

1 **Screening The Pathogen Box for Identification of New Chemicals Agents With Anti**
2 ***Fasciola hepatica* Activity**

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22

23 **Abstract**

24 Fascioliasis is an infectious parasitic disease distributed globally and caused by the liver
25 flukes *Fasciola hepatica* or *F. gigantica*. This neglected tropical disease affects both
26 animals and humans and it represents a latent public health problem due to the significant
27 economic losses related to animal husbandry. For decades, Triclabendazole has been the
28 unique anti-*Fasciola* drug that can effectively treat this disease. However, triclabendazole
29 resistance in Fascioliasis has been more recently reported around the world, and thus the
30 discovery of novel drugs is an urgent need. The aim of this study was to investigate the
31 fasciocidal properties of 400 compounds contained in the Pathogen Box. The first stage of
32 the screening was carried out by measuring the fasciocidal activity on metacercariae at a
33 concentration of 33 μM of each compound (standard dose). Subsequently the 50%
34 inhibitory concentration (IC_{50}) values of the most active compounds ($n=33$) were assayed
35 on metacercariae and resulted in 13 compounds with $\text{IC}_{50} \leq 10 \mu\text{M}$. The second stage
36 queried these compounds at 33 μM on adult flukes where seven showed high mortality
37 rates $> 50\%$. Four hit compounds were selected based on predicted nontoxic properties and
38 IC_{50} values obtained on adult worms resulted $< 10 \mu\text{M}$ thus representing the best
39 fasciocidal compounds tested here. Cytotoxicity assay on four types of cell lines
40 demonstrated that three compounds are nontoxic at its most active concentration. In
41 conclusion, three hit compounds identified in this proof-of-concept study are potential
42 candidates in the discovery of new fasciocidal drugs. Further studies are warranted.

43 **INTRODUCTION**

44 *Fasciola hepatica* is the etiological agent of fascioliasis, the most widespread trematodiasis
45 that affects both humans and herbivorous mammals such as sheep, cattle, goats and other
46 species (1, 2). In humans, fascioliasis can be acquired by the consumption of contaminated
47 vegetables. Up to 17 million people in 51 countries are estimated to be infected with *F.*
48 *hepatica* worldwide and more than 91 million are at risk of infection by this parasite (3, 4).
49 Among all continents, the Andean Region of South America is the most affected by
50 *Fasciola* where prevalence rates above 10% have been documented (5-8) and national
51 treatment programs are being scaled up.

52 Triclabendazole (TCBZ) is the most single effective fasciocidal drug, with activity against
53 both the infective larvae (*Metacercaria*) and adult worms, and an efficacy that exceeds 90%
54 in humans after a single oral dose (9, 10). Nonetheless, after decades of successful efficacy,
55 TCBZ resistance has developed in both animals and humans (11). Cases of TCBZ-resistant
56 *Fasciola* in both animals and humans have been reported in Australia, Europe and Latin
57 America (12-18). The development of TCBZ resistance represents an important public
58 health concern throughout the world that mainly affects animal husbandry and leads to
59 enormous economic losses (19). As a consequence, the discovery of novel effective drugs
60 and vaccines against *Fasciola* is an urgent need for the global control of fascioliasis.
61 Repurposing of praziquantel (PZQ) as anti-*Fasciola* drug failed whereas oxfendazole
62 showed to be an effective drug in animals (20, 21). Currently, there is no other fasciocidal
63 drug in clinical practice for humans, and thus TCBZ remains the unique treatment against
64 this infectious disease.

65 Open-access drug discovery provides a substantial resource in the research of those
66 diseases that affect primarily people living in low-resources locations. The Medicines for
67 Malaria Venture (MMV) foundation assembled a set of compounds, called “Malaria Box”,
68 which has been tested against various infectious agents including *Cryptosporidium parvum*
69 (22), *Plasmodium falciparum* (23, 24) *Schistosoma mansoni* (25, 26), *Toxoplasma gondii*
70 (27), and mycobacteria (28, 29). Later, a new set of chemical entities was assembled and
71 named the Pathogen Box collection. It contains 400 drug-like compounds that have showed
72 inhibitory activity on various infectious diseases such as haemonchosis, toxoplasmosis,
73 tuberculosis, neosporosis, malaria, sleeping sickness, Chagas, leishmaniasis and
74 trypanosomiasis (30-36). The Pathogen Box has been tested also in fungal diseases caused
75 by *Cryptococcus neoformans* and *Candida albicans* (37-39). The aim of this study was to
76 identify the fasciocidal activity of 400 compounds contained in the Pathogen Box by *in*
77 *vitro* testing.

78

79 MATERIALS AND METHODS

80 Study design. The study was conducted in three stages: (i) bioassays on metacercariae; (ii)
81 bioassays on adult worms; and (iii) cytotoxicity on cell (Figure 1). The best ffasciocidal
82 compounds were selected in each stage to be tested in the next phase based on *in vitro*
83 biological activity. To complement our knowledge on the active compounds obtained by
84 the experimental assays, computational resources were consulted to describe the chemical
85 properties as well as the *in silico* toxicology features and biological targets from these
86 active compounds.

87

88 Drugs and Media

89 The Pathogen Box was provided by the MMV agency (Geneva, Switzerland) and
90 manufactured by Evotec (USA). The 400 drug-like molecules were supplied in 96-well
91 plates as stock solutions of 10 mM dissolved in dimethyl sulfoxide (DMSO). Full data of
92 The Pathogen Box compounds is available at <https://www.pathogenbox.org> (40). TCBZ
93 was purchased from Sigma-Aldrich (Buchs, Switzerland). All of the compounds of The
94 Pathogen Box were dissolved in DMSO (Sigma-Aldrich, Irvine, UK) for drug stock
95 solutions of 200 μ M. Additional vials of MMV063404, MMV003270, MMV085210,
96 MMV676380, MMV687730, MMV687251, MMV1030799, MMV690102, MMV1029203,
97 MMV676053, MMV688179, MMV023969 and MMV688921 were manufactured by
98 Evotec (France). RPMI 1640 culture medium (Sigma- Aldrich, St. Louis, US) was used for
99 both stages, metacercariae and adult worm; supplemented with penicillin (100 U/ml) and
100 streptomycin (100 μ g/ml) (Sigma-Aldrich, St. Louis, US).

101 Parasites

102 Metacercariae of *F. hepatica* were obtained following the protocol described by Ortiz et al.
103 (16), at the Immunology and Research Laboratory of the Faculty of Veterinary Sciences of
104 the *Universidad Nacional de Cajamarca* in Peru. Eggs of *F. hepatica* were collected
105 directly from the gallbladder of sheep slaughtered in a popular abattoir in the city of
106 Cajamarca (TCBZ-resistant endemic area for fascioliasis). Miracidia were from *Fasciola*
107 eggs after incubation for 15 days at 25°C. Afterwards, they were used to infect *Lymnaea sp.*
108 snails (5-6 mm) in a proportion of two miracidia per snail. The infected snails were kept in

109 plastic containers for 45 to 60 days at room temperature. After this time, the snails were
110 stimulated by direct solar exposure and water at 4-8°C to produce metacercariae.
111 Approximately 20,000 metacercariae were obtained for this study and stored in cryovials
112 on distilled water at 4-8°C. Adult worms were collected from bile ducts of infected cattle
113 from a slaughterhouse in Lima, Peru; and maintained at 37°C until usage (within 2h).
114 Before its incubation, three washes with phosphate-buffered saline (PBS) (HiMedia, India)
115 and one additional with supplemented RPMI were performed to remove host debris. All the
116 incubations, for both metacercariae and adults, were carried out at 37°C with 5% CO₂.

117 *In vitro* screening on metacercariae

118 The 400 compounds were initially tested at 33 µM on *F. hepatica* metacercariae. Drug
119 stock solutions were diluted in 96-well plates (BD Falcon, US) with RPMI 1640
120 supplemented with antibiotic up to a final volume of 180 µL. In all *in vitro* assays, positive
121 and negative controls were run in parallel for each assay batch. A range between seven and
122 ten metacercariae were added to each well, previously analyzed microscopically to confirm
123 its viability (microscopic features intact). The metacercariae viability considered some
124 physical properties of the parasite determined by microscopy as described previously (41,
125 42). The MCs viability was surveyed as a function of both damage in membrane and fluke
126 colour (translucence). Therefore, a low viability corresponded to big damage and high
127 translucency. The viability scale was scored as follows: +++, total damage (dead parasite,
128 shattered membrane and mostly translucent); ++, partial damage (partial membrane damage
129 and highly translucent); +, mild damage (partial membrane damage, poorly translucent) and
130 no damage (intact membrane, dark metacercariae, lack of translucency).

131 Positive-control wells contained TCBZ 10 μ M whereas *F. hepatica* metacercariae
132 incubated in the presence of the highest concentration of DMSO served as negative control.
133 Each test was performed in triplicate. Culture plates were incubated at 37°C in a humidified
134 5% CO₂ atmosphere for 72 h. First, metacercariae were evaluated by inverted microscopy
135 (PhotoZoom, Cambridge Instruments) at magnification 10X and 20X at 24, 48 and 72 h
136 post drug exposure to determine its viability. Only the compounds that caused, on average,
137 at least 25% of metacercariae mortality at 72 hours were considered for IC₅₀ (50%
138 inhibitory concentration) determination. Experiments were run in sets of triplicates. The
139 mean mortality percent of the study compounds were compared to that of DMSO. A
140 standard deviation (SD) was also estimated.

141 In the second part, we determined the IC₅₀ of the selected compounds chosen in the
142 previous bioassay. Drugs were tested at concentrations of 2.1, 4.2, 8.4 and 33 μ M using
143 supplemented culture medium. The incubation was done under the conditions described
144 above, by triplicate and considering TCBZ and DMSO as controls. Anti-parasite activity
145 was evaluated at 24, 48 and 72 h post exposure, using the above-mentioned metacercariae
146 viability scale. Viability (mean % of viable parasites) at 72h was considered for the
147 estimation of IC₅₀. IC₅₀ values of test compounds were determined by linear regression
148 analysis using CompuSyn software (Version 3.0.1, 2007; ComboSyn Inc., USA). The linear
149 correlation coefficient (r) was obtained.

150 Assessment of anti-*Fasciola* activity *in vitro* on adult worms

151 Those compounds that showed activity IC₅₀ \leq 10 μ M on metacercaria were subsequently
152 tested in adult stage of *F. hepatica*. In all *in vitro* assays, positive and negative controls
153 were run in parallel for each assay batch. First, the selected compounds were tested at 33

154 μM by triplicate, using drug stock solutions diluted in supplemented RPMI on 6-well plate
155 up to a final volume of 4 ml. Adult worms were thoroughly washed with PBS to remove
156 host debris and then three worms were placed in each well. The incubation was done under
157 the same conditions as those applied in bioassays with metacercariae. Positive control
158 consisted of 50 μM TCBZ and the negative control was DMSO at the highest
159 concentration. The viability of adult flukes was scored after 24 and 48 h using a motility
160 criterion previously described (43) and also color and rigidity criteria previously applied by
161 our team (data not published). Motility was assessed only in adults and not in MC because
162 this latter has no movements. Rigidity was a parameter used to confirm the damage caused
163 by the drug once the incubation time finished. In general, low motility level corresponded
164 to transparent and rigid worms. Those changes were attributed to the damage caused by a
165 drug. The viability scale was determined as follows: (i) worm motility: 3, normal
166 movements; 2, reduced movements; 1, very weak movements and 0, absence of movements
167 (i.e. death of worm); (ii) worm color: +++, dark red; ++, pink; +, slightly transparent and -,
168 totally transparent; and (iii) worm rigidity: -, no rigidity; +, rigidity and ++, cell break when
169 touched. Assessments at 72h post drug exposure were not done because death of worms
170 always occurred \leq 48h. Experiments were run in sets of triplicates. The mean mortality
171 percent and SD of the study compounds were estimated. The selected compounds were
172 those that caused an average mortality $>$ 50% in adult parasites. Then IC_{50} assays were
173 conducted by testing the selected compounds at five different concentrations 0.31, 0.93,
174 2.78, 8.33 and 25.0 $\mu\text{g}/\text{ml}$. DMSO and TBZ were used as negative and positive controls,
175 respectively. Parasite viability at 24 h were estimated based on survival in DMSO. The
176 IC_{50} s and 95% CI were estimated using GraphPad Prism 7.0 software using a variable
177 slope of the sigmoidal curve from normalized percent activity values and \log_{10} -

178 transformed concentrations. Top and bottom values were constrained to 100 and 0,
179 respectively. The fasciocidal activity was determined by considering the adult viability
180 scale described before.

181 Computational analysis.

182 Evaluation of biological targets of small compounds. To learn about biological targets,
183 those compounds that showed promising anti-fasciola activity in the adult stage as well as
184 TCBZ were entered in the ChEMBL database (<https://www.ebi.ac.uk/chembl/>) (44). First,
185 the SMILES (Simplified Molecular-Input Line-Entry System) of each selected compound
186 were obtained from the supplementary material provided by the MMV (also available at
187 www.mmv.org). Then the SMILES were entered in ChEMBL and known targets of each
188 compound were retrieved. ChEMBL compares the query compound to a large database of
189 compounds and their targets available from multiple sources including the projects funded
190 by MMV (45). The target name, organism and protein target classification were collected.

191 *In silico* cytotoxicity prediction. Lazar (lazy structure–activity relationships), a modular
192 framework for predictive toxicology, was consulted to predict the toxic effects of the
193 selected compounds that showed activity on metacercariae (46-48). Lazar was accessed
194 through <https://lazar.in-silico.de/predict> and SMILES of each compound were entered.
195 Relevant data including carcinogenicity in rodents, mutagenicity in *Salmonella typhi* and
196 acute toxicity on *Fathead minnow*, Blood Brain Barrier Penetration and the maximum
197 recommended daily dose in humans were predicted.

198 Cell Growth Inhibition Bioassay.

199 Cytotoxicity of the compounds was evaluated in tumor and non-tumor cell lines using the
200 sulforhodamine B (SRB) assay method (49, 50). Cell lines tested include BALB/3T3 (Non-
201 tumorigenic, BALB/c mouse embryo cells), H460 (human lung large cell carcinoma),
202 DU145 (human prostate carcinoma) and HT-29 (human colon adenocarcinoma).

203 To determine the cytotoxicity of the compounds, cells were plated into 96-well tissue
204 culture plates and in their corresponding growth medium Dulbecco's Modified Eagle
205 Medium (DMEM) at approximately 10% confluency (BALB/3T3 at 3,500 cells/well, H460
206 at 1,500 cells/well, DU145 at 3,500 cells/well and HT-29 at 3,000 cells/well) and incubated
207 at 37°C in a 5% CO₂ and 95% air humidified atmosphere for 24 h to allow cells to attach. A
208 plate containing each of these cells was fixed *in situ* with trichloroacetic acid (TCA) in
209 order to obtain the cell values at zero time before adding the compounds. The rest of the
210 plates containing the different cell lines received serial dilutions of the compound to be
211 tested at the following final concentrations: 4, 1, 0.25 and 0.0625 µg/mL. The plates were
212 then incubated at 37°C in a 5% CO₂ and 95 % air humidified atmosphere for 48 h. The
213 assay was terminated by the addition of cold TCA. TCA treated plates were incubated at
214 4°C for 1 hour and then washed five times with tap water to remove TCA and air dried.
215 Background optical densities were measured in wells incubated with growth medium
216 without cells. TCA-fixed cells were stained for 20 minutes with 0.4% (w/v) SRB dissolved
217 in 1% acetic acid. At the end of the staining period unbound dye was removed by washing
218 four times with 1% acetic acid. After air drying the plates, bound dye was solubilized with
219 10 mM Tris base (pH 10.5) and the absorbance read on an automated plate reader at a
220 wavelength of 550 nm. The GI₅₀ value was defined as the concentration of test sample
221 resulting in a 50% reduction of absorbance as compared with untreated controls that

222 received a serial dilution of the solvent in which the test samples were dissolved and was
223 determined by linear regression analysis. The optical density values obtained were used to
224 determine the cell growth and cytotoxicity from each compound.

225 Ethics. This study was approved by the Animal Ethics Committee of the Universidad
226 Peruana Cayetano Heredia (Approval ID Code 41-07-16).

227 **RESULTS**

228 *In vitro* activity of The Pathogen Box determined on *F. hepatica* metacercariae

229 In the first stage of the study, the 400 compounds contained in the Pathogen Box were *in*
230 *vitro* screened against *F. hepatica* metacercaria. A total of 33 compounds showed mean
231 mortality rates above 25% at 33 μM but all these resulted being less active than TCBZ
232 (mortality rate of 90%) as shown on Table 1. Fasciocidal activity of these 33 compounds
233 was then assessed by determining the IC_{50} values (Table 1). As a result, 13 compounds
234 showed potent inhibitory activities with IC_{50} values between 0.31 μM and 8.23 μM and
235 were then assayed in adult worms although its low r values (Table 1).

236 *In vitro* activity of selected compounds on *F. hepatica* adult worms and *in silico* toxicology
237 prediction.

238 The 13 selected compounds listed on Table 2 were assayed at 33 μM in adult worms. Seven
239 compounds produced moderate or high mean mortality rates (> 50%) (Table 2). These
240 were MMV003270, MMV676380, MMV690102, MMV1029203, MMV063404,
241 MMV1030799, and MMV688921. Six compounds showed low mortality rates (<50%) and
242 for that reason these were not considered in the next assays. Before to proceed with the IC_{50}
243 assay, *in silico* safety profiles of the seven selected compounds were predicted by Lazar

244 program (Table 1). Whereas MMV003270 and MMV676380 were predicted non-
245 carcinogenic and non-tumorigenic compounds, MMV690102 was deemed non-
246 carcinogenic and tumorigenic (Table 1). MMV1029203, MMV063404, MMV1030799, and
247 MMV688921 were predicted carcinogenic and tumorigenic substances. Thus, the three
248 deemed non-carcinogenic compounds as well as MMV1029203, a predicted carcinogenic
249 substance that had the highest mean mortality rate (78%), were tested in adult worms. Such
250 four compounds constitute our hit compounds.

251 To determine which of the four hit compounds were most potent at inhibiting the growth of
252 *F. hepatica* adult worms, the IC₅₀ values were determined. The hit compounds had IC₅₀
253 values < 10 µM in adult worms (Table 3, Fig. S1, Table S1). These four hit compounds
254 were tested in the cytotoxicity study on cell cultures.

255 *In vitro* cytotoxicity on cell lines.

256 Cytotoxicity of the four hit compounds against cell lines was evaluated in culture (Table 3).
257 The GI₅₀ values ranged from 0.95 and >23.73 µM across the four types of cell lines assayed
258 (Table 3). MMV003270, MMV676380, MMV1029203 and TCBZ presented GI₅₀ values
259 above its IC₅₀ values thus meaning that these compounds are not toxic at their active
260 concentrations. In one of the four cell lines, MMV690102 had a GI₅₀ value below its IC₅₀
261 value thus suggesting that it may cause a level of toxicity in certain cell types at its active
262 concentration (Table 3).

263 Computational recognition of targets.

264 As a result of the search in the ChEMBL database, a total of 27 targets were recognized for
265 TCBZ whereas MMV003270 resulted to have 19 known target, most of them in humans

266 (Table 4). MMV003270 and TCBZ have common human targets that comprise Nuclear
267 factor erythroid 2-related factor 2, Microtubule-associated protein tau and TAR DNA-
268 binding protein 43. According to the data deposited in ChEMBL, MMV003270 targets a
269 number of cytochrome p450 of family 1, 2 and 3. MMV676380 and MMV023969 have
270 identical cell targets that include human glucose transporter and hexose transporter of
271 *Plasmodium falciparum* and *Leishmania mexicana* (Table 4). Targets for MMV1029203
272 and MMV676053 also resulted to have known targets including human ferrochelatase and
273 Inosine-5'-monophosphate dehydrogenase of *Cryptosporidium parvum*, respectively. The
274 remaining eight compounds had no known targets according to the ChEMBL database
275 (Table 4).

276

277 DISCUSSION

278 In the present study, the Pathogen Box was queried to identify compounds with *in vitro*
279 anti-Fasciola activity against both metacercariae and adult worms (Figure 1). We found 13
280 compounds with potent inhibitory activity on metacercariae ($IC_{50} < 10 \mu M$), meaning that
281 3% of the substances within the Pathogen Box are effective against the infective form of *F.*
282 *hepatica*. Two out of the 13 compounds (MMV687730 and MMV687251) had the most
283 potent activity against metacercariae with IC_{50} values below $1 \mu M$ but showed mild effects
284 on adult worms (Tables 1 and 2). Since we were interested in identifying hit compounds
285 that were active on larvae and adult stages, these two compounds were not further studied
286 (Table 2). When assayed on adult worms, seven promising compounds showed mortality
287 rates above 50% (Table 2). As a criterion for hit prioritization during the screening on adult
288 worms, we prepared a list of hit compounds that mostly excluded the predicted

289 carcinogenic/tumorigenic compounds (Table 3). Thus, three (MMV676380, MMV003270
290 and MMV690102) of the seven most promising candidates were included in the list of hit
291 compounds since they were predicted noncarcinogenic agents (Table 1). One additional
292 compound (MMV1029203) that was predicted as a carcinogenic compound was also
293 included due to its very high effect on adult worms. According to our results, the four hit
294 compounds resulted potent inhibitory molecules both on MC and adult stages (Table 3).
295 The cytotoxicity assay revealed that three hit compounds (MMV676380, MMV003270 and
296 MMV1029203) were non-toxic agents at its most active concentrations when assayed on
297 cell lines (Table 3). In contrast, MMV690102 may cause cell cytotoxicity at its most active
298 concentration meaning that it is not a primary candidate for drug development (Table 3).
299 Our results are consistent with previous cytotoxicity assays on HepG2, HL60 and MRC5 as
300 shown on Table 3 (data provided by the MMV as part of the supporting information for the
301 Open Access Malaria Box).

302 Repurposing of hits, using the Pathogen Box, against *F. hepatica* is highly relevant since
303 TCBZ is the only existing effective drug for which resistance is known (51-53). Previous
304 works tried to repurpose albendazole, nitroxylnil and closantel as candidate fasciocidal
305 drugs but treatment failed (54, 55). In the present study, four out of 400 compounds
306 contained in The Pathogen Box showed potent inhibitory activity against the infective form
307 of *F. hepatica* as well as its adult form (Table 3). Such finding represents a relevant
308 contribution in the identification of dual drug candidates that are able to act against the
309 initial stages of the infective larvae (metacercaria) and adult forms of liver flukes, similar to
310 TCBZ. Additionally, other 13- compounds showed biological activity at $< 20 \mu\text{M}$ against
311 metacercaria (Table 1). Since MC represents the initial infective form of parasites, it should

312 be primarily controlled through potent compounds such as those identified here (Table 1).
313 Future exploration of The Pathogen Box in newly juvenile metacercaria is desirable given
314 that some compounds may have not penetrated the cyst wall of larvae. By testing
315 compounds on juvenile worms, some additional molecules might be recognized that are
316 active in adult worms.

317

318 The four hit compounds identified in this study have been previously characterized against
319 *Plasmodium falciparum*, *Ancylostoma ceylanicum*, *Trypanosoma cruzi* and *Leishmania*
320 *donovani* (data provided by the MMV as part of the supporting information for the Open
321 Access Malaria Box). Therefore, a common mechanism of action or target is plausible
322 among the hit compounds across such pathogens. For instance, MMV676380 has
323 previously shown to have a lethal effect on *P. falciparum* and here was found to be a potent
324 inhibitory compound against *F. hepatica* (36, 56). Known targets of MMV676380 are the
325 glucose and hexoses transporters suggesting that such mechanism may be affected in both
326 parasites in presence of such compound (Table 4). In the other hand, MMV003270
327 (Zoxazolamine), that is also active against *A. ceylanicum*, resulted to have 19 targets
328 including three human proteins that are also targeted by TCBZ (Table 4). Two of these
329 proteins are transcription regulators (Nuclear factor erythroid 2-related factor 2 and TAR
330 DNA-binding protein 43) whose disruption may affect the gene expression. Such finding is
331 in accordance with a hypothetical mechanism of action of TCBZ that involves a direct
332 effect of the drug on protein synthesis (11, 57). Similarly, the microtubule-associated
333 protein tau is a known target both of TCBZ and MMV003270. TCBZ is a benzimidazole-
334 derivative that disrupts the assembly of microtubules in helminths by binding to tubulin

335 molecules (58). Our results suggest that MMV003270 also affects the microtubules
336 formation mechanism. Common targets of TCBZ and MMV003270 may be partially
337 explained by the similar scaffold structures. MMV1029203, one of the four hit compounds,
338 targets a human ferrochelatase that is a mitochondrial factor involved in protoheme
339 biosynthesis. This latter is a vital process that exists also in *F. hepatica* and whose
340 disruption may be lethal. Some known targets of the hit compounds here identified
341 correspond to human proteins which suggests that a level of toxicity may exist in humans.
342 However according to our results with cell lines, the compound concentrations needed to
343 kill *F. hepatica* (IC₅₀) are considerably less than that to cause cell death (GI₅₀) which means
344 that these are nontoxic (Table 3), except for MMV690102. Although no *F. hepatica* target
345 is recognized for our hit compounds, the demonstration of inhibitory activity of such
346 chemical agents both in metacercariae and adult forms suggests that common targets may
347 exist in both liver fluke stages. The identification of drug targets becomes an important step
348 that drive the discovery of novel antiparasitic agents administered by various ways (34).
349 For that reason, further studies to identify potential *F. hepatica* targets of hit compounds are
350 desirable. Such a study should consider the recognition of human homologs in *F. hepatica*
351 according to our results (Table 4).

352 Our study has some limitations. First, TCBZ metabolites (TCBZ-sulfoxide and TCBZ-
353 sulfone) that are quickly released *in vivo* were not included in this pilot study. However
354 given that TCBZ has a moderate *in vivo* and *in vitro* fasciocidal effects it is suitable as
355 positive control in bioassays (59, 60). A second limitation is that alive *F. hepatica* worms
356 were collected from a local abattoir where some animals may have been infected by various
357 other pathogens or may have been treated with TCBZ. To guarantee the best quality of

358 adult worms for bioassays, we performed a quality control on adult fasciolas before using
359 these in the experiments. Thus, only worms that presented intense brown or red color and
360 that have active motility were selected. All the remaining were discarded. A third limitation
361 is the low number of parasites used for the assays, that did not allow performing formal
362 statistical comparisons of activity between TCBZ and test drugs. Obtaining MC and adults
363 was a challenging task since both MC and adult worms were collected from natural
364 reservoirs. Therefore we had limited access to parasites for bioassays. However, our
365 exploratory study aimed to identify fasciocidal compounds, we found that negative controls
366 were enough for such purposes.

367 In conclusion, we identified three promising non-cytotoxic drug-like compounds,
368 MMV003270, MMV676380 and MMV1029203, that showed a potent biological activity
369 against *F. hepatica* metacercaria and adult worms. Such compounds represent new lead
370 candidates to potentially become future anti-*F. hepatica* drugs. By acting both on infective
371 form and adult worms, such agents may provide an appropriate treatment against
372 fascioliasis.

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375

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381

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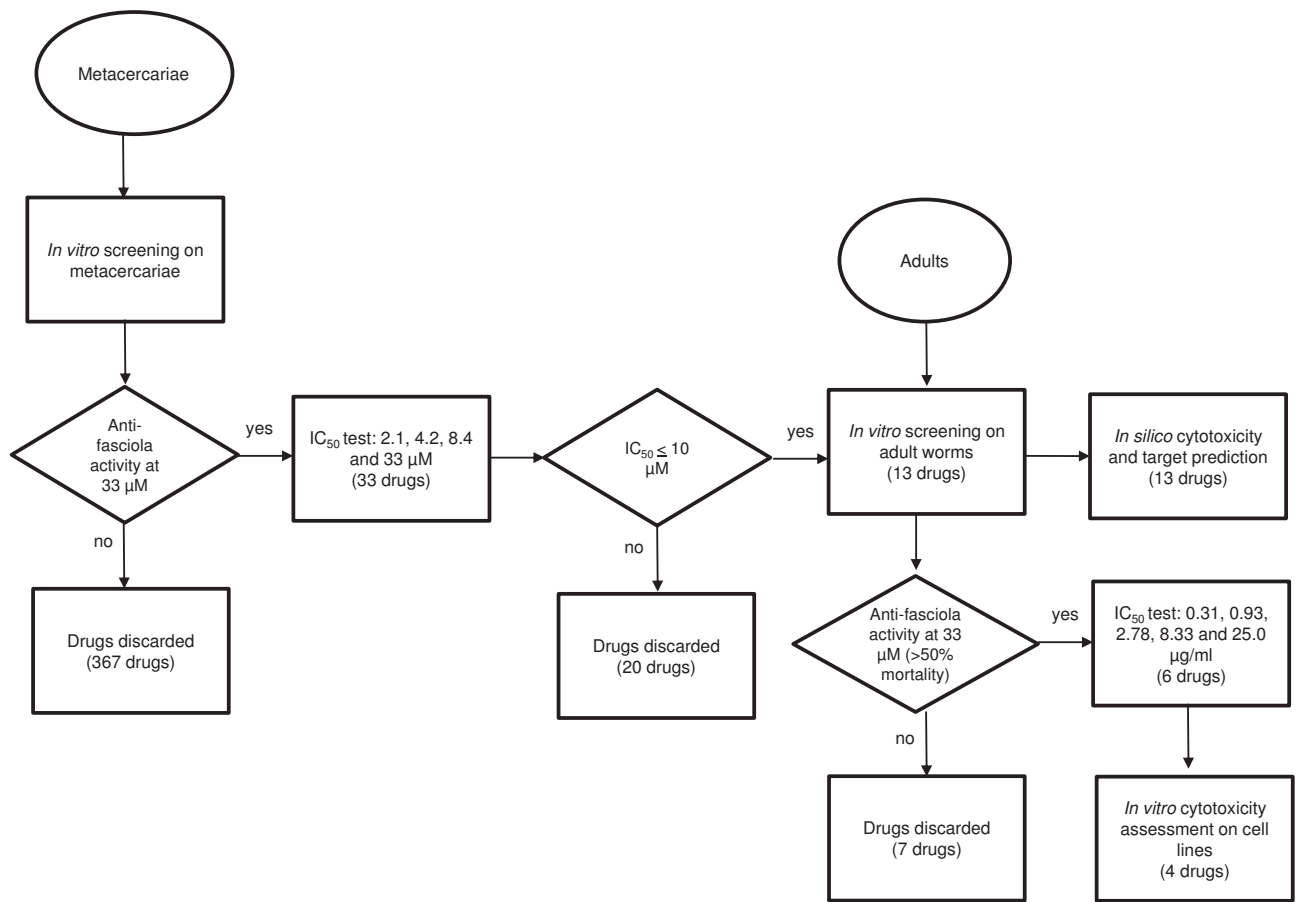


Figure 1. Flowchart of the tests carried out during the study

Table 1. Summary of the chemical compounds that showed the best biological activity against metacercariae.

| Compound plate code ¹ | MMV ID ² | Molecular Formula | Molecular weight (g/mol) | Mean % mortality on MC ³ | SD(%) | IC50 μ M ⁴ | R | Other infectious microorganisms ⁵ | In silico toxicity features ⁶ | | | | |
|----------------------------------|---------------------|-------------------|--------------------------|-------------------------------------|-------|---------------------------|------|--|--|---|--------------------------|--|--|
| | | | | | | | | | Acute cytotoxicity (Fathead minnow) | Blood Brain Barrier Penetration (Human) | Carcinogenicity (Rodent) | Mutagenicity (<i>Salmonella typhimurium</i>) | Maximum Recommended Daily Dose (Human) |
| TCBZ | N.A. | C14H9Cl3N2O5 | 359.7 | 100 † | 0 † | 15* | N.A. | Schistosoma | 4.57 (mg/L) | penetrating | non-carcinogenic | non-mutagenic | N.A. |
| PA2 | MMV010764 | C14H16N4O5 | 320.4 | 22 | 38.5 | 24.1 | -0.3 | Plasmodium | N.A. | N.A. | non-carcinogenic | non-mutagenic | N.A. |
| PA4 | MMV676388 | C15H14N4O3S | 330.4 | 29 | 24.7 | 16.9 | 0.8 | Mycobacterium | 254.0 (mg/L) | penetrating | carcinogenic | mutagenic | 2.44 (mg/kg_bw/day) |
| PA5 | MMV202553 | C15H15N3O2 | 269.3 | 29 | 24.7 | 14.9 | 0.9 | Kinetoplastids | 7.58 (mg/L) | penetrating | non-carcinogenic | mutagenic | 0.993 (mg/kg_bw/day) |
| PAG6 | MMV63404 | C19H24N3OCl | 345.9 | 54 | 7.2 | 5.3 | 1.0 | Mycobacterium | N.A. | penetrating | carcinogenic | mutagenic | N.A. |
| PAH6 | MMV676539 | C20H16N2O3 | 332.4 | 17 | 28.9 | 24.7 | -1.0 | Mycobacterium | 25.9 (mg/L) | penetrating | carcinogenic | mutagenic | 4.05 (mg/kg_bw/day) |
| PBD3 | MMV637953 | C51H40N6O23S6 | 1435.3 | 25 | 9.9 | 21.8 | -0.6 | Trypanosoma and Onchocerca | N.A. | penetrating | non-carcinogenic | non-mutagenic | N.A. |
| PBD7 | MMV019838 | C18H18N4O6F6 | 412.3 | 26 | 11.6 | 12.4 | 0.0 | Plasmodium | N.A. | penetrating | non-carcinogenic | mutagenic | N.A. |
| PBF4 | MMV003270 | C7H5N3OCl | 168.6 | 26 | 25.1 | 8.2 | -0.7 | Ancylostoma | 6.75 (mg/L) | penetrating | non-carcinogenic | non-mutagenic | 15.5 (mg/kg_bw/day) |
| PBF6 | MMV688853 | C19H23N5O2 | 389.9 | 25 | 22.5 | 31.9 | -0.8 | Cryptosporidium | N.A. | non-penetrating | non-carcinogenic | mutagenic | N.A. |
| PBF11 | MMV085210 | C22H24N3O3Cl | 446.0 | 40 | 15.3 | 2.4 | 0.8 | Plasmodium | N.A. | penetrating | non-carcinogenic | non-mutagenic | 1.64 (mg/kg_bw/day) |
| PBH10 | MMV676380 | C18H15N4O3Cl | 370.8 | 33 | 33.3 | 1.3 | 0.1 | Plasmodium | 132.0 (mg/L) | penetrating | non-carcinogenic | non-mutagenic | 101.0 (mg/kg_bw/day) |
| PCA2 | MMV675997 | C24H29N4O2F | 424.5 | 22 | 38.4 | 18.1 | -0.2 | Kinetoplastids | N.A. | penetrating | non-carcinogenic | mutagenic | 1.51 (mg/kg_bw/day) |
| PCA6 | MMV688852 | C16H17N5ClF | 333.8 | 29 | 37.4 | 17.2 | -0.7 | Toxoplasma | N.A. | penetrating | non-carcinogenic | mutagenic | N.A. |
| PCC2 | MMV688508 | C19H19N4O4F | 358.4 | 26 | 3.7 | 16.9 | -0.5 | Mycobacterium | N.A. | penetrating | non-carcinogenic | mutagenic | N.A. |
| PCC5 | MMV687730 | C23H22N4O2 | 384.5 | 28 | 13.4 | 0.4 | -0.5 | Mycobacterium | N.A. | penetrating | carcinogenic | non-mutagenic | N.A. |
| PCC6 | MMV687251 | C8H9N3O4S2 | 275.3 | 30 | 12.0 | 0.3 | -0.5 | Mycobacterium | N.A. | penetrating | non-carcinogenic | non-mutagenic | 13.3 (mg/kg_bw/day) |
| PCC9 | MMV688361 | C21H19N5O | 357.4 | 32 | 11.5 | 17.2 | -0.7 | Kinetoplastids | N.A. | penetrating | carcinogenic | mutagenic | N.A. |
| PCD10 | MMV689029 | C26H26N4O4S | 490.6 | 33 | 19.1 | 10.5 | 0.8 | Kinetoplastids | N.A. | penetrating | carcinogenic | mutagenic | 11.9 (mg/kg_bw/day) |
| PCD11 | MMV1030799 | C20H18N4O | 330.4 | 28 | 11.7 | 1.5 | -0.3 | Plasmodium | 6.62 (mg/L) | non-penetrating | carcinogenic | mutagenic | N.A. |
| PCE5 | MMV687146 | C19H26N2O | 298.4 | 21 | 25.8 | 15.6 | 0.6 | Mycobacterium | N.A. | penetrating | non-carcinogenic | mutagenic | N.A. |
| PCE6 | MMV687696 | C29H28N4O2ClF3 | 557.0 | 26 | 20.6 | 18.2 | -0.7 | Mycobacterium | N.A. | non-penetrating | carcinogenic | mutagenic | N.A. |
| PCE7 | MMV687170 | C17H13N4O2Cl | 340.8 | 34 | 25.3 | 13.1 | 0.0 | Mycobacterium | N.A. | penetrating | carcinogenic | mutagenic | N.A. |
| PCE8 | MMV690102 | C22H23N7O2 | 417.5 | 38 | 15.6 | 2.1 | 0.7 | Kinetoplastids | N.A. | penetrating | non-carcinogenic | mutagenic | 3.27 (mg/kg_bw/day) |
| PCE11 | MMV1029203 | C20H17N5O5 | 375.5 | 33 | 29.7 | 7.1 | -0.4 | Plasmodium | 100.0 (mg/L) | penetrating | carcinogenic | mutagenic | N.A. |
| PCF2 | MMV676053 | C18H16N3O3Cl | 357.8 | 38 | 12.5 | 1.9 | 0.6 | Cryptosporidium | 194.0 (mg/L) | penetrating | non-carcinogenic | mutagenic | 0.991 (mg/kg_bw/day) |
| PCF3 | MMV688179 | C18H16N6OCl2 | 476.2 | 35 | 32.0 | 3.1 | -0.1 | Kinetoplastids | 4.62 (mg/L) | penetrating | carcinogenic | mutagenic | 1.41 (mg/kg_bw/day) |
| PCF4 | MMV023969 | C24H24N4O5 | 453.0 | 48 | 21.8 | 1.5 | 0.3 | Mycobacterium | N.A. | N.A. | carcinogenic | mutagenic | N.A. |
| PCF5 | MMV687138 | C19H17N3O3S | 339.4 | 26 | 11.6 | 14.7 | -0.2 | Mycobacterium | 524.0 (mg/L) | penetrating | non-carcinogenic | mutagenic | 89.7 (mg/kg_bw/day) |
| PCF11 | MMV688921 | C23H18N3O5Cl | 451.9 | 31 | 43.0 | 2.4 | -0.4 | Aedes aegypti - chikungunya | N.A. | penetrating | carcinogenic | mutagenic | N.A. |
| PCG9 | MMV688891 | C18H11N4O4F3 | 442.2 | 25 | 10.9 | 25.7 | -0.5 | Mycobacterium | N.A. | penetrating | carcinogenic | mutagenic | 1.25 (mg/kg_bw/day) |
| PDH11 | MMV688980 | C16H18N3O2F5 | 335.4 | 33 | 38.2 | 21.2 | 0.2 | Plasmodium | N.A. | penetrating | carcinogenic | mutagenic | N.A. |
| PEC8 | MMV687765 | C25H26N6O | 463 | 28 | 25.5 | 20.6 | -0.8 | Mycobacterium | N.A. | penetrating | non-carcinogenic | mutagenic | N.A. |
| PE69 | MMV084864 | C17H12N6O | 316.3 | 40 | 18.7 | 17.3 | 0.8 | Plasmodium | 14.2 (mg/L) | penetrating | non-carcinogenic | mutagenic | N.A. |

¹ Coordinates used to identify compounds in each plate. TCBZ is Triclabendazole.² ID codes assigned by the Medicines for Malaria Venture (MMV) agency. N.A. is not applicable³ Measured at 72-hr post drug exposure on metacercariae (MC) stage. Results are the mean and standard deviation of triplicate experiments at a concentration of 33 μ M.⁴ Compounds were serially diluted and tested in culture. Results are means from triplicate experiments. Fasciolicidal activity values determined by CompuSyn. R is the correlation coefficient⁵ Activity shown in agents causing others infectious diseases as obtained from www.mmv.org.⁶ Predictions using lazar program (<https://lazar.in-silico.de/predict>). N.A. is not available

SD is standard deviation

† Mean and standard deviation of 10 individual experiments performed in 5 plates

* Data obtained from <https://drugs.ncats.io>Molecular formula and weights were obtained from www.mmv.org. For TCBZ these values were obtained from ChEMBL (<https://www.ebi.ac.uk/chembl/>).

Table 2. Biological activity of the compounds screened on adult worms.

| Compound plate code ¹ | MMV ID ² | Mean % Mortality on adults ³ | SD (%) |
|----------------------------------|---------------------|---|--------|
| TCBZ | N.A. | 100 † | 0 |
| PAG6 | MMV063404 | 67 | 33.3 |
| <i>PBF4</i> | <i>MMV003270</i> | 67 | 0 |
| PBF11 | MMV085210 | 0 | 0 |
| <i>PBH10</i> | <i>MMV676380</i> | 78 | 19.2 |
| PCC5 | MMV687730 | 11 | 19.2 |
| PCC6 | MMV687251 | 33 | 33.3 |
| PCC11 | MMV1030799 | 67 | 33.3 |
| <i>PCE8</i> | <i>MMV690102</i> | 56 | 19.2 |
| <i>PCE11</i> | <i>MMV1029203</i> | 78 | 19.2 |
| PCF2 | MMV676053 | 0 | 0 |
| PCF3 | MMV688179 | 22 | 19.2 |
| PCF4 | MMV023969 | 33 | 33.3 |
| PCF11 | MMV688921 | 67 | 33.3 |

¹ Coordinates used to identify compounds in each plate. TCBZ is Triclabendazole.

² ID codes assigned by the MMV agency.

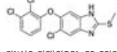

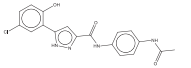
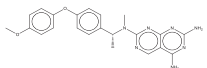
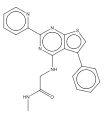
³ Measured at 48-hr post drug exposure on adult worms. Results are means and standard deviation from triplicate experiments at a concentration of 33 μ M.

† Mean and standard deviation of 6 individual experiments performed in 3 plates

Compounds in italics were selected for IC50 on adult worms and cytotoxicity assay on cell lines.

N.A. is not applicable

Table 3. Hits compounds selected for their fasciolicidal activity as a new effective drugs against *F. hepatica*.

| Structure/Smiles ¹ | Compound ID/Drug name ² | Drug name | Molecular weight | AlogP ³ | In vitro fasciolicidal assesment ⁴ | | In vitro cytotoxicity ⁴ | | | | Cytotoxicity data from other studies ⁵ | | |
|---|------------------------------------|--|------------------|--------------------|---|---------------|------------------------------------|----------------|-----------------|----------------|---|----------------|----------------|
| | | | | | Adult IC ₅₀ (µM) | CI 95% | 3T3 GI50 (µM) | H460 GI50 (µM) | DU145 GI50 (µM) | HT29 GI50 (µM) | HepG2 CC50 (µM) | HL60 CC50 (µM) | MRC5 CC50 (µM) |
|  <chem>CSC1=NC2=C(N1)C=C(C1C)C(OC1=CC=CC(C)=CC1)=C2</chem> | Triclabendazole | 6-Chloro-5-(2,3-dichlorophenoxy)-2-(methylthio)-1H-benzod[<i>e</i>]imidazole | 359.66 | 6 | 15* | 22.80 | 32.62 | 35.22 | 37.80 | | N.A. | N.A. | N.A. |
|  <chem>ClC1=CC(N=C(N)O2)=C2=C1</chem> | MMV003270/Zovaxolamine | 2-Amino-5-chlorobenzoxazole | 168.58 | 2.1 | 9.37 | 1.45 to 53.88 | > 23.73 | > 23.73 | > 23.73 | > 23.73 | > 80 | N.A. | N.A. |
|  <chem>CC(=O)Nc1ccc(NC(=O)c2cc(nh)2)c3cc(C)ccc3O)c1</chem> | MMV676380 | N-(4-Acetamidophenyl)-3-(5-chloro-2-hydroxyphenyl)-1H-pyrazole-5-carboxamide | 370.79 | 3.7 | 6.68 | 4.39 to 10.06 | > 10.79 | > 10.79 | > 10.79 | > 10.79 | > 80 | > 50 | N.A. |
|  <chem>CC1ccc(Oc2ccc(cc2)C@@H(C)N(C)C3cnc4c(N)nc(N)nc4n3)cc1</chem> | MMV690102 | 2-N-[1-[4-(4-methoxyphenoxy)phenyl]ethyl]-2-N-methylpyrimido[4,5-d]pyrimidine-2,5,7-triamine | 417.46 | 3.6 | 2.14 | 1.16 to 4.82 | 4.86 | 0.95 | 9.58 | 11.00 | 2.87 | N.A. | 5.44 |
|  <chem>CNC(=O)C1c1nc(nc2sc(C3CCCC3)C12)C4CCCC4</chem> | MMV1029203 | N-methyl-2-[[5-phenyl-2-[2-pyridyl]thieno[3,2- <i>e</i>]pyrimidin-4-yl]amino]acetamide | 375.45 | 3.58 | 4.32 | 2.82 to 6.60 | > 10.65 | > 10.65 | > 10.65 | > 10.65 | 22 | N.A. | N.A. |

¹ ID codes assigned by the MMV agency.² Data obtained from the MMV agency website 'Biological activity'.³ Compounds were serially diluted and tested in culture. Results are mean from triplicate assays.⁴ Cytotoxicity assays on cancer cell lines 3T3, H460, DU145 AND HT29. Results are mean from duplicate tests.⁵ As obtained from www.mmv.org. HepG2 is hepatocellular carcinoma, HL60 is Human promyelocytic leukemia cells, MRC5 is fibroblasts derived from lung.* Data obtained from <https://drugs.ncats.io>

Table 4. Potential targets of the 13 hits and TCBZ tested in adult worms assays.

| Compound plate code ¹ | MMV code ² | Targets predicted | Target ³ | | | |
|----------------------------------|-----------------------|-------------------|---------------------|--|-------------------------------|--|
| | | | CHEMBL ID | Preferred name | Organism | Protein target classification |
| PBH10 | MMV676380 | 3 | CHEMBL2535 | Glucose transporter | <i>Homo sapiens</i> | transporter > electrochemical transporter > slc superfamily of solute carriers > slc02 family of hexose and sugar alcohol transporters |
| | | | CHEMBL4697 | Hexose transporter 1 | <i>Plasmodium falciparum</i> | |
| PCE11 | MMV1029203 | 1 | CHEMBL3431938 | Glucose transporter | <i>Leishmania mexicana</i> | transporter |
| | | | CHEMBL3879831 | Ferrochelatase | <i>Homo sapiens</i> | unclassified protein |
| PCF2 | MMV676053 | 1 | CHEMBL6145 | Inosine-5'-monophosphate dehydrogenase, probable | <i>Cryptosporidium parvum</i> | enzyme |
| PCF4 | MMV023969 | 3 | CHEMBL2535 | Glucose transporter | <i>Homo sapiens</i> | transporter > electrochemical transporter > slc superfamily of solute carriers > slc02 family of hexose and sugar alcohol transporters |
| | | | CHEMBL4697 | Hexose transporter 1 | <i>Plasmodium falciparum</i> | |
| PBF4 | MMV003270 | 19 | CHEMBL3431938 | Glucose transporter | <i>Leishmania mexicana</i> | unclassified protein |
| | | | CHEMBL340 | Cytochrome P450 3A4 | <i>Homo sapiens</i> | enzyme > cytochrome p450 > cytochrome p450 family 3 > cytochrome p450 family 3a > cytochrome p450 3a4 |
| | | | CHEMBL289 | Cytochrome P450 2D6 | <i>Homo sapiens</i> | enzyme > cytochrome p450 > cytochrome p450 family 2 > cytochrome p450 family 2d > cytochrome p450 2d6 |
| | | | CHEMBL3397 | Cytochrome P450 2C9 | <i>Homo sapiens</i> | enzyme > cytochrome p450 > cytochrome p450 family 2 > cytochrome p450 family 2c > cytochrome p450 2c9 |
| | | | CHEMBL3622 | Cytochrome P450 2C19 | <i>Homo sapiens</i> | enzyme > cytochrome p450 > cytochrome p450 family 2 > cytochrome p450 family 2c > cytochrome p450 2c19 |
| | | | CHEMBL3356 | Cytochrome P450 1A2 | <i>Homo sapiens</i> | enzyme > cytochrome p450 > cytochrome p450 family 1 > cytochrome p450 family 1a > cytochrome p450 1a2 |
| | | | CHEMBL4040 | MAP kinase ERK2 | <i>Homo sapiens</i> | enzyme > kinase > protein kinase > cmgc protein kinase group > cmgc protein kinase mapk family > cmgc protein kinase erk subfamily |

| | | | | | | |
|------|------|----|---------------|--|---------------------|---|
| | | | CHEMBL2903 | Arachidonate 15-lipoxygenase | <i>Homo sapiens</i> | enzyme |
| | | | CHEMBL2756 | Monoamine oxidase B | <i>Bos taurus</i> | enzyme |
| | | | CHEMBL3254 | Monoamine oxidase A | <i>Bos taurus</i> | enzyme |
| | | | CHEMBL1075094 | Nuclear factor erythroid 2-related factor 2 | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1293224 | Microtubule-associated protein tau | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL2362981 | TAR DNA-binding protein 43 | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1293235 | Prelamin-A/C | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1781865 | 78 kDa glucose-regulated protein | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1977 | Vitamin D receptor | <i>Homo sapiens</i> | transcription factor > nuclear receptor > nuclear hormone receptor subfamily 1 > nuclear hormone receptor subfamily 1 group i > nuclear hormone receptor subfamily 1 group i member 1 |
| | | | CHEMBL1947 | Thyroid hormone receptor beta-1 | <i>Homo sapiens</i> | transcription factor > nuclear receptor > nuclear hormone receptor subfamily 1 > nuclear hormone receptor subfamily 1 group a > nuclear hormone receptor subfamily 1 group a member 2 |
| | | | CHEMBL1697668 | Solute carrier organic anion transporter family member 1B1 | <i>Homo sapiens</i> | transporter > electrochemical transporter > slc superfamily of solute carriers > slc21/slco family of organic anion transporting polypeptides |
| | | | CHEMBL1743121 | Solute carrier organic anion transporter family member 1B3 | <i>Homo sapiens</i> | |
| | | | CHEMBL1741193 | Chromobox protein homolog 1 | <i>Homo sapiens</i> | epigenetic regulator > reader > methyl-lysine/arginine binding protein > chromodomain |
| TCBZ | N.A. | 27 | CHEMBL1293278 | Geminin | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1075094 | Nuclear factor erythroid 2-related factor 2 | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1293224 | Microtubule-associated protein tau | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL1293258 | Mothers against decapentaplegic homolog 3 | <i>Homo sapiens</i> | unclassified protein |
| | | | CHEMBL2362981 | TAR DNA-binding protein 43 | <i>Homo sapiens</i> | unclassified protein |

| | | | |
|---------------|---|---|---|
| CHEMBL2146310 | Aberrant vpr protein | <i>Human immunodeficiency virus 1</i> | unclassified protein |
| CHEMBL2029198 | Rap guanine nucleotide exchange factor 4 | <i>Homo sapiens</i> | unclassified protein |
| CHEMBL6152 | Alpha-synuclein | <i>Homo sapiens</i> | unclassified protein |
| CHEMBL1293191 | Transcriptional regulator ERG | <i>Homo sapiens</i> | unclassified protein |
| CHEMBL2007624 | Peripheral myelin protein 22 | <i>Rattus norvegicus</i> | unclassified protein |
| CHEMBL1795086 | HSP90 | <i>Plasmodium falciparum 3D7</i> | unclassified protein |
| CHEMBL5567 | Luciferin 4-monoxygenase | <i>Photinus pyralis</i> | enzyme |
| CHEMBL2007625 | Isocitrate dehydrogenase [NADP] cytoplasmic | <i>Homo sapiens</i> | enzyme |
| CHEMBL3563 | Cruzipain | <i>Trypanosoma cruzi</i> | enzyme > protease > cysteine protease > cysteine protease ca clan > cysteine protease c1a family |
| CHEMBL1293248 | 4'-phosphopantetheinyl transferase ffp | <i>Bacillus subtilis</i> | enzyme |
| CHEMBL1795087 | Ubiquitin carboxyl-terminal hydrolase 1 | <i>Homo sapiens</i> | enzyme |
| CHEMBL1293234 | Putative fructose-1,6-bisphosphate aldolase | <i>Giardia intestinalis</i> | enzyme |
| CHEMBL1293228 | Streptokinase A | <i>Streptococcus pyogenes serotype M1</i> | enzyme > kinase |
| CHEMBL2524 | Alpha-galactosidase A | <i>Homo sapiens</i> | enzyme |
| CHEMBL1784 | Glucagon-like peptide 1 receptor | <i>Homo sapiens</i> | membrane receptor > family b g protein-coupled receptor > peptide receptor (family b gpccr) > glucagon-like receptor > glucagon-like peptide receptor |
| CHEMBL1793 | Parathyroid hormone receptor | <i>Homo sapiens</i> | membrane receptor > family b g protein-coupled receptor > peptide receptor (family b gpccr) > parathyroid hormone receptor > parathyroid hormone receptor |
| CHEMBL5162 | Neuropeptide S receptor | <i>Homo sapiens</i> | membrane receptor > family a g protein-coupled receptor > peptide receptor (family a gpccr) > short peptide receptor (family a gpccr) > neuropeptide receptor |

| | | | | | | |
|-------|------------|-----------|---------------|--|---------------------|---|
| | | | CHEMBL1293231 | Nuclear receptor ROR-gamma | <i>Mus musculus</i> | transcription factor > nuclear receptor > nuclear hormone receptor subfamily 1 > nuclear hormone receptor subfamily 1 group f > nuclear hormone receptor subfamily 1 group f member 3 |
| | | | CHEMBL1871 | Androgen Receptor | <i>Homo sapiens</i> | transcription factor > nuclear receptor > nuclear hormone receptor subfamily 3 > nuclear hormone receptor subfamily 3 group c > nuclear hormone receptor subfamily 3 group c member 4 |
| | | | CHEMBL3880 | Heat shock protein HSP 90-alpha | <i>Homo sapiens</i> | other cytosolic protein |
| | | | CHEMBL6032 | Histone-lysine N-methyltransferase, H3 lysine-9 specific 3 | <i>Homo sapiens</i> | epigenetic regulator > writer > protein methyltransferase |
| | | | CHEMBL4377 | Guanine nucleotide-binding protein G(s), subunit alpha | <i>Homo sapiens</i> | other membrane protein |
| PAG6 | MMV063404 | No target | N.A. | N.A. | N.A. | N.A. |
| PCC5 | MMV687730 | No target | N.A. | N.A. | N.A. | N.A. |
| PCC6 | MMV687251 | No target | N.A. | N.A. | N.A. | N.A. |
| PCD11 | MMV1030799 | No target | N.A. | N.A. | N.A. | N.A. |
| PCE8 | MMV690102 | No target | N.A. | N.A. | N.A. | N.A. |
| PBF11 | MMV085210 | No target | N.A. | N.A. | N.A. | N.A. |
| PCF3 | MMV688179 | No target | N.A. | N.A. | N.A. | N.A. |
| PCF11 | MMV688921 | No target | N.A. | N.A. | N.A. | N.A. |

¹ Coordinates used to identify compounds in each plate

² ID codes assigned by the MMV agency.

³ By consulting ChEMBL.

N.A is not available