

Accepted Manuscript

Title: Downstream Effects of Haemoglobinase Inhibition in *Plasmodium falciparum*-Infected Erythrocytes

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PII: S0166-6851(10)00130-1
DOI: doi:10.1016/j.molbiopara.2010.05.007
Reference: MOLBIO 10436

To appear in: *Molecular & Biochemical Parasitology*

Received date: 26-6-2009
Revised date: 28-4-2010
Accepted date: 8-5-2010

Please cite this article as: Naughton JA, Nasizadeh S, Bell A, Downstream Effects of Haemoglobinase Inhibition in *Plasmodium falciparum*-Infected Erythrocytes, *Molecular & Biochemical Parasitology* (2008), doi:10.1016/j.molbiopara.2010.05.007

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Downstream Effects of Haemoglobinase Inhibition in *Plasmodium falciparum*-Infected Erythrocytes

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Our data suggest that the likely primary downstream effect of inhibition of hemoglobin degradation in erythrocytic *Plasmodium falciparum* is blockade of protein synthesis rather than premature host cell lysis.

QuickTime™ and a
decompressor
are needed to see this picture.

1 **Downstream Effects of Haemoglobinase Inhibition in**
2 ***Plasmodium falciparum*-Infected Erythrocytes**

3
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13
14 *Abbreviations:* DMSO, dimethylsulphoxide; E-64, L-transepoxy-succinyl-leucylamido-
15 (4-guanidino)-butane; IC₅₀, 50% inhibitory concentration; iRBC, infected red blood cell
16 (erythrocyte); MACS, magnet-activated cell sorting; PBS, phosphate-buffered saline; p.i.,
17 post-invasion; PM-I, plasmepsin inhibitor I; SSC, salt sodium citrate; Z-FA-FMK, N-
18 CBZ-phenylalanyl-alanyl-fluoromethyl ketone.

19
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22

23 **Abstract**

24

25 Blood-stage malarial parasites (*Plasmodium falciparum*) digest large quantities of host
26 haemoglobin during their asexual development in erythrocytes. The haemoglobin
27 digestion pathway, involving a succession of cleavages by various peptidases, appears to
28 be essential for parasite development and has received much attention as an antimalarial
29 drug target. A variety of peptidase inhibitors that have potent antimalarial activity are
30 believed to inhibit and/or kill parasites by blocking haemoglobin digestion. It has not
31 however been established how such a blockage might lead to parasite death. The answer
32 to this question should lie in identifying the affected physiological function, but the
33 purpose of excess haemoglobin digestion by *P. falciparum* has for many years been the
34 subject of debate. The process was traditionally believed to be nutritional until Lew VL
35 *et al.* [Blood 2003;101:4189-94] suggested that it is linked to volume control of the
36 infected erythrocyte and is necessary to prevent premature osmotic lysis of the host cell.
37 Their model predicts that sufficient inhibition of haemoglobin degradation should result
38 in premature haemolysis. In this study we examined the downstream effects of reduced
39 haemoglobin digestion on osmoprotection and nutrition. We found that inhibitors of
40 haemoglobinases (plasmepsins, falcipains and aminopeptidases) did not cause premature
41 haemolysis. The inhibitors did however block parasite development and this effect
42 corresponded to a strong inhibition of protein synthesis. The effect on protein synthesis
43 (i) occurred at inhibitor concentrations and times of exposure that were relevant to
44 parasite growth inhibition, (ii) was observed with different chemical classes of inhibitor,
45 and (iii) was synergistic when a plasmepsin and a falcipain inhibitor were combined,

46 reflecting the well-established antimalarial synergism of the combination. Taken
47 together, the results suggest that the likely primary downstream effect of inhibition of
48 hemoglobin degradation is amino acid depletion, leading to blockade of protein synthesis,
49 and that the parasite probably degrades globin for nutritional purposes.

50

51 *Keywords:* malaria; *Plasmodium falciparum*; antimalarial drug; haemoglobin; peptidase;
52 protein synthesis.

53

54

55 **1. Introduction**

56

57 Erythrocytic stages of the malarial parasite, *Plasmodium falciparum*, have developed a
58 complex proteolytic machinery for haemoglobin digestion [1]. This is a highly energy-
59 consuming process and presents the parasite with a serious toxic waste disposal problem.
60 Why these parasites exert such efforts to ingest and digest haemoglobin and detoxify
61 haem has been the subject of considerable investigation and debate [2, 3]. This question
62 is particularly important because interference with haemoglobin digestion is presumed to
63 be the basis of action of a number of antimalarial agents in use or in development [4].

64

65 After invading erythrocytes, *P. falciparum* ingests erythrocyte cytoplasm at the
66 cytostome and transports it to the digestive (food) vacuole (DV) [5-7]. The *P. falciparum*
67 DV is optimised for haemoglobin digestion. This catabolic process results in the
68 digestion of approximately 75% of infected red blood cell (iRBC) haemoglobin [5]. The

69 haemoglobin tetramer is broken down to two components: globin and haem [1]. A
70 cascade of different haemoglobinasases degrades globin into its constituent amino acids [4].
71 Haem, which is toxic to parasites, is detoxified by the formation of haemozoin (malaria
72 pigment) [8]. The peak of haemoglobin degradation occurs in the trophozoite and early
73 schizont stages of parasite development.

74

75 In the DV, four different groups of peptidases have been shown to play a role in
76 haemoglobin degradation [4]: aspartic peptidases (plasmepsins I, II, IV and histoaspartic
77 peptidase [HAP, plasmepsin III]) [9, 10], cysteine peptidases (falcipains 2, 2' and 3) [11,
78 12], a metallopeptidase (falcilysin) [13] and a dipeptidylpeptidase (dipeptidyl
79 aminopeptidase I) [14]. Aminopeptidases are proposed to be involved in the terminal
80 stages of haemoglobin degradation [15-19], though there is scant evidence for this
81 contention in studies of intact cells. Genetic knock-out [15, 20, 21-25] and inhibitor [4,
82 9, 11, 26-29] studies suggest that some of these peptidases are essential for parasite
83 growth but in other cases there is a high degree of redundancy between them. In spite of
84 the substantial literature on the antimalarial effects of haemoglobinase inhibitors, it has
85 not been established how blockage of haemoglobin digestion might lead to arrest of
86 intraerythrocytic development and/or death of the parasite. Presumably the answer to this
87 question relates to the normal physiological function of haemoglobin digestion.

88

89 It has long been considered that the likely function of haemoglobin digestion is to supply
90 malaria parasites with amino acids, especially since they have limited ability for *de novo*
91 amino acid synthesis [4]. This assumption relies on different observations (as reviewed

92 in [1]), particularly (i) the detection of amino acids from radiolabeled haemoglobin in
93 parasite proteins and (ii) the increased susceptibility to cysteine and aspartic peptidase
94 inhibitors of parasites grown in medium with just the 5 amino acids that are absent
95 (isoleucine) or in low numbers (cysteine, glutamine, glutamate and methionine) in
96 haemoglobin compared with parasites grown in full medium. Recent observations have
97 however questioned the nutritional significance of haemoglobin hydrolysis. *P.*
98 *falciparum* can import all the amino acids from the culture medium (or serum *in vivo*) and
99 have a significant though limited capacity for *de novo* amino acid synthesis [1]. A large
100 proportion of the amino acids resulting from haemoglobin hydrolysis are exported to the
101 erythrocytic cytoplasm [5] and measurements by Krugliak et al [30] indicated that the
102 proportion of amino acids derived from hydrolyzed haemoglobin that was incorporated
103 into parasite proteins was only ~16%.

104

105 Lew et al [2, 31, 32] have integrated the question of the function of haemoglobin
106 degradation with that of the mechanism by which iRBCs retain their osmotic stability and
107 integrity during the 48-h asexual cycle despite a significant increase in the permeability
108 of the erythrocyte plasma membrane to different ions and nutrients (see also the article by
109 Allen & Kirk [33]). Using a mathematical model that included different factors known to
110 influence erythrocyte volume and homeostasis of iRBCs, they predicted volume changes
111 of the iRBC at different stages of parasite growth. The predicted volume changes were
112 experimentally tested and their results supported the model's predictions. They
113 concluded that excess haemoglobin consumption, which reduces the colloid-osmotic
114 pressure within the iRBC, is essential for maintaining the osmotic stability of the infected

115 cell for the 48 h of parasite development in the erythrocyte. An important implication, as
116 shown in Lew et al [31] (Fig. 5), is that inhibition of haemoglobin digestion should
117 increase the osmotic stress on the iRBC resulting in premature lysis of the iRBC. This
118 should be readily demonstrable using relevant concentrations of the peptidase inhibitors
119 previously shown to block haemoglobin digestion. Moreover, we contend here that if the
120 primary purpose of haemoglobin digestion is osmotic protection, the osmotic lysis effect
121 should be apparent at lower concentrations of inhibitors than those that affect
122 intraerythrocytic development.

123

124 In this work, we have investigated the downstream effects of inhibitors of plasmepsins,
125 falcipains and aminopeptidases as inhibitors of haemoglobin digestion. Under our
126 experimental conditions, no premature osmotic lysis was observed but protein synthesis
127 was significantly reduced at inhibitor concentrations and times of exposure that were
128 relevant to the known antimalarial effects. Our results support the notion that the primary
129 function of haemoglobin degradation is to provide amino acids for protein synthesis.

130

131

132 **2. Materials and Methods**

133

134 *2.1. Reagents*

135

136 All inhibitors were purchased from Sigma Aldrich (Dublin, Ireland) except for the
137 plasmepsin inhibitor PM-I (Fig. 1), which was kindly provided by Drs. C. Binkert and C.

138 Boss (Drug Discovery, Chemistry & Biology, Actelion Pharmaceuticals Ltd., Allschwil,
139 Switzerland). PM-I, bestatin ([[(2S, 3R)-3-amino-2-hydroxy-4-phenylbutanoyl]-L-
140 leucine), N-CBZ-Phe-Ala-fluoromethyl ketone (Z-FA-FMK) and taxol were prepared in
141 dimethylsulphoxide (DMSO) and stored at -20°C . L-trans epoxy-succinyl-leucylamido-
142 (4-guanidino)-butane (E-64) was dissolved in purified water (Millipore Synthesis AQ,
143 Carrigtwohill, County Cork, Ireland) and sterile filtered. Inhibitors were diluted in
144 complete medium freshly before each experiment. As a control for inhibitors prepared in
145 DMSO, the same amount of DMSO solvent was diluted in complete medium and added
146 to the parasite culture. L-[^{35}S] Methionine (specific activity of 1110 Ci/mmol) and [^3H]
147 isoleucine (99 Ci/mmol) were obtained from Amersham (GE Healthcare Ltd., UK).

148

149 2.2. Parasite Culture

150

151 *P. falciparum* clone 3D7 (obtained from M. Grainger, National Institute of Medical
152 Research, London, UK) was cultivated in human O^+ erythrocytes as previously described
153 [34]. Age-selection of the parasites was performed by two steps of magnet-activated cell
154 sorting (MACS: Miltenyi Biotec, Surrey, UK) [35] which isolated late trophozoite- and
155 schizont-infected erythrocytes from uninfected erythrocytes and those infected with less
156 mature forms. The MACS operations were timed to give parasites of an age range of ~8–
157 10 h. After age-selection the parasites were re-cultured to the desired approximate ages
158 for the experiment (see individual experiments below). The age ranges of the parasite
159 cultures were estimated by measuring the time between the appearance of the first rings
160 and the disappearance of the last segmenters. The parasites were continuously cultured in

161 human erythrocytes in RPMI 1640 supplemented with HEPES (25 mM), Albumax® II
162 (0.5% w/v) (Gibco, Auckland, New Zealand), hypoxanthine (50 µg/ml), glucose (0.08%
163 w/v), NaHCO₃ (0.18% w/v) and gentamicin (0.02 mg/ml) [36]. The parasites were
164 cultured at 2.5% hematocrit in candle jars at 37°C. Cultures were examined
165 microscopically with the aid of Giemsa staining. The different stages of erythrocytic
166 parasites were defined as previously described [37].

167

168 *2.3. Inhibitor Susceptibility Assays*

169

170 Fifty-% inhibitory concentrations (IC₅₀) after 72 h were determined on asynchronous
171 parasites in 96-well plates by using the parasite dehydrogenase assay [38] and were
172 determined graphically.

173

174 For all other inhibitor assays, cultures consisting mainly of young or young-mid
175 trophozoites (estimated ages as shown in the figure legends and text, obtained by MACS
176 as described above) at *circa* 8% parasitemia and 2.5% hematocrit were treated with
177 different concentrations of inhibitors (1X, 5X and 25X IC₅₀) in 96- or 24-well plates.
178 Controls were parasite cultures incubated in inhibitor-free medium with or without
179 solvent (DMSO). Growth and survival of parasites was assessed by microscopic
180 examination of Giemsa-stained smears.

181

182 *2.4. Haemoglobin Release Assay*

183

184 *P. falciparum* parasites were cultured and treated as described in section 2.3 in 24-well
185 plates (with a total volume of 500 μ l) except that phenol-red-free medium was used.
186 Parasites were harvested after specific incubation times by taking samples from wells to
187 Eppendorf tubes, followed by centrifugation (800 g) for 5 min at room temperature. The
188 supernatant was removed to another tube and the above steps were repeated until no
189 pellet was observed. Two hundred μ l of the supernatant were used to measure the
190 absorbance at 405 nm in 96-well plates.

191

192 2.5. Protein Synthesis Assay

193

194 As described in section 2.3, *P. falciparum* parasites were cultured and treated with
195 inhibitors in 96-well plates and incubated for different periods. At the end of designated
196 time points, cultures were metabolically labeled with L[³⁵S]methionine (50–100 mCi/ml)
197 for 1 h. After labeling and three steps of washing with unlabelled medium, crude extracts
198 of iRBCs were prepared by three cycles of freezing-thawing. The extracts were stored at
199 -70°C until further use.

200

201 Proteins from labeled cell extracts were precipitated with trichloroacetic acid/acetone.
202 Samples were then mixed with scintillation cocktail (ICN Biomedicals Inc., Aurora,
203 Ohio, U.S.A.) and a liquid scintillation spectrometer (Packard Tri-Carb model 2100TR)
204 was used to count radioactivity incorporated into macromolecules.

205

206 2.6. Assay of Haemozoin Formation

207

208 Haemozoin was purified and quantified as described previously [23, 39]. Briefly,
209 parasites were isolated from iRBCs using 0.1% (w/v) saponin in ice-cold salt sodium
210 citrate (SSC) [17] and washed in SSC. Harvested parasites were stored at -70°C in PBS.
211 The thawed, harvested parasites were centrifuged for 20 min at 10,000 g at 20°C . The
212 pellet was resuspended in 100 μl buffer A (Tris-HCl (100 mM, pH 8.0) containing 2.5%
213 SDS) and incubated for 30 min at 37°C with vortexing after 10 min. After a
214 centrifugation step as above, the pellet was resuspended in 100 μl alkaline bicarbonate
215 (100 mM, pH 9.2). The pellet obtained after centrifugation was dissolved in 10 μl 1 M
216 NaOH and 90 μl of buffer A and incubated for 1 h at 50°C . The haemozoin content was
217 determined by measuring the absorbance at 405 nm (Multiskan Ex, ThermoScientific,
218 Basingstoke, UK) in 96-well plates. The amount of haem was calculated using the
219 extinction coefficient of $91,000\text{ cm}^{-1}\text{M}^{-1}$.

220

221 2.7. Measurement of isoleucine uptake

222 Isoleucine uptake experiments were carried out as described previously [3] with some
223 modifications. Briefly, *P. falciparum* parasites were cultured and treated as described in
224 section 2.3 in 6-well plates (with a total volume of 3 ml). At specific time points (5.5 and
225 12.5 hours) cells were depleted of amino acids by repeated washing in PBS and
226 incubation for 30 minutes at 37°C in Solution A (130 mM NaCl, 25 mM HEPES, 5 mM
227 KCl, 20 mM glucose, 0.2 mM hypoxanthine, 25 mg/l gentamicin sulphate, RPMI 1640
228 vitamins [Sigma Aldrich] and 3.25 μM glutathione) with the appropriate inhibitor
229 concentration. Cells were then magnet purified and resuspended to a parasitaemia of 80–

230 90% and haematocrit of 5% in Solution A supplemented with amino acids (alanine (356
231 μM), arginine (88 μM), asparagine (13 μM), aspartate (13 μM), cysteine (37 μM),
232 glutamate (57 μM), glutamine (476 μM), glycine (217 μM), histidine (85 μM),
233 hydroxyproline (8 μM), [^3H] isoleucine (70 μM , 6.93 mCi/ml), leucine (100 μM), lysine
234 (163 μM), methionine (17 μM), phenylalanine (100 μM), proline (165 μM), serine (128
235 μM), threonine (112 μM), tryptophan (50 μM), tyrosine (62 μM), and valine (190 μM)
236 and incubated at 20°C for 5 minutes. Duplicate 200- μl samples were taken and the
237 reaction terminated upon centrifugation of the cells (14,000 g for 2 min) through silicon
238 oil (Dow Corning) and processed for scintillation counting as described previously [42].
239 Parasite-specific influx was calculated by subtracting the accumulation into an equal
240 number of uninfected erythrocytes.

241

242

243 **3. Results**

244

245 *3.1. Effects of Peptidase Inhibitors on Osmotic Stability of iRBCs*

246

247 According to Lew et al [31] haemoglobin hydrolysis is necessary to maintain the osmotic
248 stability of iRBCs during the maturation of parasites. By using peptidase inhibitors, the
249 occurrence of premature lysis of iRBCs due to inhibition of haemoglobin degradation
250 was tested. Inhibitors of three different classes of peptidases were used. PM-I (Fig. 1),
251 which is closely related to the plasmepsin inhibitors previously described [40], is a non-
252 peptidic aspartic peptidase inhibitor that is potent against three of the DV plasmepsins (I,

253 II and IV). Given that parasites can survive without these three enzymes [21, 23, 41] but
254 PM-I nonetheless has potent antimalarial activity, this compound is likely to target other
255 plasmepsins/aspartic peptidases. E-64 and Z-FA-FMK are cysteine peptidase inhibitors
256 and bestatin is an aminopeptidase inhibitor. The IC_{50} of the compounds used were as
257 follows: PM-I (0.8 μ M), E-64 (3 μ M), Z-FA-FMK (5 μ M) and bestatin (2 μ M).

258

259 Young–mid-trophozoite parasite cultures (21–29 h p.i.), which were at the early stage of
260 haemoglobin degradation, were treated with different peptidase inhibitors. The method
261 of age-selection used in this experiment and all other experiments reported here was
262 magnet-activated cell sorting (MACS). This method was chosen in preference to the
263 more prevalent method using sorbitol lysis of mature stages in order to diminish the
264 possible negative effects on the iRBC membrane. Following incubation for various time
265 periods, the medium of the cultures was analyzed spectrophotometrically to quantify the
266 amount of released haemoglobin. According to the graph presented in Fig. 5 in [31], if
267 haemoglobin degradation decreased from 70% to ~50%, then premature lysis should take
268 place at ~45 h p.i. Since this is an estimate and it would be hard to predict the exact time
269 of maximal osmotic lysis, treated and control cultures were harvested at different time
270 points. Cultures were harvested at 14, 16, 18 and 20 h after drug treatment (39 ± 4 -, 41 ± 4 -,
271 43 ± 4 - and 45 ± 4 h p.i., respectively). Fig. 1 shows that the amount of haemoglobin
272 release in the control cultures increased as the parasites were maturing (it more than
273 tripled from 16 h to 20 h of incubation), as expected from the increasing lysis of the host
274 erythrocytes as the parasites completed their ~46-48-h cycle and released new
275 merozoites. In most of the inhibitor-treated cultures, the amount of haemoglobin release

276 remained constant during the total 20 h incubation period and in no case did it exceed that
277 of the control. In none of our experiments, regardless of concentration of inhibitor or
278 time of incubation, did we observe premature iRBC lysis (data not shown). Significant
279 differences in haemoglobin release from iRBCs in the control- and inhibitor-treated
280 cultures were however already observable after 16 h of treatment. Microscopic analysis
281 of these cultures showed that progression of parasite development was blocked. After a
282 further 13 h of incubation, ~65% of the control population consisted of rings and
283 schizonts while the inhibitor-treated cultures contained mainly what looked like mid-
284 trophozoites (i.e. the same as the starting culture). The results of control experiments
285 indicated that parasites treated with any of the three haemoglobinase inhibitors for 13 h
286 could not subsequently be revived (data not shown). These observations indicated that
287 not only was there no premature osmotic lysis, but levels of normal lysis at the time of
288 merozoite release were considerably lowered.

289

290 *3.2. Effects of Peptidase Inhibitors on Parasite Protein Synthesis*

291

292 Protein synthesis was determined by measuring the incorporation of L[³⁵S]methionine
293 into newly synthesized proteins (erythrocytes themselves lacking the capacity for protein
294 synthesis). The negative effects of peptidase inhibitors on cultures with initial ages ~16–
295 24 h p.i. were observed after 6 h (26±4 h p.i.) treatments at the 5X- and 25X IC₅₀ for E-64
296 and all concentrations for bestatin and PM-I (Fig. 2). PM-I at concentrations 5X and 25X
297 IC₅₀ also caused inhibition of protein synthesis after only 4 h incubation (~45% and
298 ~70%, respectively) (results not shown). Inhibition of protein synthesis was observed for

299 all concentrations of the inhibitors after 13 h treatment (33 ± 5 h p.i.). For the most part
300 the inhibition of protein synthesis was progressive with time, but in a few cases (e.g. 1X
301 IC_{50} bestatin) the effect was greater at 6 h than at 13 h. This observation may be
302 connected with the upregulation of protein degradation by the ubiquitin/proteasome
303 pathway in response to amino acid starvation [43].

304

305 In order to confirm that the effect of the peptidase inhibitors was not just peculiar to the
306 specific inhibitors discussed above, Z-FA-FMK (another cysteine peptidase inhibitor)
307 was applied to *P. falciparum* cultures as described in Materials and Methods and its
308 effects on protein synthesis were tested. Treatment of parasite cultures with Z-FA-FMK
309 for 6 and 13 h resulted in a decrease in protein synthesis at concentrations 5X- and 25X
310 IC_{50} (~50%) (Fig. 2). As a negative control, taxol (a drug whose antimalarial action is
311 apparently unrelated to haemoglobin digestion or protein synthesis ([44] and see section
312 3.3) had no or only marginal effects at 5X IC_{50} in parallel experiments (Fig. 2), indicating
313 that the effects of the haemoglobinase inhibitors on protein synthesis were unlikely to be
314 merely secondary consequences of growth inhibition by some other mechanism.

315

316 Plasmepsin and falcipain inhibitors have been reported to display strong synergistic
317 effects on growth of *P. falciparum* [26, 45-47]. The data in Fig. 3 confirm that the
318 combination of E-64 and PM-I also had synergistic antimalarial activity. If the effects on
319 protein synthesis shown here are relevant to growth inhibition, they should also be
320 synergistic. The combined effect of E-64 and PM-I on the inhibition of protein synthesis
321 was therefore tested by comparing parasite cultures treated with 2.5X IC_{50} of E-64 and

322 2.5X IC₅₀ PM-I in combination with those treated with 5X IC₅₀ concentrations of each
323 agent alone. If the effect were merely additive, levels of protein synthesis in cultures
324 treated with the combination would be expected to be intermediate between those in
325 cultures treated with 5X IC₅₀ of E-64 alone and those treated with 5X IC₅₀ of PM-I alone.
326 In fact, they were much lower than this (Fig. 2), indicating a synergistic effect.

327

328 *3.3. Effects of Inhibitors on Haemozoin Formation*

329

330 To confirm that the inhibitors at the concentrations and for the times used were inhibiting
331 haemoglobin degradation, the haemozoin contents in the treated and untreated cultures
332 were measured. After 6 hours of incubation (final ages 26±4 h p.i.), the higher
333 concentrations of bestatin, E-64 and PM-I had slight negative effects on haemozoin
334 formation in the parasites (~30% decrease) (data not shown). When the parasites were
335 treated for 13 h (33±4 h p.i.), there were more significant decreases in haemozoin
336 formation in those cultures treated with 5X and 25X IC₅₀ bestatin (~20% and ~45%,
337 respectively) and E-64 (~50% and ~65%, respectively) and all concentrations of PM-I
338 (~70–80%) (Fig. 4). Treatment of parasite cultures with Z-FA-FMK for 6 and 13 h
339 resulted in a decrease of haemozoin formation at all concentrations (~40–60%) (Fig. 4
340 and data not shown). These results show that the peptidase inhibitors used did inhibit
341 haemozoin formation and therefore imply that haemoglobin digestion was inhibited.
342 These were important control observations and, interestingly, we were unable to
343 demonstrate a significant effect of relevant concentrations of the aspartic peptidase
344 inhibitor pepstatin A on haemozoin accumulation (data not shown) so were therefore

345 unable to include it in the experiments above. Moreover, the degrees of inhibition
346 obtained with the various inhibitors were the same as or greater than that expected to
347 induce osmotic lysis of iRBCs according to Lew et al [31]. The negative control agent,
348 taxol, had a negligible effect on haemozoin accumulation ($92.6 \pm 2.2\%$ of control at 5X
349 IC_{50} ($0.25 \mu\text{M}$), not shown in Fig. 4).

350

351 *3.4. Effect of bestatin on isoleucine uptake, a surrogate for free leucine concentration*

352

353 The results described above suggested that the haemoglobinase inhibitors were not acting
354 by increasing osmotic lysis of iRBC but that they affected protein synthesis. The most
355 likely explanation for this effect was that inhibition of haemoglobin digestion reduced the
356 supply of amino acids required for protein synthesis. We were unable to prove this
357 contention because of the technical difficulties associated with measuring (possibly
358 transient) changes in the free amino acid concentrations within parasites. The work of
359 Martin & Kirk [3], showing the dependence of uptake of isoleucine on free intraparasitic
360 leucine, did however suggest a measurable 'surrogate' for free leucine, one of the most
361 abundant amino acids in haemoglobin. We therefore measured radiolabelled isoleucine
362 uptake in parasites treated with bestatin. As an aminopeptidase inhibitor, bestatin would
363 be expected to act most closely to the point of release of free amino acids from globin.
364 The data in Fig. 5 show that bestatin at 5X and 25X IC_{50} reduced the uptake of isoleucine
365 by ~20-25% and ~50% respectively, while the apparently irrelevant inhibitor taxol had no
366 such effect.

367

368

369 **4. Discussion**

370

371 The peptidase inhibitors active on haemoglobin digestion in *P. falciparum* are attractive
372 targets for designing anti-malarial drugs and have been explored extensively by several
373 groups [4, 9, 11, 26-29]. It is therefore surprising that the downstream consequences of
374 inhibition of haemoglobin digestion, and the mechanism by which this metabolic
375 inhibition actually kills parasites or blocks their development, have not been elucidated.
376 We addressed this question with reference to the two main theories regarding the primary
377 biological purpose of haemoglobin degradation: maintaining the osmotic stability of
378 iRBCs or providing amino acids for protein synthesis.

379

380 The haemoglobin release experiment did not show any premature iRBC lysis in cultures
381 treated with the peptidase inhibitors used in this study (Fig. 1 and data not shown).
382 Rather, the peptidase inhibitors caused a blockade of the growth and development of the
383 parasites and eventual death before they could reach the later stages of parasite growth.
384 This was also observed by Lew et al [2] when they treated *P. falciparum* at the age range
385 of 22–28 h p.i. with E-64. They concluded that using peptidase inhibitors as a tool for
386 testing the osmotic protection hypothesis of excess haemoglobin digestion was not
387 appropriate. However, if the primary purpose of haemoglobin digestion is to prevent
388 premature iRBC lysis and the model of Lew et al [31] (Fig. 5) is correct, it should be
389 possible to induce this lysis with concentrations of peptidase inhibitors lower than those
390 that affect progression through the cycle. Our data show that at a range of concentrations

391 above and below IC_{50} , including those that are capable of inhibiting haemoglobin
392 degradation (as measured by haemozoin accumulation, Fig. 4) by 50% or more, no
393 measurable premature lysis was detected. We conclude that the primary target of the
394 inhibitors lies at an earlier stage of parasitic development. It is relevant to mention here
395 that since inhibition of DV plasmepsins alone is unlikely to be sufficient to arrest parasite
396 growth [48], the primary target of PM-I is likely to be an aspartic peptidase that resides
397 outside the vacuole and therefore is not directly involved in haemoglobin digestion. Yet
398 PM-I is a powerful inhibitor of haemoglobin degradation and is in fact the most potent of
399 the peptidase inhibitors, relative to its IC_{50} , in this respect (Fig. 4). A possible
400 explanation is that PM-I may inhibit one or more peptidases catalysing proteolytic
401 activation of one or more of the haemoglobinases.

402

403 Given the proposed nutritional role of haemoglobin degradation and the fact that globin-
404 derived amino acids are incorporated into parasite proteins, we explored the possibility
405 that the reduced amino acid supply might cause arrest of parasite protein synthesis. This
406 was investigated by measuring the incorporation of L-[^{35}S]methionine into newly
407 synthesized proteins (Fig. 2). Protein synthesis inhibition was observed for all four
408 peptidase inhibitors tested after 6 hours' incubation (26 ± 4 h p.i.) and was in most cases
409 increased after 13 h incubation (33 ± 4 h p.i.). The inhibitor concentrations and times of
410 exposure were sufficiently low relative to their IC_{50} to persuade us that the effect on
411 protein synthesis was relevant to the antimalarial action of the inhibitors. This conclusion
412 was further strengthened by the observed synergistic effect of E-64 and PM-I on protein
413 synthesis inhibition (Fig. 2), which mirrors that seen on parasite growth (Fig. 3).

414

415 The most logical interpretation of our results would be that protein synthesis is a
416 downstream effect of inhibition of haemoglobin digestion, resulting from reduced supply
417 of amino acids from globin, and perhaps the primary mechanism by which
418 haemoglobinase inhibitors block the development of trophozoites and early schizonts.
419 We have not been able to establish here that amino acid depletion actually occurs as a
420 result of inhibition of haemoglobin digestion. The reduction of isoleucine uptake, which
421 is known to depend on free leucine, following treatment with bestatin is however
422 suggestive of such a connection. If this idea is correct, our results also imply that the
423 primary role of the haemoglobinases and haemoglobin digestion is to provide the parasite
424 with free amino acids, supplementing those available via uptake and *de novo* synthesis
425 and allowing protein synthesis to proceed at the level required for normal development.
426 Whatever the cause of the reduced protein synthesis, it appears that the role if any of
427 haemoglobin digestion in maintaining the osmotic stability of iRBCs is a secondary one.

428

429 An important caution is however that the inhibitors employed may have effects on targets
430 other than those involved in haemoglobin digestion. Therefore it is conceivable that the
431 effect on protein synthesis is working by some hitherto unknown mechanism. We
432 consider this unlikely, because (i) the effect was seen with four different chemical types
433 of compound directed to three different classes of peptidase, but not with control
434 compounds, and (ii) haemoglobin digestion is unquestionably the major peptidase-
435 dependent metabolic event occurring in trophozoite-stage parasites. Nonetheless we have
436 to admit the possibility of a hitherto unsuspected pathway, and for this reason it would be

437 useful to repeat these experiments with new haemoglobinase inhibitors as they become
438 available, and in different genetic backgrounds such as parasites lacking specified DV
439 peptidases.

440

441 One notable, related finding in the present study is the effect of the aminopeptidase
442 inhibitor bestatin on haemozoin accumulation (Fig. 4). Previous studies [15-17] have
443 been highly suggestive of a role of bestatin-susceptible aminopeptidases in haemoglobin
444 digestion but the experiments have been performed in cell-free systems. Moreover,
445 relevant concentrations of bestatin did not visibly affect any aspect of haemoglobin
446 metabolism as judged by electron microscopy [17]. The data in Fig. 4 are therefore the
447 first demonstration that bestatin affects haemoglobin digestion in intact parasites. That
448 this effect leads to a reduction of supply of amino acids such a leucine for protein
449 synthesis remains to be proven, but the effect of bestatin on isoleucine uptake (Fig. 5) is
450 consistent with this idea.

451

452 The presumed nutritional role of haemoglobin degradation has been demonstrated here in
453 parasites in culture with medium containing high concentrations of amino acids. Since
454 the levels of amino acids in blood plasma of malaria-infected patients in poverty-stricken
455 areas are significantly lower [48] it can be postulated that this role of haemoglobin
456 digestion may be of even more importance *in vivo*. If true, this would strengthen the
457 argument that *P. falciparum* haemoglobinases or the enzymes regulating them are
458 promising targets for new anti-malarial drugs.

459

460

461 **Acknowledgements**

462

463 This work was supported by a Health Research Board project grant to A.B. (RP/2005/57).

464 We thank Drs. Christoph Binkert and Christoph Boss of Actelion for the kind gift of PM-

465 I and Dr. Binkert for helpful discussions.

466

467

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- 606

607 **Figure legends**

608

609 **Fig. 1. Effects of peptidase inhibitors on osmotic stability of iRBCs.** Parasites age-
610 selected to young-mid trophozoites (21–29 h) post invasion were cultured in the presence
611 of different protease inhibitors and harvested after periods of 14, 16, 18 and 20 h of
612 incubation. iRBC lysis was analyzed spectrophotometrically by measuring the amount of
613 haemoglobin released into the medium (at 405 nm). These results are representative of at
614 least three separate experiments. *Inset:* Structure of Plasmepsin Inhibitor I (PM-I).

615

616 **Fig. 2. Effects of inhibitors on protein synthesis.** *P. falciparum* cultures consisting of
617 young trophozoites (16–24 h post invasion) were treated with peptidase inhibitors for
618 periods of 6 h (white bars) and 13 h (black bars). Taxol (IC₅₀ 50 nM) was included as a
619 control. L[³⁵S]Methionine was then added and the cultures were harvested after 1 h. The
620 results are shown as % of control and are an average of two to three independent
621 experiments (6 h and 13 h, respectively) and n=2 in each experiment. The error bars
622 show SEMs.

623

624 **Fig. 3. Isobologram showing interaction between E-64 and PM-I on growth of *P.***
625 ***falciparum* in culture.** Fifty-% inhibitory concentrations of E-64 alone and in the
626 presence of various concentrations of PM-I, and of PM-I alone in the presence of various
627 concentrations of E-64, were determined as described in section 2.3. The solid diagonals
628 in the isobolograms represent the theoretical line of additivity (i.e., no interaction), while
629 the values below this line indicate a synergistic effect between the two compounds. The

630 concave isobole (dashed line) was fit by inspection. Each point is a geometric average of
631 four separate experiments and error bars represent the SEM.

632

633 **Fig. 4. Effects of peptidase inhibitors on haemozoin accumulation.** Parasite cultures
634 consisting of young trophozoites (16–24 h p.i.) were incubated for 13 h with protease
635 inhibitors. At the end of the incubation time parasites were isolated and haemozoin was
636 purified as described in section 2.6. The amount of extracted haemozoin was quantified
637 spectrophotometrically at 405 nm. The results from inhibitor-treated cultures were plotted
638 as % of control (inhibitor-free culture). The amount of haemozoin in the control was 0.3–
639 1 fmol/parasite, assuming a trophozoite volume of 74 fl [49]. The data are averages from
640 two independent experiments (n=2 per experiment). The bars show SEMs.

641

642 **Fig. 5. Effect of bestatin on [³H] isoleucine uptake.** Parasite cultures consisting of
643 young trophozoites (16–24 h p.i.) were treated with bestatin or taxol at 1X, 5X or 25X
644 IC₅₀ (20X IC₅₀ for taxol because 25X IC₅₀ caused substantial parasite destruction) and
645 after 5.5 and 12.5 h cells were depleted of amino acids as described in section 2.7.
646 Infected erythrocytes were then purified by MACS, resuspended in [³H] isoleucine plus
647 other (unlabelled) amino acids, and incubated at 20°C for 5 min. Duplicate 200-μl
648 samples were taken and labelled isoleucine uptake determined as described in section 2.7.
649 The data are averages from three independent experiments (n=2 per experiment). The
650 bars show SEMs.

651

652

Fig. 1

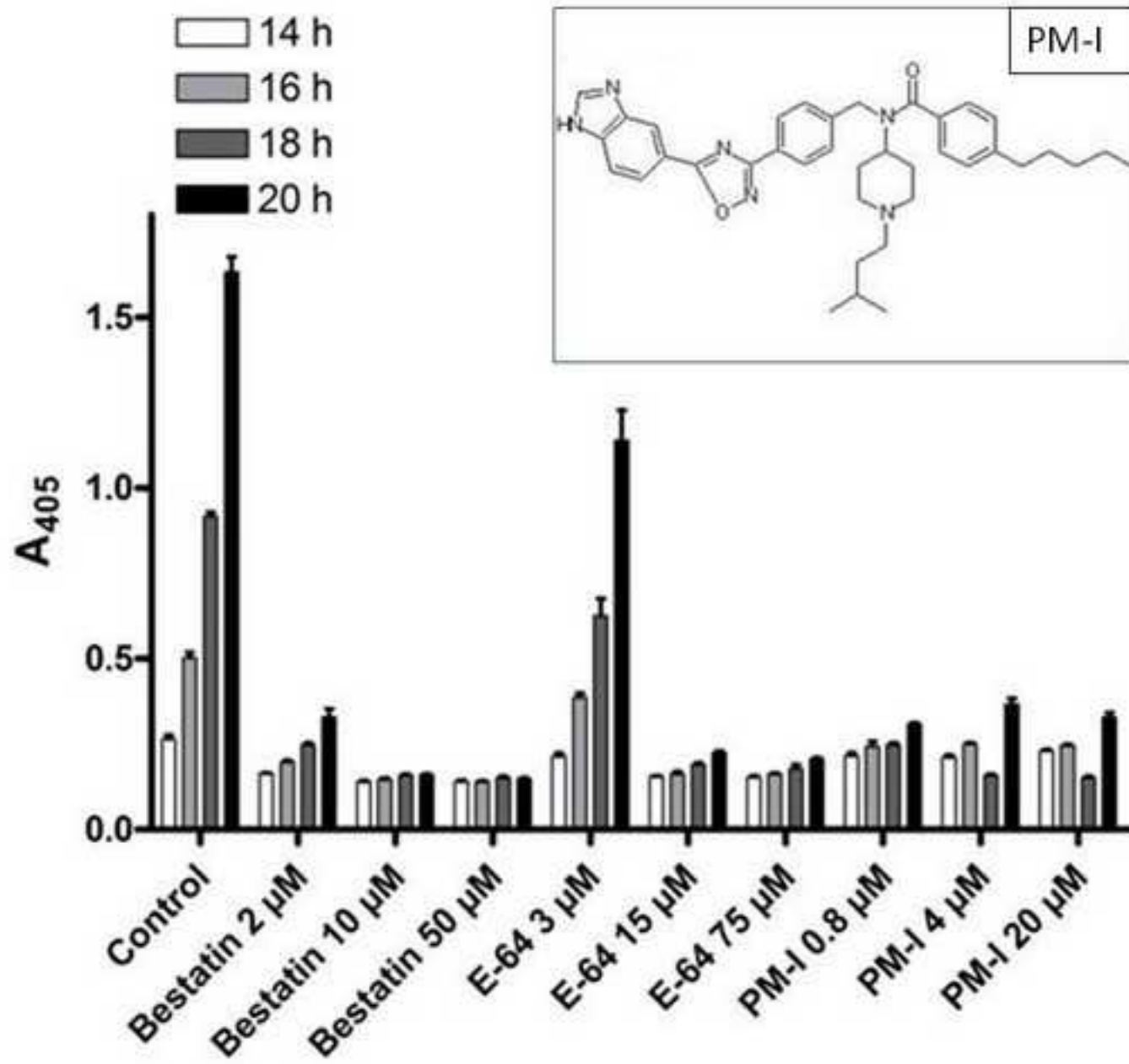


Fig. 2

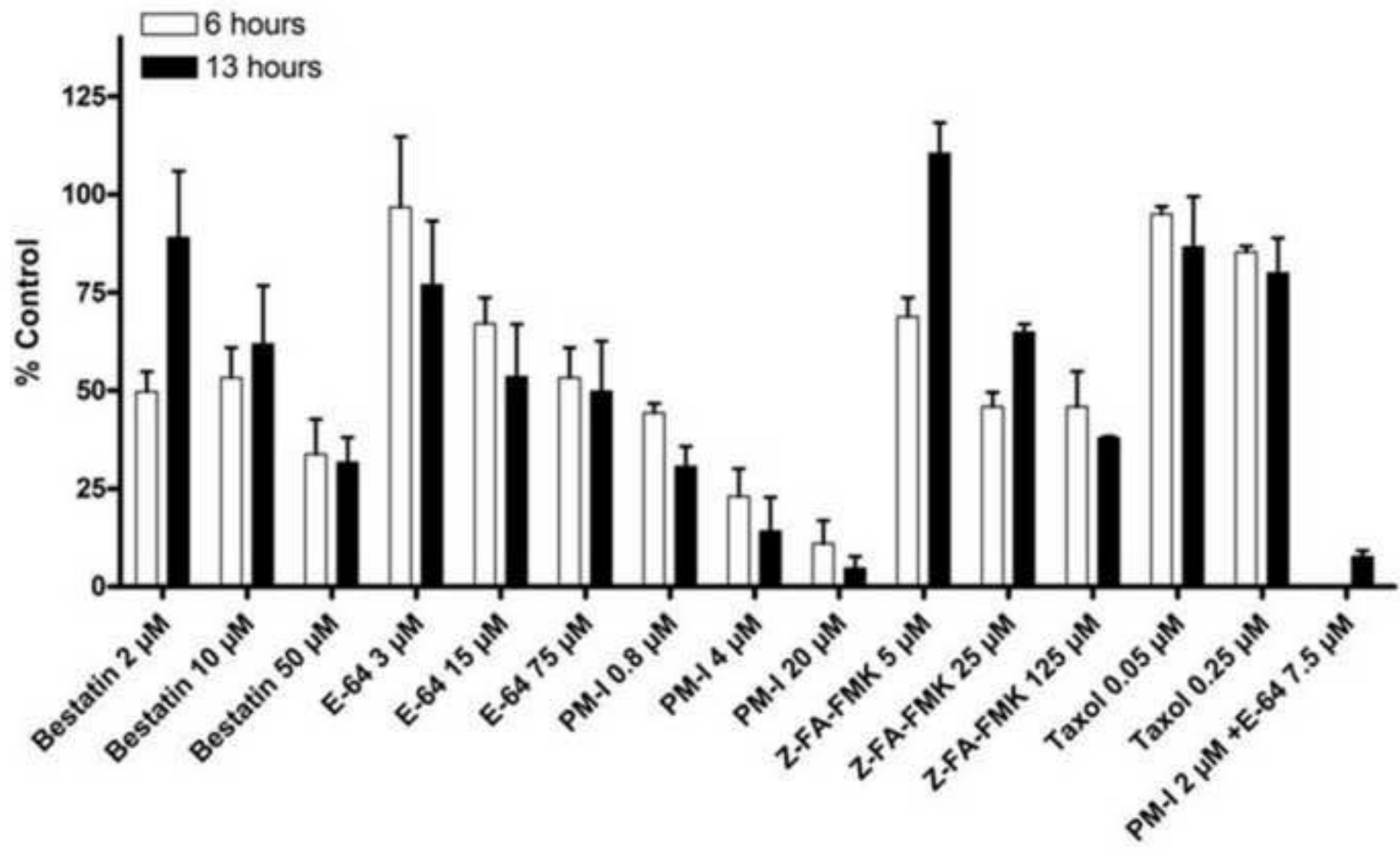


Fig. 3

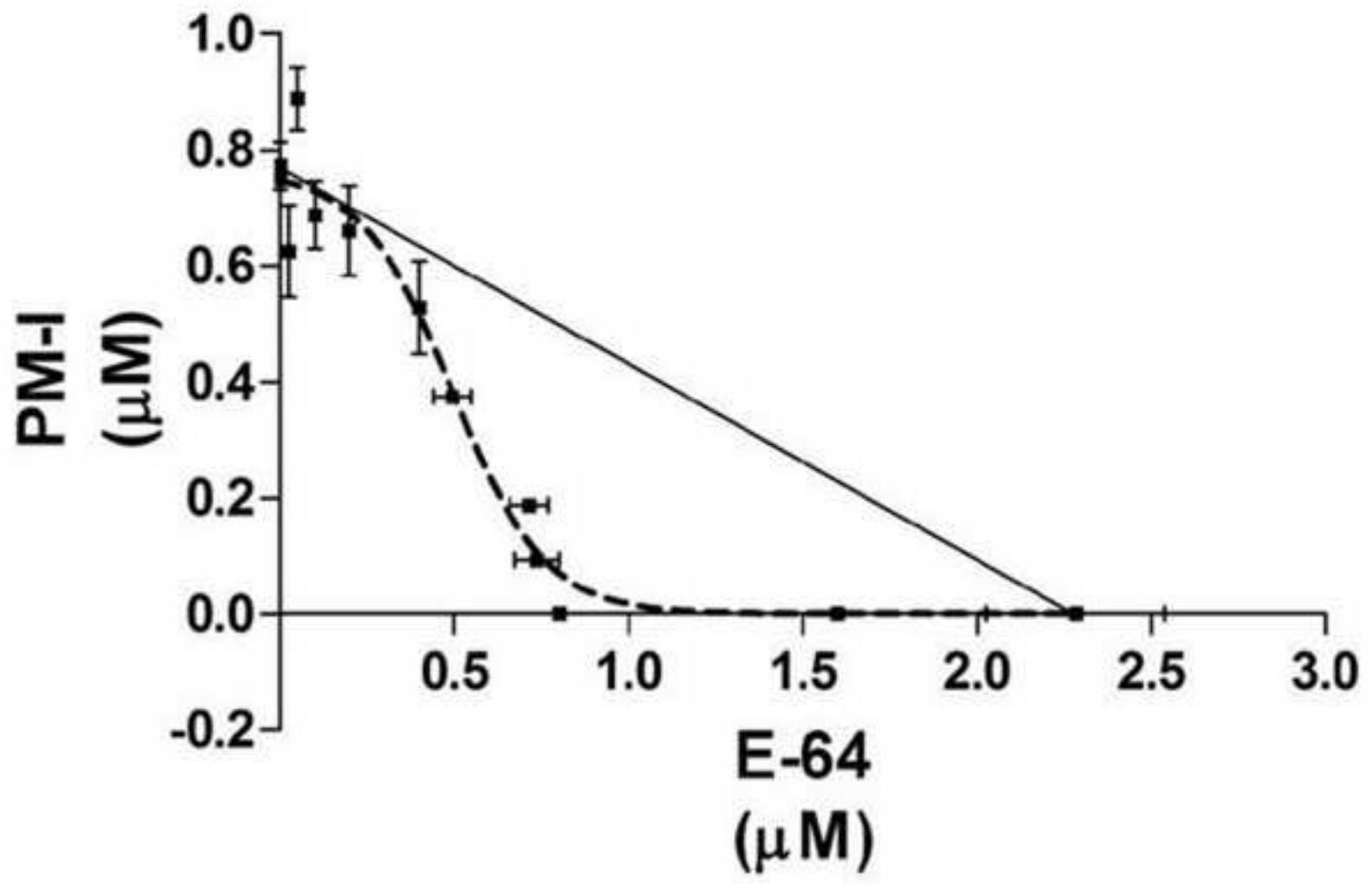


Fig. 4

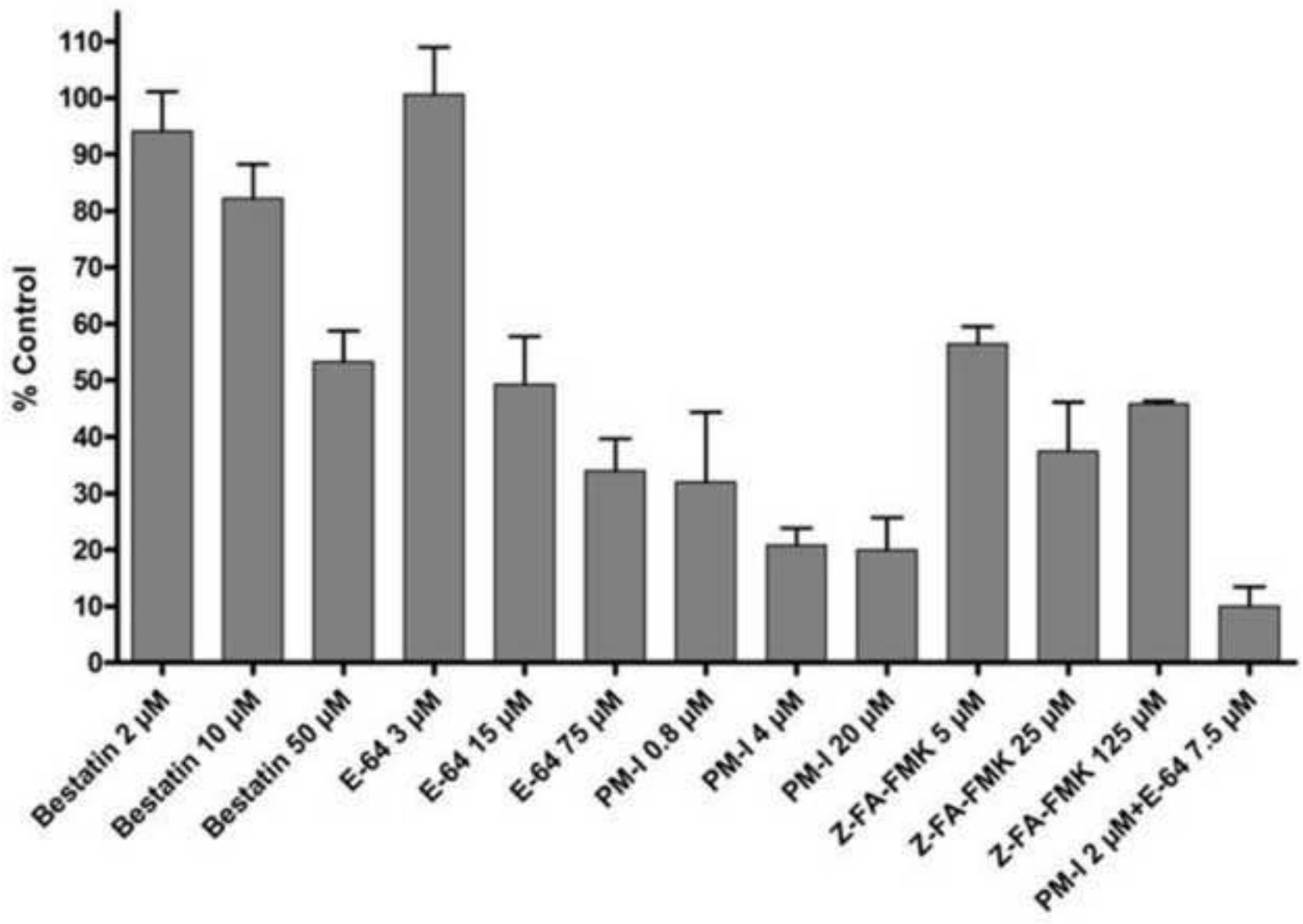


Figure 5

