9 mM glutathione, 1 mM hematin in 0.1
191 mM Tris buffer (pH 8.0) and 0.24 mM tetramethyl-p-phenylenediaminedihydrochloride (TMPD).
192 Then various concentrations (31.25 to 1000 µg/ml) of test sample (20 µl) plus enzyme solution
193 (60 µl) were maintained at room temperature for 5 to 10 min, followed by initiation of the reaction
194 by adding 30 mM arachidonic acid (20 µl) and keeping this mixture at 37 °C for a duration of 15
195 min. Afterwards, the reaction was terminated by addition of hydrochloric acid (HCL) and
196 absorbance was measured via a UV-visible spectrophotometer at 570 nm. COX-2 percentage
197 inhibition was calculated from the absorbance value per unit time. In the study Celecoxib was
198 utilized as the standard inhibitor agent.

199 2.3.3. Reverse transcription polymerase chain reaction (RT-PCR)

200 Post-mortem mouse paw sub plantar tissues were removed 5 h after carrageenan administration
201 and RNA was extracted using TRIzol reagent according to the manufacturer's protocol. The total
202 RNA was reverse transcribed to cDNA following a standard protocol. The primers for targeted
203 genes use were;

204 **COX-2:** F-5'-GGAGAGACTATCAAGATAGTGATC -3', R- 5'- ATGGTCAGTAGA-CTTT-
205 TACAGCTC-3'. **TNF-α:** F-5'-CTTCTCCTTCCTGATCGTGG-3'; R-5'-GCTGGTTAT-

206 CTCTCAGCTCCA-3'. **IL-1 β** : F-5'-AGAAGCTTCCACCAATACTC-3', R-5'-AGCACCTAG-
207 TTGTAAGGAAG-3'. **GAPDH**: F-5'-TGCACCACCAACTGCTTAGC-3'; R- 5'-GGCATG-
208 GACTGTGGTCATGAG3' was used as a housekeeping gene (Cheon et al., 2009; Khalid et al.,
209 2018). Amplified products were separated using 1.5% Agarose gel electrophoresis, analyzed with
210 image J software (Almeer et al., 2019; Ullah et al., 2021).

211 *2.4. In vivo pharmacological evaluation*

212 *2.4.1. Animals*

213 Mice (Balb-C) of either sex weighing 18-30 g were used during the investigation unless otherwise
214 stated. Animals were maintained on standard laboratory food and water *ad libitum* at an ambient
215 temperature of $22 \pm 2^\circ\text{C}$ through a thermostatically controlled air conditioning system on a 12/12
216 h light and dark cycle and they were habituated to laboratory conditions for two h before
217 experiments.

218 *2.4.2. Ethical approval*

219 The study and all *in vivo* protocols were conducted under a project entitled “Studies on the
220 nociceptive, inflammatory and neuropathic pain relieving potential of a cyclohexenone
221 derivative.” It was approved by the Research Ethical Committee of the Department of Pharmacy,
222 University of Peshawar, Pakistan which issued a certificate number of 01/EC/18/Pharm.
223 Furthermore, animal experiments were performed in compliance with the Animals Scientific
224 Procedure Act UK (1986).

225

226

227

228 *2.4.3. Acute toxicity study of compound CHD*

229 The acute toxicity profile of CHD was evaluated after intraperitoneal (i.p) injection of selected
230 doses (on a sequential dose-doubling increasing scale viz 15, 30, 60, 120 or 240 mg/kg (n=6)).
231 Animals were observed at 30-60 min and 24-72 h for any abnormal behaviour (Akbar et al., 2016).

232 *2.4.4. Anti-nociceptive activity*

233 *2.4.4.1. Anti-nociceptive activity of compound CHD and a standard drug in the acetic acid*
234 *abdominal constriction test*

235 Food and water were withdrawn 120 min prior to animal experiments. One percent acetic acid (10
236 ml/kg) i.p injection was used to induce abdominal constriction as a reflection of tonic nociception.
237 Five min after acetic acid i.p injection, the incidence of abdominal constrictions was recorded over
238 a 20 min period (Abbas et al., 2011). The animals were randomly allocated to different
239 investigational groups (n=6). Group I received normal saline as vehicle, group II-IV received
240 standard aspirin (15-45 mg/kg), group V-VII received test compound (CHD) (15-45 mg/kg) via
241 i.p injection, 30 min prior to 1% acetic acid injection. Percentage analgesia was calculated using
242 the following formula:

243 $\% \text{ protection} = (1 - \text{mean number of abdominal constrictions of the treated drug} / \text{mean number}$
244 $\text{of abdominal constrictions of the vehicle control}) \times 100$

245 *2.4.4.2. Anti-nociceptive activity of compound CHD compared to a standard drug in the hot plate*
246 *test*

247 The hot plate analgesiometer, was kept at a constant temperature of $54 \pm 0.1^{\circ}\text{C}$. After placement on
248 the plate, animal nociceptive reaction latencies (s) were determined to the following escape end
249 points: paw licking, flinching or jumping and a 30 s cut off time was imposed after which mice

250 were removed from the stimulus (Ahmad et al., 2017; Rukh et al., 2020). Animals were randomly
251 assigned to groups (n=6) and administered saline or drug treatment intraperitoneally. Group I
252 received normal saline as vehicle, group II received standard drug (tramadol, 30 mg/kg, i.p),
253 groups III-V received the trial compound (CHD, 15-45 mg/kg, i.p).

254 *2.4.4.3. Pharmacological antagonism study of CHD compared to a standard drug*

255 In order to evaluate the possible involvement of GABA_A or opioid receptors in the anti-nociceptive
256 activity of CHD, mice were administered PTZ (15 mg/kg; i.p) or naloxone (1 mg/kg;
257 subcutaneously (s.c)) 10-20 min prior to i.p dosing with saline, CHD or standard drug. Hot plate
258 latencies were recorded 30,60 and 90 min after administration of each drug (Muhammad et al.,
259 2012). Percentage protection against nociception was determined using the following formula:

$$260 \text{ \% protection} = (\text{Test latency} - \text{baseline latency}) / (\text{cut off time} - \text{baseline latency}) \times 100$$

261 *2.4.4.4. Anti-nociceptive activity of CHD compared to a standard drug in the tail immersion test*

262 Each animal was gently held in a vertical position and half of the tail was immersed in a water
263 bath maintained at a temperature of 55±0.5 °C. A nociceptive reaction latency (s) was determined
264 to a tail flick end point and a cut off time of 15 s imposed after which animals were removed from
265 the stimulus. Any non-responders within the cut-off time were excluded from the study. The
266 vehicle, standard tramadol (30 mg/kg) and test compound (15-45 mg/kg) were administered i.p to
267 their respective groups. The readings were taken at 30, 60, 90 and 120 min after drug
268 administration (Sewell and Spencer, 1976).

269

270

271 *2.4.4.5. Anti-nociceptive activity of CHD and standard drug in the formalin induced biphasic pain*
272 *model*

273 Mice were administered a sub plantar injection of 20 μ l of freshly prepared 2 % formalin in the
274 right hind paw. Thirty min prior to formalin injection, groups I-VI, received intraperitoneally
275 normal saline as vehicle, standard drugs indomethacin (10 mg/kg) or diclofenac (10 mg/kg), and
276 CHD (15-45mg/kg). The nociceptive reaction time (s) (latency to biting, licking, paw lifting or
277 flinching) was measured in two phases: first phase (0 to 5 min) and second phase (10-30 min) after
278 the formalin injection (Silva et al., 2017; Maione et al., 2020).

279 *2.4.5. Anti-inflammatory activity*

280 *2.4.5.1. Anti-inflammatory activity of compound CHD and standard in a carrageenan induced paw*
281 *edema model*

282 Mice were treated with normal saline, aspirin (50-150 mg/kg) or test compound (CHD, 15-45
283 mg/kg, i.p) 30 min before s.c injection of 0.05 ml of freshly constituted carrageenan (1 %) in the
284 right hind paw. A digital Plethysmometer was utilized to determine the inflammation in terms of
285 paw edema volume (ml) at hourly intervals up to 5 h post carrageenan injection (Ali et al., 2013).

286 *2.4.5.2. Anti-inflammatory activity of compound CHD and standard drug in a histamine induced*
287 *paw edema model*

288 Inflammation was induced in mice (25-30 g) by sub plantar injection of 0.1 ml freshly constituted
289 histamine (1 mg/ml) in the right hind paw. Paw inflammation swelling was measured by means of
290 plethysmometer previously described in the carrageenan test (Mequanint et al., 2011).

291

292 *2.4.5.3. Anti-edema activity of CHD and a standard drug in the serotonin induced paw volume*
293 *model*

294 Mice were administered serotonin (0.001 mg/ml s.c) into the plantar surface of the right hind paw.
295 The ensuing inflammation and paw edema was measured by plethysmometer (Masresha et al.,
296 2012).

297 *2.4.5.4. Anti-inflammatory action of compound CHD and standard drug in the xylene provoked*
298 *ear edema model*

299 In mice weighing 25-35 g, ear edema was evoked by topical application of 0.03 ml of xylene to
300 the internal and outer surface of the right ear while the left ear was used as control. Thirty min
301 before induction of xylene edema, saline vehicle was administered i.p to group I, standard
302 indomethacin (10 mg/kg) or diclofenac (15 mg/kg) to groups II-III and test compound CHD (15-
303 45 mg/kg) to groups IV-VI respectively. Subsequently, 15 min after xylene application, animals
304 were killed and the ears were amputated then weighed. The mean weight difference between right
305 and left ears was then determined (Manouze et al., 2017).

306

307

308

309 *2.5. In silico activity*

310 *2.5.1. Docking studies*

311 Docking studies were executed through the Molecular Operating Environment (MOE) version
312 2016.08 docking program. Three-dimensional (3D) structures of the enzymes, GABA_A and opioid
313 receptors with their co-crystalized ligands were obtained from the Protein Data Bank as listed in

314 Table 1. The docking algorithm was validated by re-docking of native ligands as shown in Table
 315 1. The computed root mean square deviation (RMSD) between the experimental and re-docked
 316 poses was found within a threshold limit $< 2 \text{ \AA}$. The 3D structures of the compound were
 317 constructed in MOE by utilizing Builder Module. Energy minimization of the ligand, preparation
 318 of structures of the downloaded enzymes and active site identification was carried out according
 319 to our earlier validated methods (Iftikhar et al., 2017; Iftikhar et al., 2018; Rashid et al., 2016).
 320 Assessment of docking outcomes and scrutiny of their surface with graphical demonstrations were
 321 accomplished with discovery studio visualizer and MOE (Systemes, 2015).

322

323 **Table 1**

324 Protein Data Bank (PDB) code numbers, names of their co-crystallized ligands and resolution for
 325 the enzymes studied.

326

Enzyme/Receptor	PDB code	Co-crystallized ligand	Resolution (\AA)
COX-1 enzyme	1EQG	Ibuprofen	2.61
COX-2 enzyme	1CX2	1-Phenylsulfonamide-3-trifluoromethyl-5-parabromophenylpyrazole (SC-558)	3.00
GABA _A receptor	4COF	Benzamidine	2.97
μ -opioid receptor	4DKL	β -Funaltrexamine (μ -opioid receptor antagonist)	2.8
δ -opioid receptor	4EJ4	Naltrindole (δ -opioid receptor antagonist)	3.4
κ -opioid receptor	4DJH	(3R)-7-Hydroxy-N-[(2S)-1-[(3R,4R)-4-(3-hydroxyphenyl)-3,4-dimethylpiperidin-1-yl]-3-methylbutan-2-yl]-1,2,3,4-tetrahydroisoquinoline-3-carboxamide (JDC)	2.9

327

328

329

330 2.5.2. Molecular Dynamic Simulation of Complexes

331 Molecular Dynamic (MD) simulations were performed using the same protocol as explained in
332 our previous study (Abbasi et al., 2016). MD simulations facilitate understanding of the binding
333 pattern and determine the stability of selected receptor-CHD docked complexes. Using AMBER
334 18 software, six different systems were prepared to run MD simulations for 50 ns each (Case et
335 al., 2010). In order to verify the structural variations and convergence of the simulated systems,
336 the CPPTRAJ module of AmberTools18's was used to estimate the RMSDs for all the studied
337 systems.

338 2.5.2.1. Binding Free Energy Calculations

339 The MMPB/GBSA methods, integrated in AMBER 18, were employed to calculate the binding
340 free energies for all six systems (Miller III et al., 2012). Binding free energy calculations were
341 performed on 100 snapshots taken from the MD trajectories as described previously by Abro and
342 Azam (Abro and Azam, 2016). The binding free energy can be expressed as:

$$343 \Delta G_{\text{bind}} = \Delta G_{\text{complex}} - [\Delta G_{\text{receptor}} + \Delta G_{\text{ligand}}]$$

344 where ΔG is the Gibb's free energy calculated by MMGB/PBSA.

345 2.6. Statistical analysis

346 The data were analyzed statistically utilizing Graph Pad Prism Software, version 5, for manifold
347 assessments via one-way analysis of variance (ANOVA) with Post-hoc Dunnett's test. Outcomes
348 were regarded as statistically significant at $P < 0.05$.

349 3. Results

350 3.1. In vitro activities

351 All enzyme suppression results are presented as the mean of triplicate determinations for each
352 concentration studied and an IC_{50} value was extrapolated from the overall inhibitory concentration
353 relationships.

354

355

356 *3.1.1. 5-LOX inhibitory activity*

357 The 5-LOX inhibitory activity of CHD was examined at various concentrations ranging from 31.25
358 to 1000 $\mu\text{g/ml}$ and the compound displayed a potent inhibition of 5-LOX with an extrapolated IC_{50}
359 value of 10.27 $\mu\text{g/ml}$ as compared to the standard 5-LOX inhibitor drug zileuton (extrapolated
360 $\text{IC}_{50} = 5.50 \mu\text{g/ml}$) over the same tested concentration range (Table 2).

361 *3.1.2. COX-2 inhibitory activity*

362 CHD disclosed a potent inhibitory action on the COX-2 enzyme as shown in Table 3. It was also
363 evident from the outcomes that CHD possessed valuable COX-2 inhibitory activity in comparison
364 with the standard COX-2 inhibitor drug celecoxib. Thus, the IC_{50} value for CHD was extrapolated
365 as 8.94 $\mu\text{g/ml}$ in contrast to that of celecoxib ($\text{IC}_{50} = 4.30 \mu\text{g/ml}$) (Table 3).

366 **Table 2**

367 5-LOX enzyme inhibitory activity of CHD in comparison with zileuton as a standard 5-LOX
368 inhibitor drug.

Compound	Conc. ($\mu\text{g/ml}$)	% 5-LOX inhibition (Mean \pm S.E.M)	Extrapolated $\text{IC}_{50} \mu\text{g/ml}$
Cyclohexenone derivative (CHD)	1000	89.44 \pm 0.55 ^b	10.27
	500	83.17 \pm 0.72 ^c	
	250	78.30 \pm 0.64 ^c	
	125	73.34 \pm 0.63 ^c	
	62.5	68.30 \pm 0.64 ^c	
	31.25	61.93 \pm 1.13 ^c	
Zileuton	1000	93.55 \pm 0.40	5.50
	500	89.37 \pm 1.65	
	250	85.50 \pm 0.40	
	125	79.60 \pm 0.90	
	62.5	74.17 \pm 0.72	
	31.25	70.35 \pm 0.45	

369 Data is represented as mean \pm S.E.M; Values were significantly different as compared to the
370 positive control (zileuton); n=3, b= $P < 0.01$, c= $P < 0.001$.

371

372

373 **Table 3**

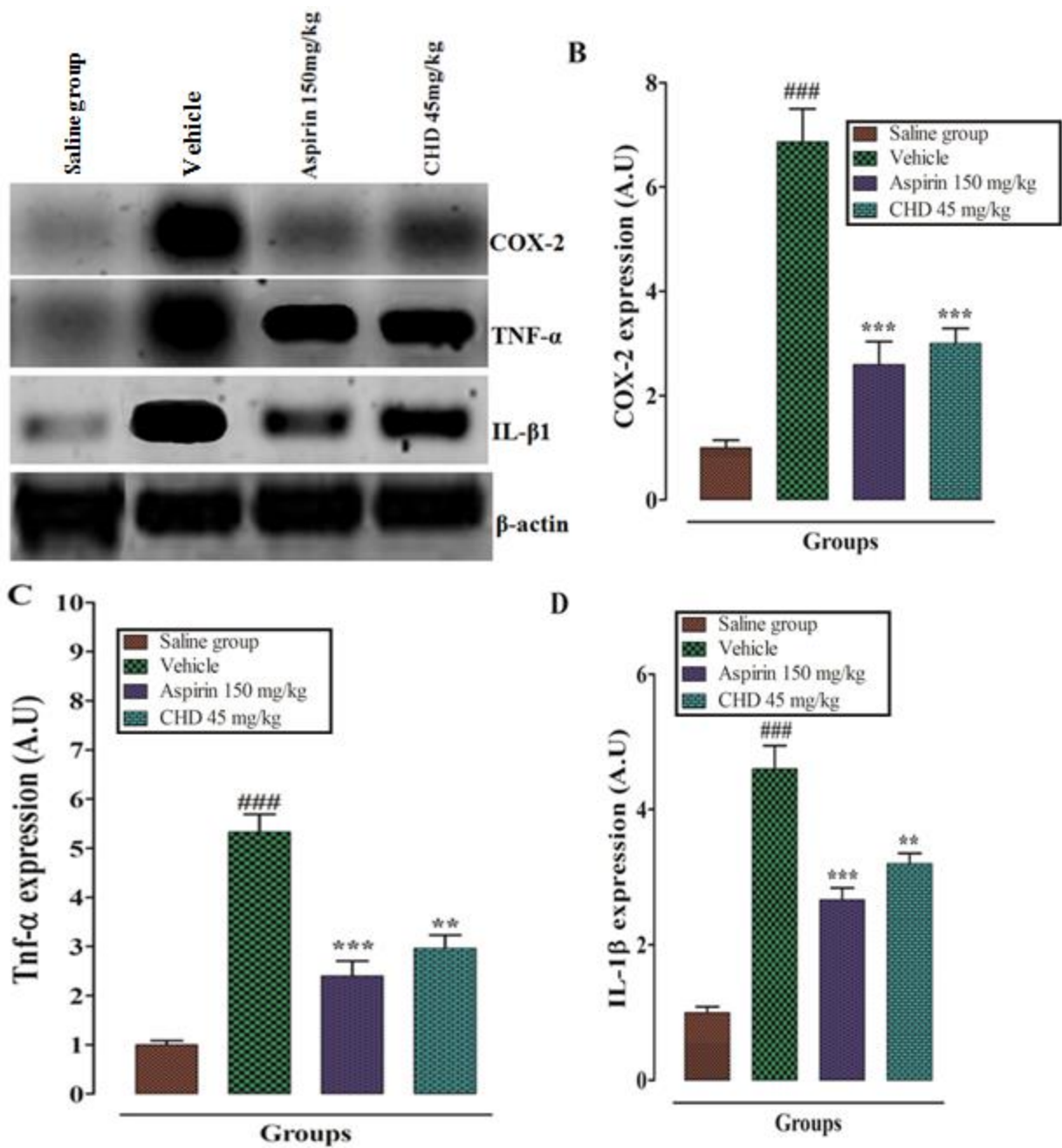
374 COX-2 enzyme inhibitory assay of CHD in comparison with celecoxib as a standard COX-2
 375 inhibitor drug.

Compound	Conc. ($\mu\text{g/ml}$)	% COX-2 inhibition (Mean \pm S.E.M)	Extrapolated IC ₅₀ $\mu\text{g/ml}$
Cyclohexenone derivative (CHD)	1000	88.91 \pm 1.30 ^c	8.94
	500	85.00 \pm 0.30 ^c	
	250	78.76 \pm 0.58 ^c	
	125	73.67 \pm 0.61 ^c	
	62.5	67.74 \pm 0.61 ^c	
	31.25	63.47 \pm 0.56 ^c	
Celecoxib	1000	95.20 \pm 0.15	4.30
	500	91.17 \pm 0.53	
	250	86.98 \pm 0.85	
	125	81.20 \pm 0.65	
	62.5	77.80 \pm 0.37	
	31.25	73.11 \pm 1.20	

376 Data is represented as mean \pm S.E.M; Values were significantly different as compared to the
 377 positive control (celecoxib); n=3, c= $P < 0.001$.

378 *3.1.3. RT- PCR*

379 To further investigate the anti-inflammatory potential of CHD, RT-PCR was utilized to assess the
 380 mRNA levels of COX-2 enzyme and the pro-inflammatory cytokines TNF- α and IL-1 β in the
 381 carrageenan induced paw edema test in mice. The outcomes of this assessment revealed that CHD
 382 (45 mg/kg) significantly reduced the mRNA expression of COX-2 ($P < 0.001$), while in the case of
 383 TNF- α and IL-1 β , CHD also produced a reduction ($P < 0.01$) compared to the carrageenan treated
 384 vehicle group. Aspirin (150 mg/kg) as the standard positive control decreased ($P < 0.001$) the
 385 expression of COX-2, TNF- α and IL-1 β as presented in (Fig. 2).



386
 387 **Fig. 2.** Agarose gel electrophoresis (A) quantification of CHD activity on the mRNA level of
 388 COX-2 (B), TNF- α (C), and IL-1 β (D) in carrageenan induced hind paw edema in mice. The results
 389 are shown in relative arbitrary units (A.U). Bars represent mean expression in A.U \pm S.E.M. ###
 390 $P < 0.001$ compared to the saline group. ** $P < 0.01$, *** $P < 0.001$ compared to the vehicle group.

391

392

393 *3.2. In vivo pharmacological activity*

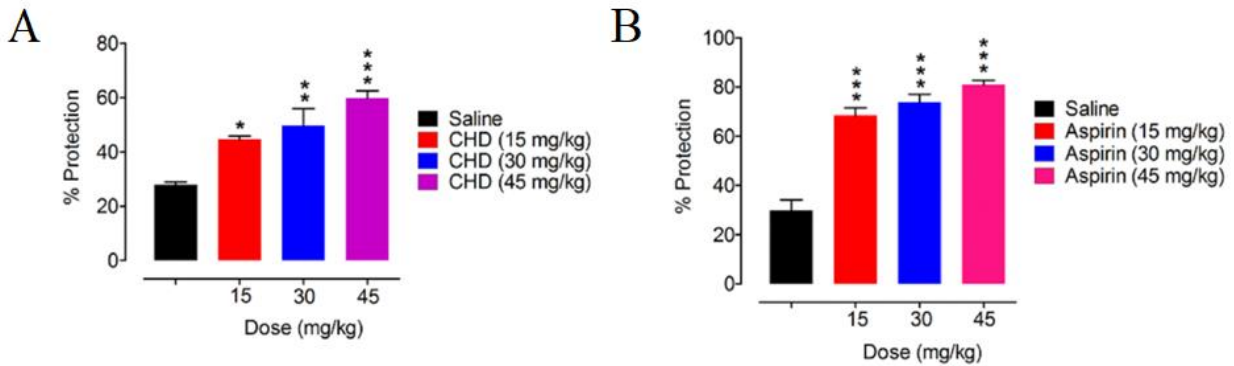
394 *3.2.1. Acute toxicity of CHD*

395 After i.p. injection of selected doses of CHD (15-240 mg/kg; $n = 6$), there was no acute toxicity
396 observed in gross animal behaviour, neither was any incidence of mortality recorded up to the
397 highest dose. Thus, the maximum tolerated dose (MTD) which was devoid of unacceptable toxicity
398 for CHD was >240 mg/kg.

399 *3.2.2. CHD attenuation of chemically induced tonic nociceptive behaviour*

400 Injection of acetic acid (1%) was accompanied by a significant rise in the nociceptive response
401 perceived as an onset increase in the incidence of abdominal constriction. The percentage
402 protection against this chemically induced tonic nociception in the group of animals treated with
403 CHD at a lower dose (15 mg/kg) decreased the nociceptive response as evidenced by an increase
404 in the percentage protection (44.66%, $P < 0.05$). Likewise, the mid-range CHD dose (30 mg/kg)
405 also protected against acetic acid evoked abdominal constriction (49.78%, $P < 0.01$). and the higher
406 dose (45 mg/kg) had an even greater anti-nociceptive effect (59.81%) reflecting dose dependent
407 activity relative to the saline treated animals. The aspirin positive control also yielded a dose
408 dependant anti-nociceptive response (15-45 mg/kg) versus the saline controls (Fig. 3).

409

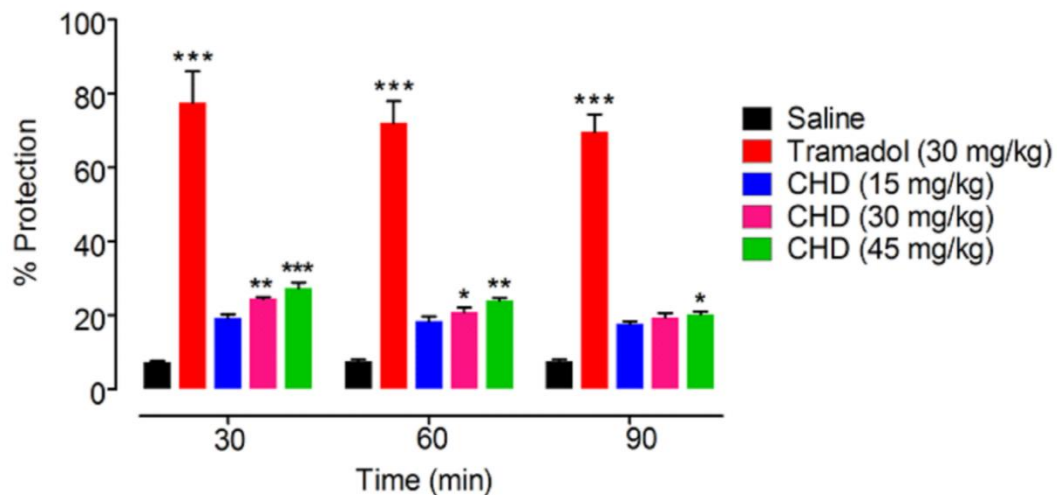


410

411 **Fig. 3.** Anti-nociceptive activity of (A) CHD and (B) the positive control, aspirin in the acetic acid
 412 (1%) induced abdominal constriction test. Each bar represents mean percentage protection \pm
 413 S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as compared to the saline treated group (one-way
 414 ANOVA followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

415 3.2.3. CHD attenuation of phasic thermal nociception

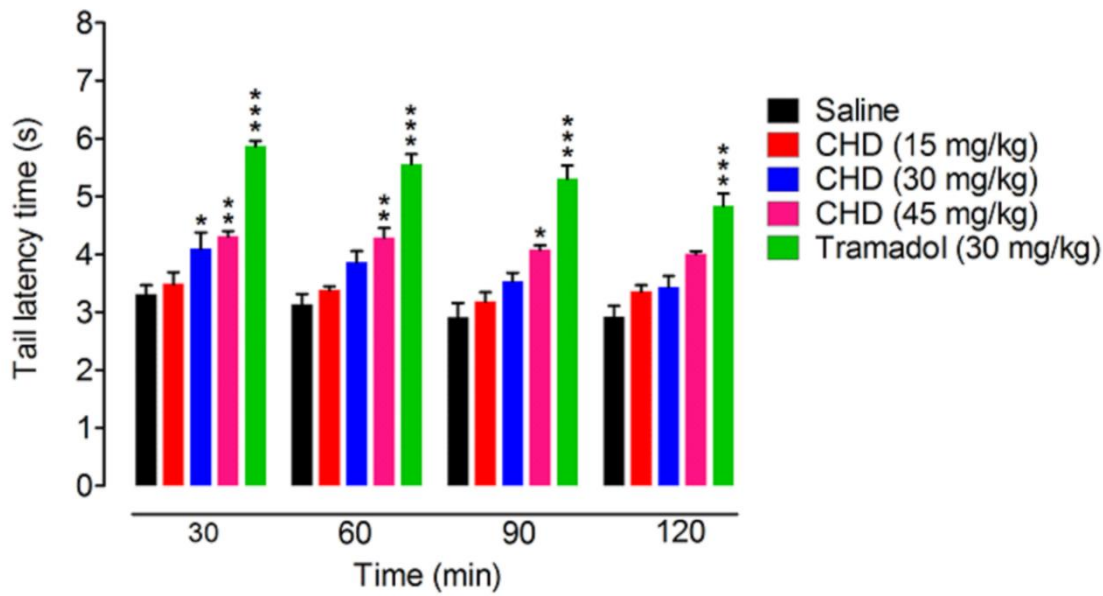
416 In the hot plate test, the saline treated animal group displayed a control escape response from the
 417 thermal nociceptive stimulus of 7.3%, 7.5% and 7.7% after 30, 60 and 90 min respectively. The
 418 lower dose of CHD was ineffective in producing any detectable anti-nociception between 30-90
 419 min (19.3%-17.6%). However, CHD at 30 mg/kg did produce an anti-nociceptive effect at 30 min
 420 (24.5%) and 60 min (20.8%) but this was not evident after 90 min (19.5%). A greater anti-
 421 nociceptive response was noted at the 45 mg/kg dose (27.3%, 24.0% and 20.3% at 30, 60 and 90
 422 min respectively) while the tramadol (30 mg/kg) positive control produced an even bigger response
 423 77.6%, 72.0% and 69.7% at 30, 60 and 90 min respectively (Fig. 4).



424
 425 **Fig. 4.** Anti-nociceptive activity of CHD and the positive control, tramadol, in the hot-plate test.
 426 Each bar represents mean percentage protection \pm S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as
 427 compared to saline treated group (one-way ANOVA followed by *post hoc* Dunnett's test), ($n = 6$
 428 mice per group).

429 *3.2.4. CHD attenuation of phasic nociception in the tail immersion test*

430 CHD produced a measurable anti-nociceptive response in the tail immersion test at the 30 mg/kg
 431 dose (30 min). However, the 45 mg/kg dose produced a peak response at 60 min which subsided
 432 by 120 min. Treatment with the positive tramadol control (30 mg/kg), produced an intense long-
 433 acting anti-nociceptive effect which lasted up to 120 min (Fig. 5).

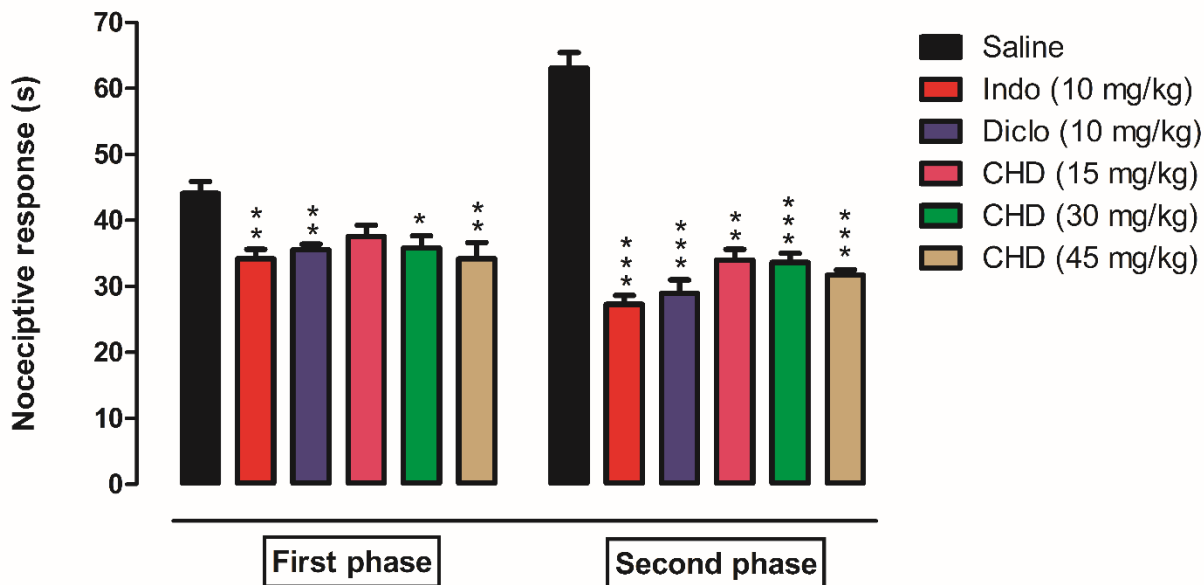


434
 435 **Fig. 5.** Anti-nociceptive activity of CHD and the positive control, tramadol in the thermal tail
 436 immersion test. Each bar represents mean withdrawal latency time in s \pm S.E.M). * $P < 0.05$, ** $P <$
 437 0.01, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
 438 Dunnett's test), ($n = 6$ mice per group).

439 *3.2.5. CHD attenuation of the formalin induced biphasic nociceptive response*

440 Administration of formalin in the sub-plantar mouse hind paw initiated a marked nociceptive
 441 response as indicated by an increase in the duration of biting, licking, lifting and flinching of the
 442 affected paw. This was observed throughout the first phase (0-5 min) and also the second phase
 443 (15-30 min) following formalin administration in the saline treated animals. Treatment with CHD
 444 (15 mg/kg) only diminished the second phase of formalin induced nociception. Conversely, the 30
 445 mg/kg CHD dose was more effective in that it markedly reduced the formalin nocifensive response
 446 in both the second and first phases ($P < 0.05$). Similarly, treatment with the higher CHD dose (45
 447 mg/kg) did induce an anti-nociceptive response in the first phase, but a more statistically

448 significant response in the second phase. The indomethacin and diclofenac positive controls both
449 at 10 mg/kg generated comparable anti-nociception to CHD in both phases (Fig. 6).

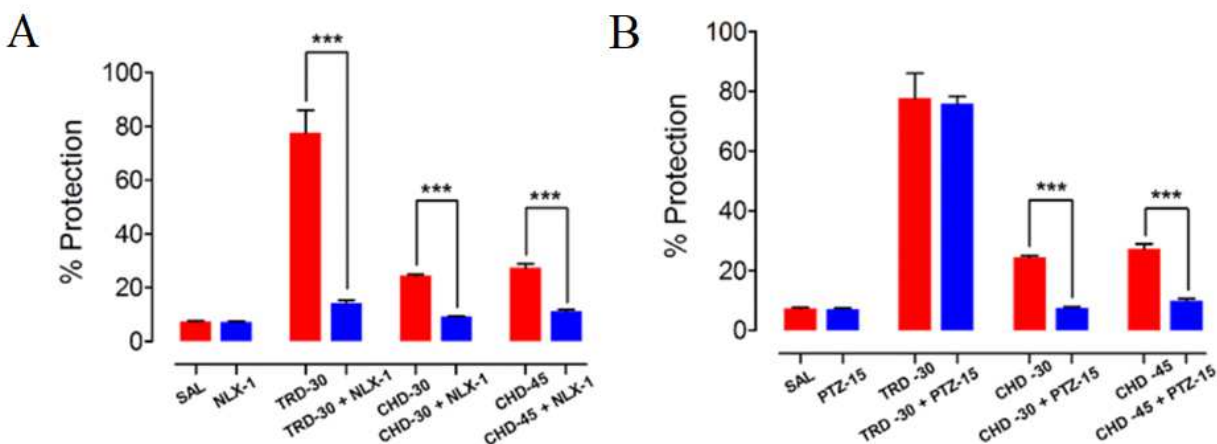


450
451 **Fig. 6.** Anti-nociceptive activity of CHD and the positive controls, indomethacin (Indo), and
452 diclofenac (Diclo) in the formalin induced paw nociceptive test. Each bar represents mean
453 noceptive response in s \pm S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as compared to the saline
454 treated group (one-way ANOVA followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

455 3.2.6. Opioidergic and GABAergic mediation of CHD anti-nociception

456 Any possibility of GABAergic or opioidergic mechanisms underlying the anti-nociceptive effect
457 of CHD in the hot-plate test were probed using pentylenetetrazole (PTZ) and naloxone as
458 respective antagonists. Hence, the anti-nociceptive effect of CHD (30 and 40 mg/kg), was
459 significantly antagonized ($P < 0.001$) by naloxone (1 mg/kg) implicating the involvement of an
460 opioidergic mechanism. Likewise, in animals treated with the opioid agonist, tramadol (30 mg/kg)
461 as a positive control, naloxone also blocked the anti-nociceptive response (Fig. 7A).
462 Administration of PTZ (15 mg/kg) did not modify the anti-nociceptive action of tramadol (30

463 mg/kg), but it did markedly decrease the anti-nociceptive response of CHD (30 and 45 mg/kg) in
 464 the hot plate paradigm. This would tend to suggest an involvement of a GABAergic mechanism
 465 in the anti-nociceptive action of CHD but not tramadol (Fig. 7B).

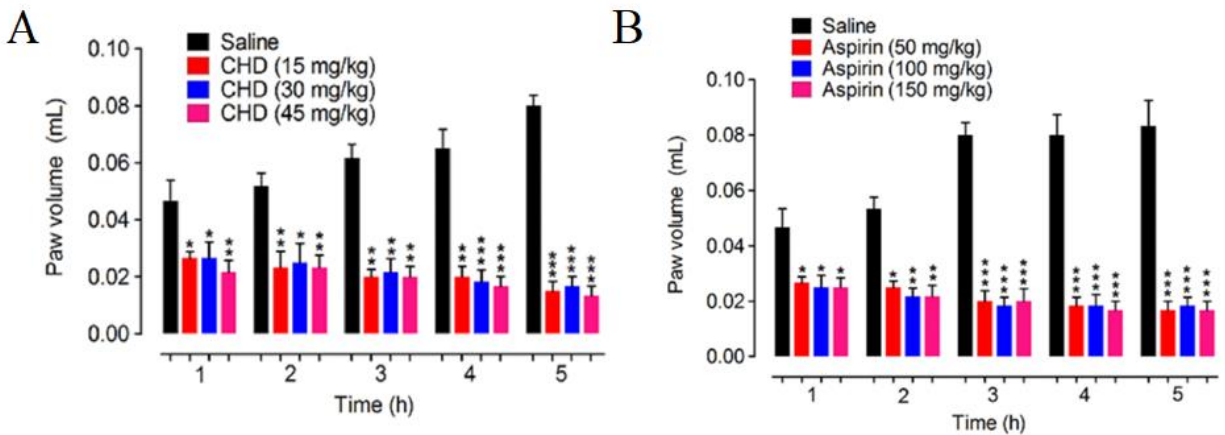


466
 467 **Fig. 7.** (A) Effect of naloxone at 1 mg/kg (NLX-1) and (B) PTZ at 15 mg/kg (PTZ-15) on the anti-
 468 nociceptive activity of CHD (30 mg/kg, CHD-30 and 45 mg/kg, CHD-45) or tramadol (30 mg/kg,
 469 TRD-30) in the mouse hot-plate test. Each bar represents mean percentage protection \pm S.E.M.
 470 *** $P < 0.001$ compared to saline control (SAL). (two sample t -test), ($n = 6$ mice per group).

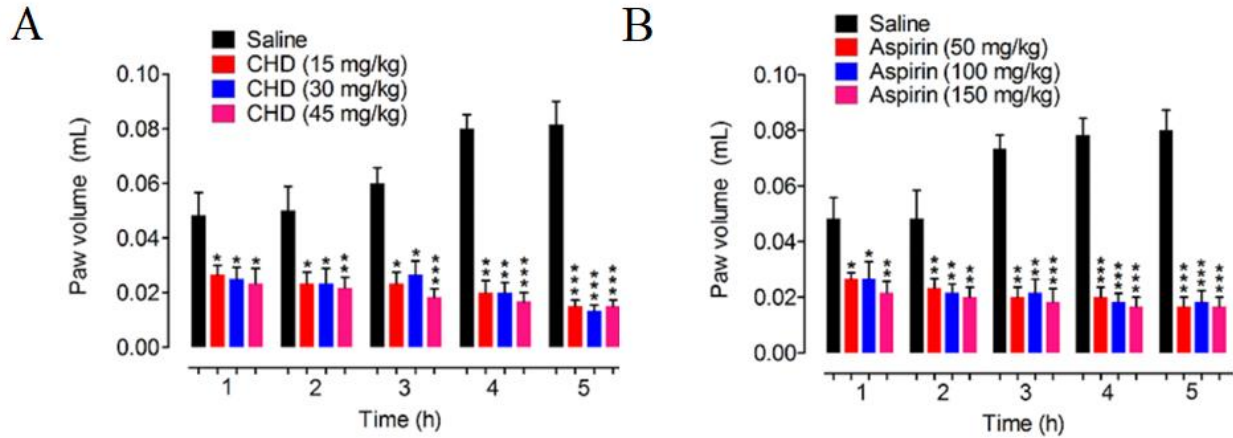
471 *3.2.7. Anti-inflammatory action of CHD against phlogistic agents (carrageenan, histamine, and*
 472 *serotonin) in the paw volume inflammation and xylene in the ear inflammation test*

473
 474 Intraplantar administration of the phlogistic agents, carrageenan, histamine, and serotonin was
 475 associated with a pronounced inflammatory response manifested by a substantial increase in the
 476 paw volume. The increased edema formation followed a temporal pattern and was first expressed
 477 during the initial h of the paradigm and maintained throughout the advanced stages of
 478 inflammation i.e. up to 5 h of the study duration. A dose dependent anti-inflammatory effect was
 479 produced by CHD in the three paradigms of paw edema. Treatment with CHD (15, 30 and 45
 480 mg/kg) reduced the inflammatory response evoked up to 5 h after administration of carrageenan
 481 (Fig. 8A), histamine (Fig. 9A), and serotonin (Fig. 10A), Treatment with the aspirin positive

482 control, (50-150 mg/kg) consistently displayed an anti-inflammatory effect up to 5 h after injection
 483 of carrageenan, serotonin or histamine in the inflammatory paradigms (Figs 8B, 9B and 10B).
 484 In the xylene provoked ear inflammatory edema paradigm, application of xylene produced a
 485 marked inflammatory response as observed by the increased ear weight recorded in the saline
 486 treated control animals (Fig 11). This marked oedematous change was significantly countered by
 487 treatment with CHD (30 and 45 mg/kg). The positive anti-inflammatory control drugs,
 488 indomethacin (10 mg/kg) and diclofenac (15 mg/kg) both produced a noteworthy decline in the
 489 augmented ear weight edema induced by xylene, as compared to the saline treated controls (Fig.
 490 11).

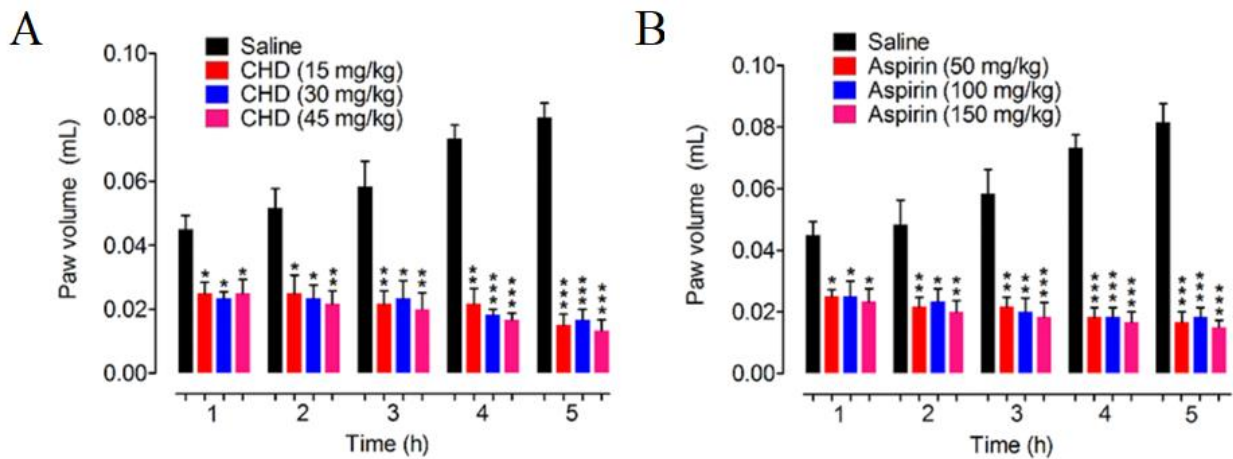


491
 492 **Fig. 8.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the
 493 carrageenan induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$,
 494 ** $P < 0.01$, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post*
 495 *hoc* Dunnett's test), ($n = 6$ mice per group).



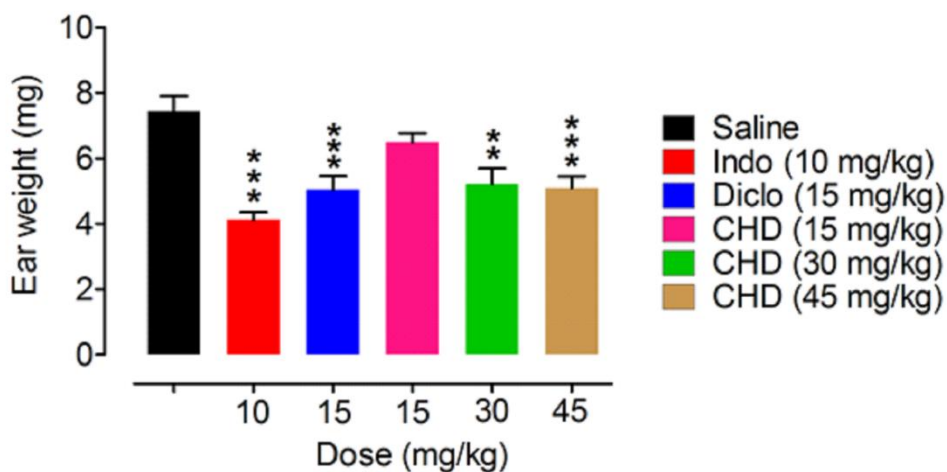
496

497 **Fig. 9.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the histamine
 498 induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$, ** $P < 0.01$,
 499 *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
 500 Dunnett's test), ($n = 6$ mice per group).



501

502 **Fig. 10.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the
 503 serotonin induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$, ** $P <$
 504 0.01, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
 505 Dunnett's test), ($n = 6$ mice per group).



506
 507 **Fig. 11.** Anti-inflammatory activity of CHD and the positive controls, indomethacin (Indo), and
 508 diclofenac (Diclo) in the xylene induced ear edema test. Each bar represents ear weight in mg \pm
 509 S.E.M. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ compared to the saline treated group (one-way ANOVA
 510 followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

511
 512
 513
 514 *3.3. In silico studies*

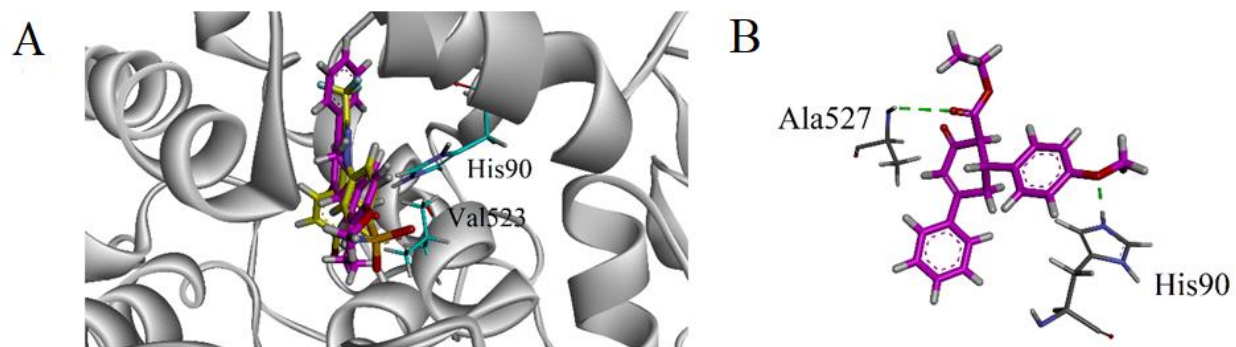
515 *3.3.1. Molecular Docking*

516 Docking studies were performed to explore any possible underlying mechanism(s) of CHD anti-
 517 nociception and anti-inflammatory activity. Accordingly, simulations were carried out on: (1)
 518 cyclooxygenase-2 enzyme (COX-2), (2) GABA receptors and (3) opioid μ -, δ - and κ - receptors
 519 using Molecular Operating Environment (MOE 2016.08, Chemical Computing Group, Canada).
 520 Data concerning three-dimensional (3D) structures of enzymes with their co-crystallized ligands
 521 were downloaded from the Protein Data Bank (PDB) listed in Table 1 and the docking algorithm
 522 was validated by re-docking native co-crystallized ligands (Table 1). The computed root mean

523 square deviation (RMSD) between experimental and re-docked poses was found to be within a
524 threshold limit $< 2 \text{ \AA}$.

525 The binding orientation of CHD and the native ligand into the binding site of the COX-2 isoform
526 is shown in Figure 12A. The three-dimensional (3-D) interaction plot of CHD showed that the
527 methoxy group formed a hydrogen bond interaction with His90, an important residue of a
528 selectivity pocket. The carbonyl oxygen formed hydrogen bond interactions with Ala527 (Figure
529 12B). The computed binding energy for the CHD-COX-2 complex was -8.1050 kcal/mol and the
530 docking score was -12.0458 .

531



532

533 **Fig. 12.** (A) Ribbon diagram of overlaid binding orientation of CHD and native ligand into the
534 binding site of the COX-2 enzyme. (B) Three-dimensional ligand-enzyme interaction plots of the
535 cyclohexenone derivative (CHD) into the binding site of COX-2 enzyme.

536

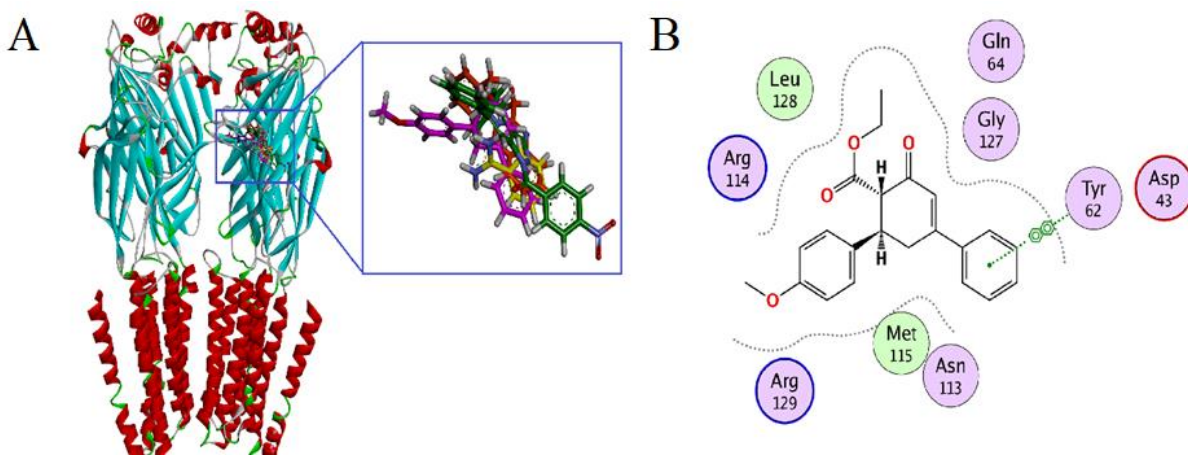
537 For the GABA receptor, the docking study was carried out on PDB code 4COF (benzamidine).

538 The computed binding energy for the ligand-GABA_A complex was obtained as -5.4853 kcal/mol

539 with a docking score of -8.4314 . The superimposed three-dimensional ribbon model of the CHD,

540 (purple), methaqualone (orange) (a positive allosteric GABA_A receptor modulator) and native

541 ligand benzamidine (yellow) is shown in (Fig. 13). The 2D interaction plot showed that the phenyl
542 ring of the synthesized compound creates π - π assembling interactions with Tyr62.



543
544 **Fig. 13.** (A) Three-dimensional superimposed binding pose of the native ligand benzamidine
545 (yellow), cyclohexenone derivative (CHD; purple) and methaqualone (orange) into the binding
546 site of the GABA_A receptor (PDB code 4COF) and (B) Two-dimensional interaction plot for CHD.

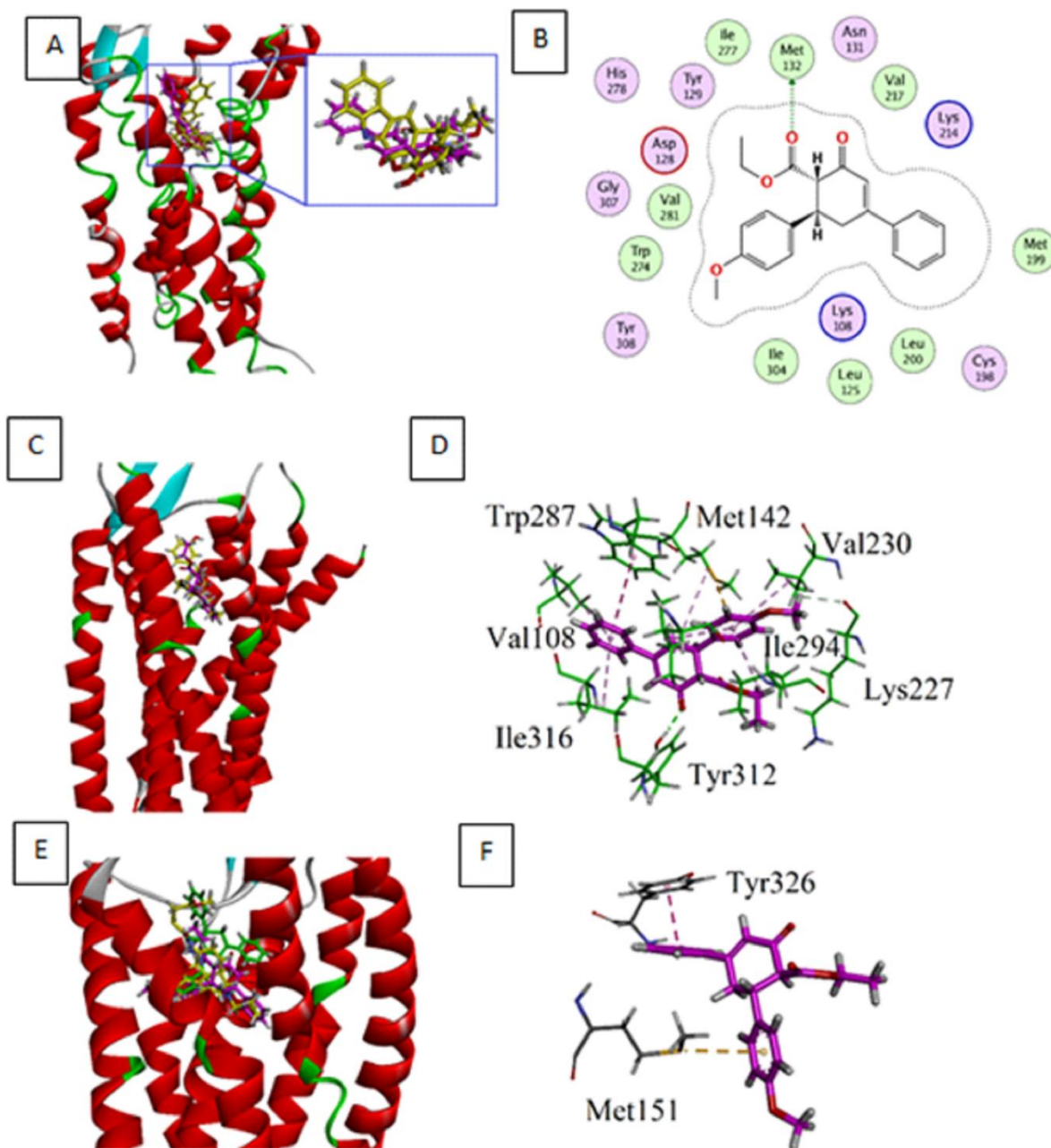
547
548 For μ -opioid receptors (μ OR), the computed binding affinity for the ligand-receptor complex was
549 -7.0501 kcal/mol and the docking score was computed as -11.4240 (Fig. 14). μ OR are important
550 opioid receptors for pain perception and are currently the target of various potent centrally-acting
551 analgesic drugs. The binding pose of CHD (purple) overlaid with β -funaltrexamine is shown in
552 (Fig. 14). The ligand enzyme complex was stabilized by hydrophobic and π -sulfur interactions.
553 The phenyl ring formed π - π stacking interactions with Tyr326, while the 4-methoxyphenyl group
554 formed π -sulfur interactions with Met151.

555 The binding affinity and docking score in the case of the κ -opioid ligand-receptor complex was
556 calculated as -8.0501 kcal/mol and -12.0240, respectively. The binding pose of the synthesized
557 compound (purple) into the κ -opioid receptor active site (PDB code 4DJH) is shown in Fig. 14.
558 CHD exhibited a binding pose similar to that of the co-crystallized ligand (JDC). The 3D interaction
559 plot showed that the ligand-enzyme complex was stabilized by a hydrogen bond, hydrophobic, π -

560 sulfur as well as π -CH type interactions. Met142 formed π -sulfur interactions with the 4-
561 methoxyphenyl ring. The phenyl ring of CHD engages in π - π stacking interactions with Trp287.
562 A hydrogen bonding interaction was found between the carbonyl oxygen of the ring with Tyr312.
563 Similarly, Val108, Val230, Val290, Ile294 and Ile316 formed some π -alkyl interactions.

564 For the δ -opioid receptor, the binding affinity for the ligand-enzyme complex was calculated as -
565 7.4000 kcal/mol and the docking score was noted as -11.0903. In the case of δ -opioid receptors,
566 the 3D structure with naltrindole as co-crystalized ligand was retrieved (PDB code = 4EJ4). The
567 superimposed 3D binding pose of CHD (purple) with naltrindole (yellow) is shown in (Fig. 14).
568 The two-dimensional interaction plot showed that it interacted with Met132 *via* hydrogen bond
569 donor interactions.

570



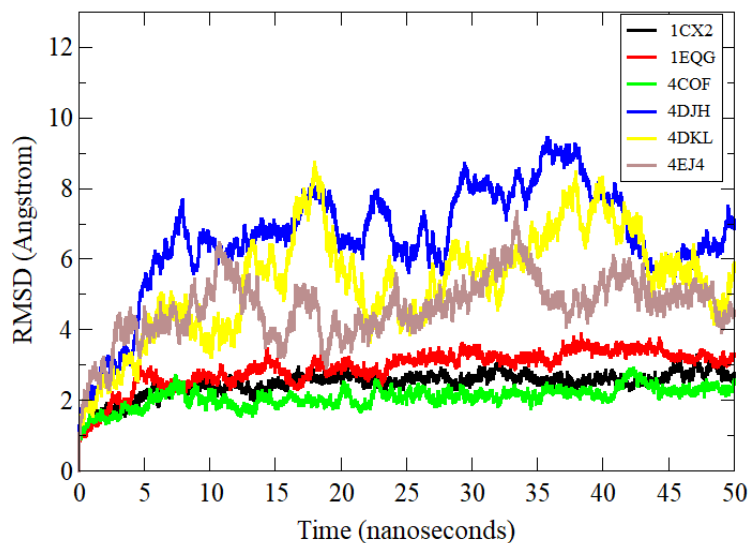
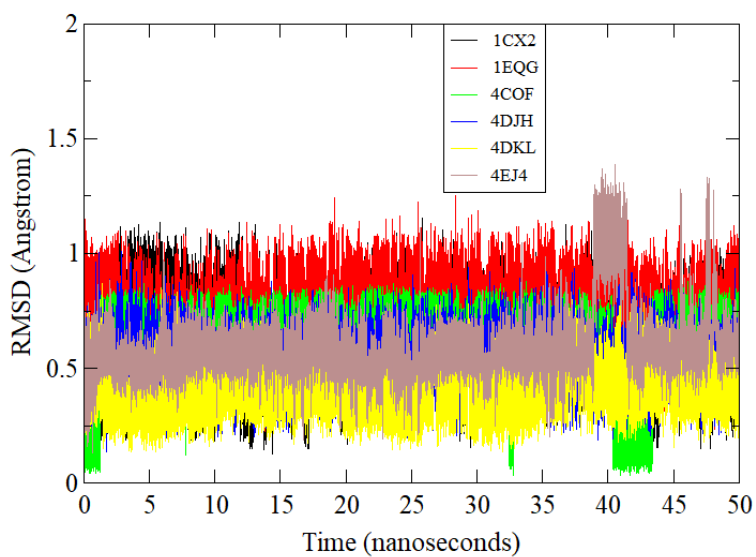
571

572 **Fig. 14.** Three and Two dimensional models of CHD binding with opioid receptors. (A) Three-
 573 dimensional and (B) Two-dimensional modeled superimposed binding pose of native ligand and
 574 CHD (purple) into the binding site of δ -opioid receptors (PDB code = 4EJ4). (C) Three-
 575 dimensional and (D) Two-dimensional model superimposed binding pose of native ligand and
 576 CHD (purple) into the binding site of κ -opioid receptor (PDB code = 4DJH). (E) Three-
 577 dimensional and (F) Two-dimensional model superimposed binding pose of the native ligand and
 578 selected compound CHD (purple) into the binding site of μ -opioid receptors (4DKL).

579

580 3.3.2. *Molecular Dynamics (MD) Simulations*

581 MD simulations were performed in order to understand the dynamics of all complexes and check
582 the stability of the CHD conformation at the docked site with respect to the backbone atoms for
583 each receptor. Among the complexes, 4COF, 1CX2, and 1EQG showed good stability in the
584 presence of CHD compared to the other three receptors (Fig. 15A). The mean RMSDs of these
585 were 2.0 Å, 2.4 Å, and 2.9 Å, respectively. These sites in the presence of CHD at the docked
586 position revealed very constant RMSD patterns throughout the simulation time, indicating a good
587 intermolecular strength of affinity and stability pattern. 4DJH (mean RMSD = 6.8 Å), 4DKL (mean
588 RMSD = 5.4 Å), and 4EJ4 (mean RMSD = 4.6 Å) showed major fluctuations in the receptor
589 structures, however, these changes do not affect the binding and conformation of the compound
590 with the receptors. In essence, these RMSD receptor fluctuations correspond to local protein
591 structure movements which are normal to their function. To substantiate compound conformation
592 stability, we additionally computed compound RMSDs in all complexes and plotted them versus
593 time. As can be seen in (Fig. 15B), the compounds were significantly stable with all receptor
594 RMSDs < 1 Å in all frames of the MD simulation.

A**B**

595

596 **Fig. 15.** (A) Root Mean Square Deviations of backbone atoms for each receptor of docked
597 complexes. (B) CHD Root Mean Square Deviations over 50-ns of MD simulation in complex with
598 receptors.

599

600

601

602 *3.3.1. MMPB/GBSA Binding Energy Calculation*

603 The free energy of binding was computed for all complexes to evaluate and revalidate the affinity
604 of intermolecular interactions and discover which type of interaction energy was dominant in
605 contributing to complex stability. All the complexes divulged robust interaction energies and were
606 dominated by gas phase energy in both MMGBSA and MMPBSA methodologies. Solvation
607 energy appeared to play less of a role in molecular interactions and was therefore non-favorable.
608 More specifically, the van der Waals energy of the gas phase disclosed by both methods played a
609 key role in complex stability whereas a minor contribution from electrostatic energy was also
610 evident except in the case of ICX2. The non-polar energy of solvation also favored docked
611 molecules as opposed to a highly unfavorable contribution from polar solvation energy. Overall,
612 the 4DKL receptor in complex with the CHD compound was highly stable with a MMGBSA
613 energy of -55.4492 kcal/mol and -47.9865 kcal/mol in MMPBSA. Details of MMGBSA and
614 MMPBSA energies of the complexes can be viewed in Table 4.

615

616 **Table 4**

617 MMGBSA and MMPBSA binding energies of the complexes

Method	Energy Component	1EQG	1CX2	4COF	4DKL	4EJ4	4DJH
MMGBSA	VDWAALS	-40.4511	52.1322	47.2564	62.2517	61.1760	-53.3142
	EEL	-6.2952	3.4110	25.3784	-8.2760	10.7373	-9.9464
	EGB	18.2375	8.9329	35.5743	21.3416	24.9894	19.8313
	ESURF	-4.3177	-5.6008	-4.8037	-6.2631	-6.2177	-5.5521
	DELTA gas G	46.7463	48.7212	72.6348	70.5277	71.9133	-63.2606
	DELTA solv G	13.9198	3.3320	30.7706	15.0785	18.7716	14.2792
	DELTA TOTAL	32.8265	45.3892	41.8641	55.4492	53.1417	-48.9814
MMPBSA	VDWAALS	-40.4511	52.1322	47.2564	62.2517	61.1760	-53.3142
	EEL	-6.2952	3.4110	25.3784	-8.2760	10.7373	-9.9464
	EPB	22.4009	14.6505	40.4885	26.2948	32.7712	26.3717
	ENPOLAR	-2.9491	-3.5859	-3.3982	-3.7537	-3.7336	-3.6908
	EDISPER	0.0000	0.0000	0.0000	0.0000	0.0000	0.0000
	DELTA gas G	46.7463	48.7212	72.6348	70.5277	71.9133	-63.2606
	DELTA solv G	19.4518	11.0646	37.0904	22.5411	29.0376	22.6810
	DELTA TOTAL	27.2945	37.6567	35.5444	47.9865	42.8757	-40.5796

618

619

620 **4. Discussion**

621 Cyclohexenone derivatives have received considerable attention over recent years not only
622 preclinically, but also clinically because of their extensive pharmacological possibilities. These
623 include: analgesic (Said et al., 2009), anti-inflammatory (Yaouba et al., 2018), anti-neuropathic
624 (Khan et al., 2019), antipyretic (Mousavi, 2016), antibacterial (Saranya and Ravi, 2012),
625 antioxidant (Okoth et al., 2016), antifungal (Kanagarajan et al., 2013), antimalarial (Ledoux et al.,

626 2017), anti-tubercular (Monga et al., 2014), anti-leishmanial (Das and Manna, 2015),
627 anticonvulsant (Said et al., 2009) tyrosine kinase inhibitory (Nazar et al., 2015) cytotoxic (Ayyad
628 et al., 1998) and anticancer (Okoth and Koorbanally, 2015) activities. Bearing in mind these wide-
629 ranging potential capacities of cyclohexenone functionality, this study was designed to examine a
630 selected cyclohexanone derivative (CHD) exemplar (Ethyl 6-(4-metohxyphenyl)-2-oxo-4-
631 phenylcyclohexe-3-enecarboxylate). This was done firstly for its safety profile; secondly, to
632 investigate any feasible *in vivo* effects in standard animal models of nociceptive and inflammatory
633 pain; thirdly, to perform molecular docking and molecular dynamic (MD) simulation studies to
634 facilitate interpretation of targeted drug-receptor interactions to corroborate the *in vivo* findings.
635 In parallel with this research approach, *in vitro* assays were conducted to examine any possibility
636 of COX-2 or 5-LOX enzyme inhibition and/or suppression of mRNA expression of TNF- α , IL-1 β
637 and COX-2 that might underlie anti-nociceptive and anti-inflammatory effects.

638 Four nociceptive and inflammatory, highly reproducible standard models were used to generate
639 the results. The findings clearly indicated that CHD possessed a noteworthy degree of safety with
640 a maximum tolerated dose above 240 mg/kg. Statistically significant anti-nociceptive and anti-
641 inflammatory activity was found in the rodent models. These effects were comparable to those of
642 aspirin, tramadol, indomethacin and diclofenac used as positive controls (Figs. 3, 4, 5, 6, 7, 8, 9,
643 10 and 11). Moreover, *in silico* docking analysis demonstrated that CHD manifested
644 favorable interactions with common pain targets i.e. COX-1/2 enzymes in addition to opioid and
645 GABA_A receptors (Figs. 12, 13 and 14.) substantiating the in-vivo results. Equally, CHD produced
646 marked inhibition of COX-2 and 5-LOX in the enzyme assays while in the case of RT-PCR, CHD
647 reduced the mRNA expression of TNF- α , IL-1 β and COX-2.

648 Administration of GABA receptor agonists either supraspinally, spinally or peripherally, has
649 been reported to reduce the nociceptive index in models of neuropathic and inflammatory pain
650 (Malan et al., 2002; Patel et al., 2001). In our study, it is postulated that CHD alleviates centrally
651 mediated nociception via GABAergic and opioidergic mechanisms (Fig. 7) alongside a capability
652 of interaction with the COX-1/2 target (Fig. 12). An involvement of GABAergic and opioidergic
653 systems was further reinforced by computational studies whereby CHD exhibited favorable
654 binding affinity for the GABA_A (Fig. 13) and opioid receptor subtypes (μ , κ and δ) (Fig. 14). It
655 has been reported that GABAergic agonists may augment the anti-nociceptive effect of a centrally
656 acting analgesic such as morphine (Sawynok, 1984), hence, it is conceivable that GABA receptor
657 agonist administration may represent a therapeutic option for the management of both chronic and
658 acute pain (McCarson and Enna, 2014) or as a combination of GABA with opioid receptor related
659 therapies.

660 The acetic acid induced abdominal constriction assay is a tonic visceral pain model frequently
661 utilized for monitoring the anti-nociceptive action of drugs (Utsunomiya et al., 1998). Although it
662 is a very sensitive test, it cannot distinguish whether the nociceptive activity is peripherally or
663 centrally mediated (Chen et al., 1995). It entails stimulation of visceral receptors followed by the
664 release of bradykinin, serotonin, cyclooxygenase, prostaglandins and interleukins which induce
665 pain and inflammation (Olonode et al., 2015; Rodrigues et al., 2012). It also implicates an
666 enhanced activation of peripheral receptors (Bentley et al., 1983) and innervated nociceptive nerve
667 terminals (Duarte et al., 1988). In the current study, CHD induced a significant reduction in
668 abdominal constrictions in a dose-dependent manner comparable to standard aspirin (Fig. 3A-B).

669 Hot plate and tail immersion nociceptive tests were employed to determine the central anti-
670 nociceptive potential of CHD. These models can specifically evaluate possible central nociception

671 (Eddy and Leimbach, 1953), where there is a non-inflammatory and acute nociceptive reaction
672 developed upon exposure to heat via spinal receptors which is evidence of centrally mediated anti-
673 nociception (Amabeoku and Kabatende, 2012; Pini et al., 1997). CHD moderately enhanced the
674 hot plate latencies of mice compared to standard tramadol, suggesting it to be a centrally acting
675 analgesic (Fig. 4). In the tail immersion test, the behavioural response is predominantly controlled
676 by supraspinal and spinal entities (Danneman et al., 1994). At the doses studied, CHD presented a
677 modest increase in tail withdrawal latency, but tramadol produced a more pronounced latency
678 elevation (Fig. 5). The duration of action of a drug depends on several factors including biological
679 half-life, first pass effect, plasma protein binding and other pharmacokinetic factors, nature of
680 formulation, co-morbid conditions such as renal impairment or liver dysfunction. Any of the above
681 cited factors, may be a potential contributor to the loss of CHD effectiveness at the doses of 30
682 and 45 mg/kg in the thermal nociception tests within 90 min (hot plate test) and 120 min (tail
683 immersion test), respectively. The formalin induced nociceptive paradigm comprises of a binary
684 phased nociceptive reaction and neuropathic pain (Salinas-Abarca et al., 2017). A neurogenic or
685 first phase (0-5 min) in which class C fibres are stimulated and an inflammatory or second phase
686 (10 to 30 min) which involves the release of inflammatory mediators (Hunskaar and Hole, 1987;
687 Tjølsen et al., 1992). Interestingly, CHD was effective in both the neurogenic and inflammatory
688 mediator phases (Fig. 6), further reinforcing the concept of a possible centrally acting anti-
689 nociceptive component mechanism in the activity of this compound. Moreover, in experiments
690 involving pharmacological antagonism of CHD anti-nociception with PTZ and naloxone, it was
691 divulged that an apparent participation of both GABAergic and opioidergic mechanisms was
692 implicated (Fig. 7A-B).

693 The anti-inflammatory activity of CHD was investigated by employing four standard models
694 of inflammation i.e., the carrageenan, serotonin, histamine and xylene mediated edema tests (Figs.
695 8, 9, 10 and 11). The carrageenan incited paw volume model is most extensively employed for
696 evaluating the anti-edematous potential of drugs (Mazzanti and Braghiroli, 1994). Localised paw
697 injection of carrageenan in mice initiates a three-phased inflammatory process. The primary phase
698 (0 to 1.5 h), is caused by the release of serotonin and histamine whereas the secondary phase (1.5
699 to 2.5 h) is mediated via bradykinin and the tertiary phase (2.5 to 5 h) is elicited mainly by the
700 generation of prostaglandins (Suba et al., 2005).

701 CHD (15 - 45 mg/kg) substantially reduced the elevated paw edema in all three phases of the
702 carrageenan-induced paw volume assay and this was comparable to the response yielded by the
703 standard drug, aspirin (Fig 8A-B). In order to authenticate the finding from the carrageenan paw
704 edema model, the anti-edematous effect of CHD was further investigated in the three other
705 standard models (histamine and serotonin induced paw volume and xylene induced ear edema).
706 Histamine and serotonin can increase vascular permeability and both are effective vasodilators
707 (Skidmore and Whitehouse, 1967) which are conducive to an ensuing edema. CHD not only
708 suppressed the edema mediated by histamine and serotonin but also that of xylene at doses
709 corresponding to standard anti-inflammatory drugs (Figs. 9A-B, 10A-B and 11). The xylene
710 induced ear edema model is extensively utilized to determine the anti-inflammatory action of
711 steroidal and non-steroidal anti-phlogistic agents (Zanini Jr et al., 1992). Studies reported in the
712 literature have revealed that xylene also promotes vascular permeability causing skin edema owing
713 to the release of inflammatory mediators leading to acute neurogenic inflammation (Bánki et al.,
714 2014). CHD markedly reduced ear edema induced by xylene comparable to the standard agents
715 (Fig. 11).

716 To further corroborate the anti-nociceptive and anti-inflammatory potential of CHD, it was
717 subjected to *in vitro* studies involving 5-LOX and COX-2 enzyme inhibition assays along with
718 RT-PCR studies. Thus, CHD substantially inhibited 5-LOX and COX-2 enzymes in comparison
719 with the standard inhibitors zileuton and celecoxib respectively as shown in (Table 2 and 3). In the
720 case of RT-PCR studies, CHD decreased the mRNA expression of COX-2, TNF- α and IL-1 β
721 compared to the carrageenan treated control group as presented in Fig. 2. This *in vitro* study
722 therefore endorsed the promising anti-nociceptive and anti-inflammatory findings with CHD in
723 both the *in vivo* and *in silico* studies which further strengthens a potential for application in pain
724 and inflammation.

725 In summary, *in silico* docking analysis demonstrated that the synthesized cyclohexanone
726 derivative has shown favourable interactions with common pain targets i.e. COX 1/2, GABA_A and
727 opioid receptors (Figs. 12, 13 and 14). The binding affinity study revealed that the intensity of
728 interactions of the CHD ligand with the COX-2 isozyme was more than that with COX-1 and this
729 was supported by the degree of 5-LOX and COX-2 enzyme inhibition observed (Table 2 and 3).
730 In addition, MD simulations of the complexes revealed that CHD was a highly stable molecule at
731 the docked site and generated robust chemical interactions underlying strong intermolecular
732 affinity (Table 4). Interactions with other pharmacological targets suggest that CHD may act as a
733 novel nociceptive and inflammatory pain reliever supported by *in vivo* studies (Figs. 3, 4, 5, 6, 7,
734 8, 9, 10 and 11).

735 **5. Conclusions**

736 This study elucidated the synthesis and pharmacological evaluation of a novel cyclohexenone
737 derivative (CHD) as a putative analgesic agent. CHD possessed not only anti-nociceptive, but also
738 anti-inflammatory activity when tested in validated models of pain and inflammation in mice.

739 These *in vivo* properties were attended by an inhibitory action on COX-2 and 5-LOX enzymes *in*
740 *vitro* in addition to a complementary *in silico* interaction with GABA_A and opioid receptors.
741 Consequently, CHD represents an innovative and noteworthy anti-nociceptive and anti-
742 inflammatory compound worthy of further pharmacological investigation and possible
743 development.

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749 Professor, Institute of Chemical Science, University of Peshawar. We are grateful to him for
750 providing a series of compounds and after preliminary study, we selected the cited compound
751 (CHD) for our study.

752 **Conflict of interest**

753 The authors have no conflict of interest.

754

755 **Authors' contributions**

756 GA conceived the research study and directed the research group as supervisor of the
757 pharmacological experimentation. GA also interpreted the results in addition to critically
758 reviewing the contents of the final version of manuscript. JK carried out the pharmacological
759 experiments and performed the statistical analyses. He likewise developed the preliminary draft of
760 the manuscript. RK helped in planning and supervising experiments related to the chemistry of our

761 selected compound. UR conducted the computational studies and performed related calculations,
762 interpretations and analysis. RU, MSJ, AAK, SA and SA helped in conducting the in vitro and in
763 silico studies. All authors read and approved the final manuscript and RDES had an intellectual
764 input in the writing of the manuscript and the interpretational outcome of the study.

765

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1000 **FIGURE LEGENDS:**

1001 **Fig. 1.** Chemical structure of Ethyl 6-(4-methoxyphenyl)-2-oxo-4-phenylcyclohex-3-
1002 enecarboxylate

1003 **Fig. 2.** Agarose gel electrophoresis (A) quantification of CHD activity on the mRNA level of
1004 COX-2 (B), TNF- α (C), and IL-1 β (D) in carrageenan induced hind paw edema in mice. The results
1005 are shown in relative arbitrary units (A.U). Bars represent mean expression in A.U \pm S.E.M. ###
1006 $P < 0.001$ compared to the saline group. ** $P < 0.01$, *** $P < 0.001$ compared to the vehicle group.

1007 **Fig. 3.** Anti-nociceptive activity of (A) CHD and (B) the positive control, aspirin in the acetic acid
1008 (1%) induced abdominal constriction test. Each bar represents mean percentage protection \pm
1009 S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as compared to the saline treated group (one-way
1010 ANOVA followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

1011 **Fig. 4.** Anti-nociceptive activity of CHD and the positive control, tramadol, in the hot-plate test.
1012 Each bar represents mean percentage protection \pm S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as
1013 compared to saline treated group (one-way ANOVA followed by *post hoc* Dunnett's test), ($n = 6$
1014 mice per group).

1015 **Fig. 5.** Anti-nociceptive activity of CHD and the positive control, tramadol in the thermal tail
1016 immersion test. Each bar represents mean withdrawal latency time in s \pm S.E.M). * $P < 0.05$, ** $P <$
1017 0.01, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
1018 Dunnett's test), ($n = 6$ mice per group).

1019 **Fig. 6.** Anti-nociceptive activity of CHD and the positive controls, indomethacin (Indo), and
1020 diclofenac (Diclo) in the formalin induced paw nociceptive test. Each bar represents mean
1021 nociceptive response in s \pm S.E.M). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ as compared to the saline
1022 treated group (one-way ANOVA followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

1023 **Fig. 7.** (A) Effect of naloxone at 1 mg/kg (NLX-1) and (B) PTZ at 15 mg/kg (PTZ-15) on the anti-
1024 nociceptive activity of CHD (30 mg/kg, CHD-30 and 45 mg/kg, CHD-45) or tramadol (30 mg/kg,
1025 TRD-30) in the mouse hot-plate test. Each bar represents mean percentage protection \pm S.E.M.
1026 *** $P < 0.001$ compared to saline control (SAL). (two sample *t*-test), ($n = 6$ mice per group).

1027 **Fig. 8.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the
1028 carrageenan induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$,
1029 ** $P < 0.01$, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post*
1030 *hoc* Dunnett's test), ($n = 6$ mice per group).

1031 **Fig. 9.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the histamine
1032 induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$, ** $P < 0.01$,
1033 *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
1034 Dunnett's test), ($n = 6$ mice per group).

1035 **Fig. 10.** Anti-inflammatory activity of (A) CHD and (B) the positive control, aspirin in the
1036 serotonin induced paw edema test. Each bar represents paw volume in ml \pm S.E.M. * $P < 0.05$, ** $P <$
1037 0.01, *** $P < 0.001$ as compared to saline treated group (one-way ANOVA followed by *post hoc*
1038 Dunnett's test), ($n = 6$ mice per group).

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1040 **Fig. 11.** Anti-inflammatory activity of CHD and the positive controls, indomethacin (Indo), and
1041 diclofenac (Diclo) in the xylene induced ear edema test. Each bar represents ear weight in mg \pm
1042 S.E.M. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ compared to the saline treated group (one-way ANOVA
1043 followed by *post hoc* Dunnett's test), ($n = 6$ mice per group).

1044 **Fig. 12.** (A) Ribbon diagram of overlaid binding orientation of CHD and native ligand into the
1045 binding site of the COX-2 enzyme. (B) Three-dimensional ligand-enzyme interaction plots of the
1046 cyclohexenone derivative (CHD) into the binding site of COX-2 enzyme

1047 **Fig. 13.** (A) Three-dimensional superimposed binding pose of the native ligand benzamidine
1048 (yellow), cyclohexenone derivative (CHD; purple) and methaqualone (orange) into the binding
1049 site of the GABA_A receptor (PDB code 4COF) and (B) Two-dimensional interaction plot for CHD.

1050 **Fig. 14.** Three and Two dimensional models of CHD binding with opioid receptors. (A) Three-
1051 dimensional and (B) Two-dimensional modeled superimposed binding pose of native ligand and
1052 CHD (purple) into the binding site of δ -opioid receptors (PDB code = 4EJ4). (C) Three-
1053 dimensional and (D) Two-dimensional model superimposed binding pose of native ligand and
1054 CHD (purple) into the binding site of κ -opioid receptor (PDB code = 4DJH). (E) Three-
1055 dimensional and (F) Two-dimensional model superimposed binding pose of the native ligand and
1056 selected compound CHD (purple) into the binding site of μ -opioid receptors (4DKL).

1057 **Fig. 15.** (A) Root Mean Square Deviations of backbone atoms for each receptor of docked
1058 complexes. (B) CHD Root Mean Square Deviations over 50-ns of MD simulation in complex with
1059 receptors.

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