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Title: Analysis of caecal mucosal inflammation and immune modulation during *Anoplocephala perfoliata* infection of horses.

Short title Immune modulation by *A.perfoliata* in horses.

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1 **Abstract**

2 **Aims**, *Anoplocephala perfoliata* is the commonest equine tapeworm, adult parasites are attached in
3 groups close to the ileo-caecal valve causing marked inflammatory pathology. This work aimed to
4 characterise the nature of the *in vivo* mucosal immune response to *A.perfoliata*, and to investigate
5 the role of *A.perfoliata* excretory secretory components in modulating *in vitro* immune responses.

6 **Methods and results**. Real time PCR detected elevation of IL13 and TGFβ transcription in early stage
7 *A.perfoliata* infection. In late stage infection IL-13, IL4 and Ifnγ transcripts were reduced while the
8 regulatory cytokines, TGFβ, IL10 and the transcription factor FOXP3 were increased in tissue close to
9 the site of *A.perfoliata* attachment; indicating down regulation of T-cell responses to *A.perfoliata*. *In*
10 *vitro*, *A.perfoliata* excretory secretory products induced apoptosis of the Jurkat T-cell line and
11 premature cell death of ConA stimulated equine peripheral blood leucocytes. Analysis of cytokine
12 transcription patterns in the leucocyte cultures showed a marked inhibition of IL-1 and IL-2
13 suggesting that a lack of T-cell growth factor transcription underlies the mechanism of the induced
14 equine T-cell death.

15 **Conclusion** These preliminary findings suggest *A.perfoliata* may have the ability to down-regulate
16 host T-cell responses.

17

18 **Introduction.**

19 Horses are the definitive host of the cestode *Anoplocephala perfoliata*, (Goeze, 1782) which
20 has a world-wide distribution¹, the reported prevalence varies from 6%-70%^{1,2,3,4}. Within the
21 horse caecum total and relative numbers of immature or adult developmental stages vary with
22 season and climate^{2,4,5,6,7,8}. The eggs are passed in faeces and develop into cystocercoids
23 within free living oribatid mites⁹ and grazing horses become infected through coincidental
24 ingestion of infected mites and subsequent *A.perfoliata* development takes place within the
25 gut lumen. Immature parasites are found superficially attached throughout the caecum¹, but
26 mature adult stages of *A.perfoliata* typically attach in groups close to the ileo-cecal
27 junction^{10,11,12}. In temperate climates, the seasonal variations in the observed numbers of
28 adult and immature *A.perfoliata* are consistent with an annual lifecycle, more immature
29 parasites are found during the late summer and autumn while fully mature adults predominate
30 during the late spring and early summer^{2,5,6,7}.

31 Historically, infection by equine cestodes has been regarded to be of little significance¹,
32 however more recent studies have linked *A.perfoliata* with colic including those caused by
33 ileal impaction, ileo-caecal intussusception and ulceration or perforation of the caecum in
34 heavily infected horses^{1,13,14,15,16,17,18,19}. Pathological changes seen in *A.perfoliata* infections
35 include epithelial and goblet cell hyperplasia, epithelial necrosis and ulceration, along with
36 thickening and eosinophil infiltration of the caecal lamina propria^{20,21,22}. Hyperplasia
37 resulting in thickening of the muscle layers, along with vascular and neural damage are also
38 described in heavy infections of more than 100 parasites²⁰.

39 The presence of such a severe localised inflammatory response to *A.perfoliata* occurs despite
40 the parasite remaining within the intestinal lumen, and it is unclear exactly what provokes the
41 host reaction. *A.perfoliata* E/S products may mediate pathological damage either by causing
42 direct injury and inflammation, or through stimulating an immune reaction which mediates
43 chronic inflammatory changes. Humoral immune responses to *A.perfoliata* excretory
44 secretory (E/S) products have been well documented and the presence of serum IgG(T)
45 antibodies to a major 10-12kD antigen has been used as a diagnostic test for tapeworm
46 infection^{23,24}. In a previous paper, we described active synthesis of both IgG(T) and IgE anti
47 *A.perfoliata* antibodies in explant cultures from caecal lamina propria of infected horses²². In
48 this paper we describe further the nature of the mucosal immune responses to *A.perfoliata*
49 and its E/S antigens.

50 **Material and methods**

51 **Caecal biopsies.**

52 Samples of caecal wall were collected from horses slaughtered at a licenced UK abattoir.
53 The autumn samples consisted of an 3-5cm cm² area of inflamed caecum close to the ileo-
54 caecal valve taken from horses with >100 *A.perfoliata* parasites (n=8), similar size control
55 samples were taken from horses with no *A.perfoliata* (n=8). The summer samples consisted
56 of control horses (n=8) and paired tissues from heavily infected horses (n=8), one taken from
57 the site of parasite attachment that showed gross pathological changes, and a sample from the
58 adjacent area of hyperplastic caecal wall 5-10 cm from the point of *A. perfoliata* attachment.
59 The samples were divided, one section was placed in RNeasyTM (www.thermofisher.com)
60 while a further section was pinned to dental wax and fixed in formalin for routine histological
61 processing.

62 ***A.perfoliata* E/S antigen preparation**

63 *A.perfoliata* collected from horses slaughtered at a licensed abattoir, were placed in a flask
64 containing caecal content for transportation to the laboratory. The *A.perfoliata* parasites were
65 washed in PBS then incubated at 37°C for six hours in serum free RPMI tissue culture media
66 without phenol red (www.Invitrogen.co.uk) containing 50 ug/ml gentamycin
67 (www.sigma.com). The medium was removed, spun at 3000g for 10 minutes, sterilised by
68 passing through a 0.2µm filter and the protein content was determined by fluorescent dye
69 binding assay (Qubit www.Invitrogen.co.uk). Endotoxin content of the preparations was
70 assayed using a *Limulus* Amoebocyte Lysate based assay (PierceTM LAL Chromogenic
71 Endotoxin Quantitation Kit (www.thermofisher.com). Aliquots of E/S were further
72 processed by one of the following methods a) heating at 56°C for 1hour, b) dialysis against
73 RPMI using a 3.5 kD membrane cut off Slide-A-LyzerTM cassette (www.thermofisher.com),
74 c) ultra-filtration using a 3 kD micro-centrifuge device (Pall Nanosep® www.sigma.com) e)
75 fractionation by reverse phase chromatography using a 1ml C-18 solid phase extraction tube
76 (Discovery DSC-18 www.Sigma-Aldridge.com/Supelco): briefly 5ml filtered E/S supernatant
77 was loaded onto a pre-wetted column, washed with PBS and eluted with a stepped
78 H₂O/methanol extraction buffer (1ml each 5%, 30% 70% 100% MeOH), the fractions were
79 freeze dried overnight then re-dissolved in serum free RPMI tissue culture medium. The size
80 and number of proteins were measured by NuPAGE gel electrophoresis using 4-12% Bis-Tris
81 gradient gels and MES buffer (www.Invitrogen.co.uk) stained with Coomassie blue.

82 **Effect of *A.perfoliata* E/S products on in vitro proliferation and viability of Jurkat cells.**

83 Jurkat J6 cells (www.ATCC.org) were maintained in RPMI supplemented with 2mM
84 glutamine and 10% FCS; cells were seeded at 10^5 /ml and split 1:10 with fresh media every 3-
85 4 days. For the growth inhibition assay 100ul aliquots of cells in log phase at 10^5 /ml added to
86 a 96 well tissue culture plate. Triplicate wells were treated with dilutions of E/S preparations
87 and incubated at 37°C for 72 hours, cell growth/viability was assayed using the Vybrant®
88 MTT assay system according to the manufacturer's instructions (www.thermofisher.com).

89 **Lamina propria cytokine quantitative RT-PCR**

90 The muscular layers of the caecal wall were stripped off, total RNA was isolated from 30mg
91 of the remaining lamina propria/epithelial tissue using Nucleospin RNA kit (Macherey-Nagel
92 www.mn-net.com) incorporating a three stage DNase treatment protocol to ensure complete
93 removal of genomic DNA as previously described²². The total RNA concentration of each
94 sample was assayed using a fluorescent dye binding assay (Quibit RNA BR
95 www.invitrogen.com). Complementary DNA was generated from 50ng of the total RNA
96 using ImProm-II™ Reverse Transcriptase (www.promega.co.uk) and random hexamer
97 primers. The cDNA was made up to a final volume of 120ul. Q-PCR reactions consisting of
98 5ul aliquots of cDNA, 12.5µl Gotaq mastermix (www.promega.co.uk), 1.25µl 50mM
99 MgCl₂, 0.5µl of 10µM forward and reverse primers plus 0.5µl of the appropriate 10µM probe
100 conjugated to 3'FAM and 5' BQ1 (www.metabion.com) were run on a Mx3005P
101 (www.genomics.agilent.com). The cycling parameters were, 95°C for 2 minutes, followed
102 by 40 cycles of 95°C for 15 seconds and 60°C for 30 seconds. The primer pairs and gene
103 specific probes for equine multiple housekeeper genes (RPL, B2M, ACTβ, GAPDH, UBB,
104 TUBA1, HPRT) along with equine cytokine gene specific primer probe sets (IL1β, IL2, IL4,
105 IL5, IL6, IL10, IL13, IL17, TGFβ, Ifnγ and FOXP3) were as previously described²⁵. For IL9
106 a new set of primers and probes were designed using primer_3 software
107 (<http://frodo.wi.mit.edu/primer3>); these were EqIL9f ctacaggagcaccaccttc, EqIL9r
108 aaggaatgggcagacacaa and EqIL9probe cggtcacattggtgctgcagc. The efficiency of this primer
109 pair was 100.1%, r = 0.998.

110 The target gene transcripts were normalised against multiple house keeper genes using
111 GeNorm software²⁶. The normalised data for cytokine transcripts are expressed as relative
112 copy number calculated by $(1 + \text{Efficiency})^{(40 - \text{CT})}$.

113

114 **Effect of *A.perfoliata* E/S products on transcription of key cytokines by *in vitro***
115 **proliferation of horse lymphocytes activated by Concanavalin-A.**

116 At a licensed abattoir, 50 ml of blood was collected immediately post stunning into 15ml acid
117 citrate dextrose anticoagulant (ACD_B USP), the absence of *A.perfoliata* infection was
118 established by examining the caecum for parasites. Peripheral blood mononuclear cells
119 (PBMCs) were separated by centrifugation through Lymphosep Lymphocyte Separation
120 Medium (www.thermofisher.com) harvested and washed in RPMI. The immune status of the
121 horses to *A.perfoliata* was determined by IgG(T) ELISA; only lymphocyte samples from
122 sero-negative horses (n=7) were used for the final analysis.

123 Aliquots of PBMC from each horse were cultured in 5ml RPMI medium containing 10% FCS
124 100U/ml penicillin/100ug/ml streptomycin at a cell density of 2×10^6 PBMC/ ml. Three wells
125 were prepared from each horse; one contained media alone, the second was stimulated with
126 10ug/ml Con-A, the third contained 10ug /ml Con-A along with *A.perfoliata* E/S supernatant
127 at a final concentration of 1:20. The PBMCs were harvested after 24 hours and mRNA
128 isolated using mTRAP™ Midi Kit (www.activemotif.com) following the manufacturers
129 protocol incorporating the optional DNase step. Complementary DNA was immediately
130 synthesised from the eluted mRNA using ImpromII reverse transcriptase
131 (www.promega.com) and random hexamer primers. The obtained cDNA was run in a
132 QRTPCR reaction using the same equine cytokine primer pairs and probes as described for
133 the lamina propria samples.

134 **Statistical analysis.**

135 Statistical analysis on the two groups of lamina propria cytokine transcripts from infected and
136 uninfected horses taken in November was carried out using Mann-Whitney test $\alpha=0.05$. For
137 the three groups of samples taken in July, Kruskal-Wallace ANOVA $\alpha=0.05$ was used to
138 determine overall significance of differences between the groups, if significant differences
139 were detected, Mann-Whitney test was then used to examine differences between the pairs of
140 groups with a Bonferroni corrected $\alpha=0.16$ for three possible comparisons. For the
141 peripheral blood cytokine results Friedman's two-way analysis of variance by rank was used
142 to determine if there were overall significant differences between treatments ($\alpha=0.05$). Post
143 hoc differences between treatments were examined using Wilcoxin signed-rank test and one-

144 sided hypothesis ConA> Media; ConA+E.S< ConA; ConA+E/S>media; with a Bonferroni
145 correction for three comparisons $\alpha=0.033$.

146

147 **Results**

148 **Histopathology of caecal wall**

149 Fig 1a shows the caecum of a horse during autumn in which there is a moderate level of
150 *A.perfoiata* infection, the parasites are variable in size and stage of maturity. Individual
151 parasites were attached in a diffuse pattern, near the ileo-caecal valve (arrow). When the
152 parasites were removed the mucosa at the site of attachment was seen to be inflamed but the
153 wall of the caecum in the surrounding area remained grossly normal in thickness with clearly
154 defined rugae. By contrast Fig 1b shows a horse during the summer, the mature parasites are
155 attached in a discrete cluster close to the ileo-caecal valve, the mucous membrane at the site
156 of attachment was grossly thickened, reddened and ulcerated; upon cutting the tissue showed
157 extensive fibrosis. In heavy infections (>100 parasites), the mucosa and caecal wall
158 surrounding the site of attachment had gross thickening and oedema with fewer, less defined
159 rugae compared to the more distant areas of unaffected caecum.

160 Histological examination confirmed the findings. Biopsies from eight control horses and
161 twenty four samples from sixteen affected horses were analysed. The healthy control
162 samples were almost identical to one another, the samples from *A.perfoliata* lesions were
163 more variable but all showed leucocyte infiltration and hyperplasia which increased in
164 biopsies associated with heavy parasite burden and those closest to the point of parasite
165 attachment. Figure 1c is taken from the caecal mucosa of a horse in which no *A.perfoliata*
166 were found, a section through the entire thickness of the caecal wall. Figure 1d depicts an
167 inflamed area of caecum close to the site of attachment of *A.perfoliata* taken during the early
168 autumn phase of infection. The caecal wall shows some hyperplasia of the epithelium, and
169 increased numbers of infiltrating eosinophils and lymphocytes were evident in the lamina
170 propria and sub mucosa. Areas of haemorrhage, oedema, disruption and necrosis of the
171 muscular layers of the intestine were also present.

172 Figure 1e depicts part of a section through the caecal mucosa from the point of attachment of
173 adult *A.perfoliata* parasites during the late summer phase of the infection. There is marked
174 hyperplasia of the epithelium, with increased numbers of goblet cells and necrosis of the

175 superficial epithelial layers, due to hyperplasia of all tissue layers, only the mucosa and
176 submucosa fit within the frame. Extensive infiltration of the sub-mucosa by eosinophils and
177 lymphoid cells was evident, the deeper layers consisted of disrupted muscle tissue with
178 extensive leucocyte infiltration, and fibrosis (not shown). Figure 1f is from a section of
179 caecal wall in the area adjacent to the site of *A.perfoliata* attachment where there is less
180 severe hyperplasia of the mucous membrane, with infiltrating eosinophils, leucocytes and
181 lymphoid follicles also frequently observed in the sub-mucosa. The muscularis is markedly
182 hyperplastic becoming almost twice the thickness of that seen in an uninfected horse Fig 1c.

183 **Cytokine gene expression in caecal mucosa.**

184 Cytokine gene expression was assayed in caecal mucosa collected during November from
185 nine *A.perfoliata* infected horses harbouring over 100 *A.perfoliata* parasites in varying stages
186 of maturity, and eight control horses in which no *A.perfoliata* were observed Fig 2. The
187 results were analysed by Mann Whitney test; *A.perfoliata* infected horses had a significant
188 increase in IL13 ($p=0.034$) and TGF β ($p=0.021$) transcripts compared to the uninfected
189 control horses, but only a trend towards increased IL4 ($p=0.073$) was detected in infected
190 mucosa and no other significant changes were observed for the other cytokine transcripts
191 ($p>0.10$).

192 The July samples from late stage infected horses had fewer IL4 transcripts at the site of
193 attachment and in the adjacent areas compared to the control ($p= 0.002$). IL13 transcription
194 showed a trend toward the same result ($p=0.052$). Transcription of Ifn γ was also significantly
195 reduced in the caecal tissues from infected horses compared to controls ($p=0.0085$) indicating
196 a general reduction in both Th1 and Th2 effector T-cell function. Regulatory T-cell
197 cytokines showed significant variation among the groups (Kruskal-Wallis $p<0.05$). IL10
198 was significantly ($p<0.01$) lower in samples taken from the site of parasite attachment
199 compared to samples taken from uninfected mucosa while samples from infected mucosa
200 adjacent to the site of attachment were highly significantly higher ($p<0.01$) than at the site of
201 attachment and also showed a trend ($p=0.052$) towards higher IL10 compared to the
202 uninfected horses. In the infected horses TGF β was significantly higher ($p<0.001$) in the
203 areas of mucosa adjacent to site of attachment, compared to both the site of attachment and
204 the uninfected controls. FOXP3 expression differed between the groups; post hoc Mann-
205 Whitney testing revealed the largest effect to be a significant increase ($p=0.021$) in the tissues
206 adjacent to the sites of *A.perfoliata* attachment compared to uninfected tissue from control

207 horses. Both IL1 β and IL6 were significantly elevated in the inflamed tissues at the site of
208 attachment compared to either the control tissue or the tissue adjacent to the site of
209 attachment ($p < 0.01$), once again IL9 did not differ between the groups.

210 **Effect of *A.perfoliata* ES on growth of Jurkat cells.**

211 *A.perfoliata* were incubated in serum-free RPMI subsequent analysis of the E/S by SDS page
212 and Coomassie staining revealed the presence of the previously described dominant 10-12kD
213 band and at least eighteen other protein bands with molecular weights between 20 and >250
214 kD Fig 4a. Preliminary experiments showed that this ES supernatant inhibited the growth of
215 the Human T-cell line Jurkat when added to the culture media (Fig 4b). To further
216 characterise and or neutralise the inhibitory activity a series of experiments were conducted
217 on the ES supernatant using Jurkat cells. Neither heat inactivation nor dialysis against a 3.5
218 kD cut off membrane completely remove the activity Fig 4c although in both cases protein
219 precipitation occurred resulting in a less intense set of bands seen in SDS page Fig 4a.
220 Ultrafiltration of the ES supernatant through a 3kD cut off membrane removed all visible
221 proteins Fig 4a but did not completely remove the inhibitory activity Fig 4c. The retentate
222 fractions > 3 kD cut-off membrane also had inhibitory activity suggesting a larger active
223 component (data not shown). Taken together the results are consistent with a small active
224 component that is bound in equilibrium with a larger carrier protein.

225 The LAL test showed that the E/S supernatants typically contained endotoxin at
226 concentrations of between 1 and 4 ug/ml. To address the question of whether endotoxin was
227 the cause of the growth inhibition, the effect of a sample of *A.perfoliata* E/S supernatant with
228 an LPS content of 1-4 ug/ml was compared to either culture media containing 100ug/ml LPS
229 (equivalent to 10 ug/ml at the 1:10 starting dilution) or as a more general control for
230 contamination by gut microflora a sample of supernatant generated by incubating culture
231 media with equine mucosal tissue in place of the *A.perfoliata*. The results shown in figure 4d
232 confirmed that only the *A.perfoliata* E/S supernatant caused significant inhibition of Jurkat
233 cells. Passing the ES supernatant through a C18 reverse phase absorbent cartridge, did not
234 remove all the inhibitory activity, nevertheless, inhibitory activity had bound to the C18
235 column as evidenced by the inhibition of Jurkat cell growth induced by the eluted fractions.
236 The maximal activity was recovered in the 75% MeOH elution fraction (after freeze drying
237 and re-dissolving in RPMI). Figure 4e.

238 Fig 5 shows the kinetics of E/S induced cell death in Jurkat cultures using flow cytometry to
239 assay annexin binding and 7-AAD to assay cell permeability. Throughout the 72 hours of
240 culture there was a gradual increase in the number of cells binding annexin and staining with
241 7-AAD indicating loss of cell membrane integrity. This pattern is consistent with a gradual
242 loss of cell viability due to an indirect apoptotic mechanism rather than an acute chemically
243 induced necrosis of the cells.

244 **Effect of *A.perfoliata* ES on cytokine transcription by ConA stimulated lymphocytes.**

245 After 24 hours, both ConA and ConA plus 5% *A.perfoliata* E/S cultures demonstrated the
246 initial clumping phase of growth. However, in the presence of *A.perfoliata* E/S the
247 lymphocyte clumps disaggregated from 24-48 hours and all the cells died prematurely
248 between 48-72 hours.

249 Messenger RNA was harvested from Con-A lymphocyte blasts after 24 hours of culture when
250 the cells were still viable. Fig 6 depicts the relative copy number of cytokine gene transcripts
251 normalised against a panel of housekeeping genes. As expected ConA stimulated a
252 significant ($p<0.025$) increase in transcription of the cytokines IL2, IL4, IL5, IL13, $\text{Ifn}\gamma$, and
253 IL17 compared to media alone. In the presence of ES supernatant, the ConA stimulated
254 transcription of IL2, IL5, IL17 and $\text{Ifn}\gamma$ was significantly reduced ($p<0.025$) compared to
255 ConA stimulated lymphocytes while IL4 and IL13 remained unchanged ($p>0.05$). No
256 overall significance was detected for $\text{TGF}\beta$, IL10 or FOXP3 (Friedman's $p>0.05$). IL1 was
257 markedly reduced following ConA stimulation $p<0.01$ and in the presence of *A.perfoliata* E/S
258 there was a further significant decrease in IL1 transcription $p=0.018$. In contrast there was no
259 effect of any treatment on IL6 (Friedman's $p>0.05$). The results for ActB are shown to
260 confirm that all samples contained cDNA.

261 **Discussion**

262 The pathological changes associated with *A.perfoliata* infection described here agree with
263 previously published reports in which more severe lesions were associated with higher
264 parasite burdens^{20,21}. Based on our previous findings of *A.perfoliata* specific IgE and IgG(T)
265 antibody synthesis within the caecal lamina-propria of infected horses²², we had expected to
266 find elevations of Th2 type cytokines, at least in the acute stage of infection. The results
267 rather confounded this expectation with a modest increase in IL13 and increased $\text{TGF}\beta$ the
268 only significant changes detected. Neither IL4 nor IL5 were significantly elevated and IL9, a

269 cytokine associated with several aspects of innate and adaptive anti-parasite or allergic
270 immune responses^{26,28,29}, was absent or at very low copy number in the majority of infected
271 horses (Figs 2,3). Differentiation of IL9 producing Th9 cells requires both IL4 and TGF β ,
272 whereas the relatively low IL4 and high TGF β seen in the early *A.perfoliata* infection (Fig 2)
273 would be expected to favour T-reg development³⁰. Th9 cells are not the only potential source
274 of IL9, Fc ϵ RI bearing mucosal mast cells can also release significant amounts of IL9, this
275 source has been shown to have a critical role in food allergy²⁸ and intestinal helminth
276 rejection by rodents³¹. Abundant Fc ϵ RI mucosal mast cells are present in the lamina propria
277 of equine colon and their numbers increase with maturity or in association with high burdens
278 of cyathostomes^{32,33}. Although Fc ϵ RI cells are also numerous throughout the caecal lamina-
279 propri of *A.perfoliata* infected horses²², their production of IL9 is not supported by our data,
280 and their importance in immunity to *A.perfoliata* remains unknown.

281 Immune regulation is a well described component of anti-parasite responses to Taeneiids, in
282 respect of the parasite cystosercoid stages within the tissues of their intermediate hosts³⁴ and
283 by adult *Hymenolepis diminuta* (Rudolphi 1819) in the intestine of its definitive host³⁵. The
284 very modest changes in Th2 cytokines and the high levels of TGF β in autumn *A.perfoliata*
285 infections suggests that even at this early stage post infection the immune response to
286 *A.perfoliata* is down-regulated. This impression was confirmed by the results from late stage
287 infections in July where the Th2 cytokines IL13 and IL4 along with The Th1 cytokine
288 Ifn γ were all markedly reduced in infected horse lamina propria compared to controls.
289 Moreover, the regulatory cytokines IL10 and TGF β , as well as the transcription factor
290 FOXP3 were all elevated in the tissues adjacent to the point of parasite attachment.
291 Conversely, at the point of attachment itself, where extensive cellular infiltration and damage
292 were observed, there was a reduction in IL10 and an increase in pro-inflammatory cytokines
293 IL1 and IL6 indicating an active inflammatory response mediated by leukocytes responding
294 to *A.perfoliata* and/or environmental antigens gaining access due to loss of intestinal barrier-
295 function.

296 Immune modulation is a common feature of nematode pathogenesis in which a wide range of
297 excretory secretory products have been identified³⁶. Rodent models of immune regulation by
298 cestodes implicated E/S components as the mediators of immune suppression, both by larval
299 stages of *Mesocostoides vogae*³⁷ and by adult *H.diminuta*³⁵. The inhibition of Jurkat T-cells

300 and the down regulation of cytokine transcription by ConA stimulated equine lymphocytes by
301 *A.perfoliata* E/S components provides another example of this type of activity.

302 The effect of the E/S in supressing Th1 cytokines, in particular IL2, is reminiscent of several
303 immunosuppressive drugs; e.g cyclosporine; or FK506 which act via calcineurin binding to
304 inhibit the dephosphorylation and nuclear localisation of NFAT (nuclear factor activated T-
305 cell) which in turn prevents IL2 transcription³⁹. While we have not yet identified the active
306 component of *A.perfoliata* E/S components or its mechanism of action, the results so far
307 favour a small MW compound or possibly a peptide and further studies to identify the active
308 component would be merited.

309

310 **Ethical approval** This article does not contain any studies with animals performed by any of the
311 authors. The use of post mortem materials taken from animals was approved by the University of
312 Bristol animal welfare committee authorisation number UIN/18/045.

313
314 **Data availability** The data that support the findings of this study are available from the
315 corresponding author upon reasonable request.

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410

411 **Figure legends**

412 **Fig 1** a) shows the caecum of a horse during autumn in which there is a moderate level of
413 *A.perfoliata* infection, the parasites are variable in size and stage of maturity, attached in a
414 diffuse pattern, close to the ileo-caecal valve (arrow), b) *A.perfoliata* infected horse caecum
415 during the summer, similar sized adult *A.perfoliata* attached in a discrete cluster close to the
416 ileo-caecal valve, c) histological section showing the entire thickness of the caecal mucosa
417 from a horse in which no *A.perfoliata* were present, x40 magnification, d) caecal wall from
418 an inflamed area close to the site of *A.perfoliata* attachment taken during the early phase of
419 infection x40 magnification, e) histological section through the caecal mucosa from the point
420 of attachment of adult *A.perfoliata* parasites during the late summer phase of the infection x
421 40 magnification. Due to the hyperplasia of all tissue layers, only the mucosa and submucosa
422 fit within the frame, f) a section of caecal wall from a late stage infection taken from the area
423 adjacent to the site of *A.perfoliata* attachment x40 magnification.

424

425 **FIG 2** Relative copy number of cytokine transcripts in the lamina propria during November
426 the early stage of *A.perfoliata* life cycle. Uninfected control horses (solid symbols n=9) and
427 *A.perfoliata* infected (open symbols n=8), a) TH2 cytokines; IL13 (circles) was significantly
428 elevated ($p=0.034$) in infected horses, IL4 (diamonds) was not significantly different from
429 control $p=0.073$, IL5 (triangles) was not significantly different from control ($p>0.1$), b) TH1
430 cytokines IL2 (circles) and $\text{Ifn}\gamma$ (diamonds) did not differ between control and infected
431 horses, nor was there any difference in IL17 (triangles), c) shows no significant difference in
432 the regulatory cytokine IL10 (circles) nor in the regulatory transcription factor FOXP3
433 (triangles) but there was a significant ($p=0.021$) increase in $\text{TGF}\beta$ (diamonds) in the mucosa
434 of *A.perfoliata* infected horses, d) shows the results for pro-inflammatory genes IL1 β
435 (circles) and IL6 (diamonds), along with IL9 (triangles) none of which showed any
436 significant change ($p>0.1$) between infected and uninfected horses

437 **Fig 3** Relative copy number of cytokine transcripts in the lamina propria in July the later
438 stage of *A.perfoliata* infection cycle. Samples taken from uninfected control horses (solid
439 symbols n=8), from the site of *A.perfoliata* attachment (open symbols n=8), and from the
440 thickened mucosa adjacent to the site of *A.perfoliata* attachment (shaded symbols n=8), a)
441 TH2 cytokines; IL13 (circles) was not significantly different from control ($p=0.052$), both at
442 the site of *A.perfoliata* attachment and in the adjacent lamina propria. IL4 (diamonds) was
443 significantly reduced in infected horses compared to controls $p=0.02$. IL5 (triangles) was not
444 significantly different from control ($p>0.1$). b) TH1 cytokine $\text{Ifn}\gamma$ (diamonds) was
445 significantly reduced in infected horses compared to control ($p=0.0085$) but there was no
446 difference in IL2 (circles) or IL17 (triangles) between the groups, c) IL10 (circles) was
447 significantly reduced ($p<0.001$) at the site of attachment ** compared either to the areas
448 adjacent to parasite attachment or to uninfected horses. IL10 was not significantly different
449 in the mucosa adjacent to the site of attachment compared to uninfected controls ($p=0.059$).
450 The regulatory transcription factor FOXP3 (triangles) was a significantly higher ($p<0.05$) in
451 the adjacent mucosa compared to the site of attachment or to normal horses. Similarly $\text{TGF}\beta$
452 (diamonds) was highly significantly increased ($p<0.001$) in the mucosa adjacent to the site of
453 *A.perfoliata* attachment, d) pro-inflammatory genes IL1 β (circles) and IL6 (diamonds)
454 showed a highly significant rise at the site of *A.perfoliata* attachment ($p<0.001$) compared to

455 either the tissue adjacent to the site of attachment or to uninfected control horses, IL9
456 (triangles)) did now show any significant change ($p>0.1$) between any group

457 **Fig 4** a) *A.perfoliata* E/S separated on an 4-12% SDS gel. Lane 1 molecular weight
458 standards, lane 2 *A.perfoliata* E/S, lane 3 *A.perfoliata* E/S after heat inactivation and removal
459 of precipitated proteins, *A.perfoliata* E/S <3kD filtrate, *A.perfoliata* E/S after dialysis against
460 3kD membrane and removal of precipitate. 4b) Inhibition of Jurkat cell growth in the
461 presence of E/S supernatant. Each bar represents mean \pm standard error of n=4 experiments
462 using different samples of *A.perfoliata* ES. Each sample dilution was cultured in triplicate
463 wells and the growth is expressed as a % of media control. 4c) Inhibition of Jurkat cells by
464 *A.perfoliata* E/S components after ultrafiltration through a 3kD membrane, heat inactivation
465 at 60°C for thirty minutes or dialysis using a 2kD cut of membrane. Each bar represents the
466 mean \pm standard error of n=4 experiments . Each sample dilution was cultured in triplicate
467 wells and the growth is expressed as a % of media control. 4d) Media containing 100 ug/ml
468 LPS had no inhibitory effect on Jurkat cell growth at a dilution of 1:10 (equivalent to 10ug/ml
469 final concentration), compared to a samples of undiluted *A.perfoliata* supernatant which had
470 1-4 ug/ml LPS contamination. Samples of culture supernatant prepared using equine caecal
471 mucosa in place of *A.perfoliata* were similarly devoid on inhibitory activity. Each bar
472 represents the mean \pm standard error of n=3 experiments each sample dilution was cultured in
473 triplicate wells and the growth is expressed as a % of media control. 4e) Inhibition of Jurkat
474 cell growth by E/S/ components eluted from a C18 column with increasing concentrations of
475 methanol. The fractions were freeze dried and dissolved in RPMI before testing. Each bar
476 represents the mean \pm standard error of n= 5 experiments. Each sample dilution was cultured
477 in triplicate wells and the growth is expressed as a % of media control.

478 **Fig 5 Data from a representative experiment showing changes indicating apoptosis of**
479 **Junkat cells following treatment with A.perfoliata E.S.** Percentage apoptotic cells in
480 cultured Jurkat cells over a 72-hour time course assayed by annexin and 7AAD binding using
481 flow cytometry. Panels a,c,e cultured in media, panels b,d,f, cultured in media with 5%
482 *A.perfoliata* E/S supernatant

483 **Fig 6** Relative copy number of cytokine transcripts in cDNA from samples of equine
484 peripheral blood lymphocytes (n=7) cultured in media alone (solid symbols), in the presence
485 of 5ug Con A (shaded symbols), or in the presence of 5ug ConA with 5% *A.perfoliata* E/S
486 supernatant (open symbols). Significant differences ($p<0.05$) between groups are indicated by
487 *

488

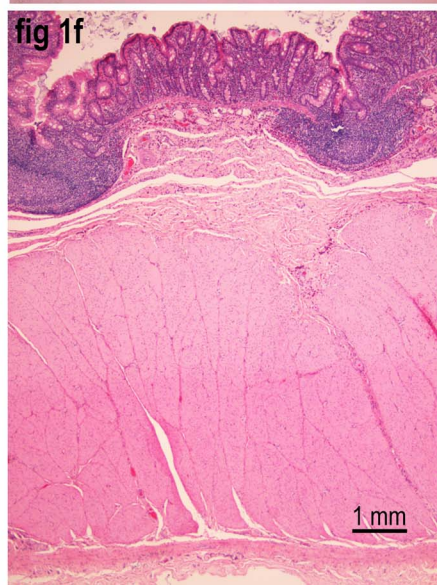
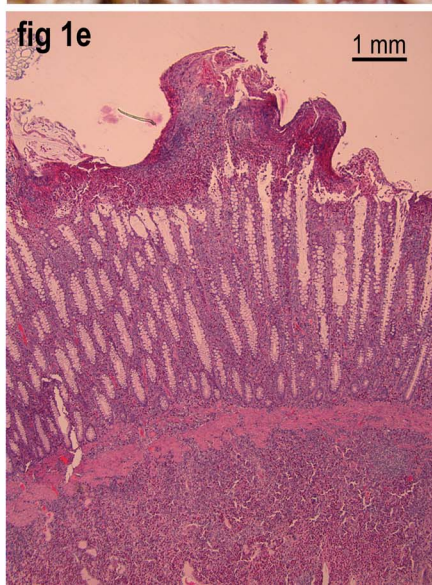
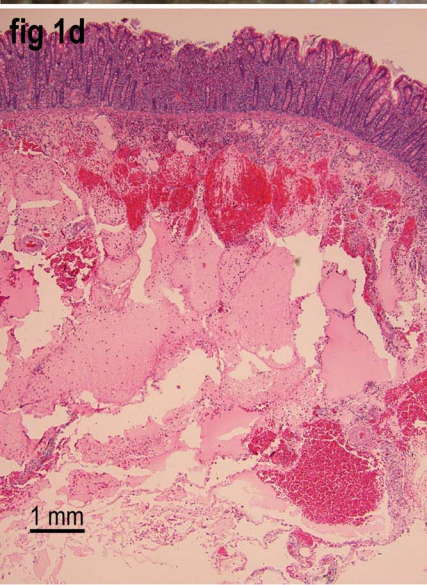
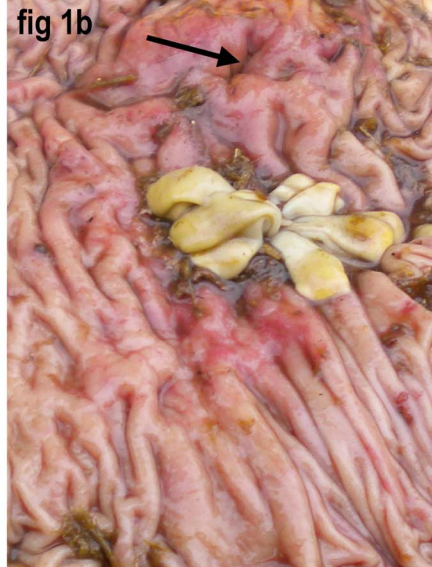
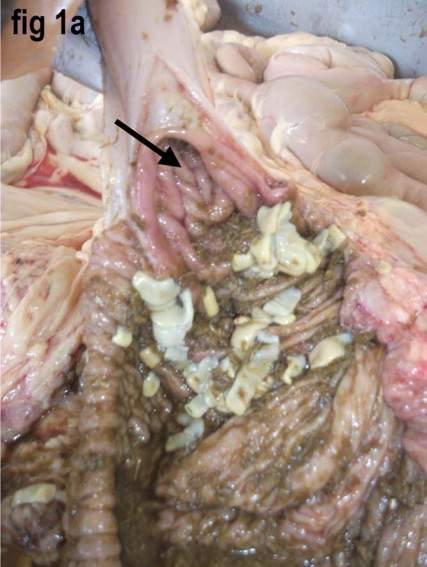


Fig 2a

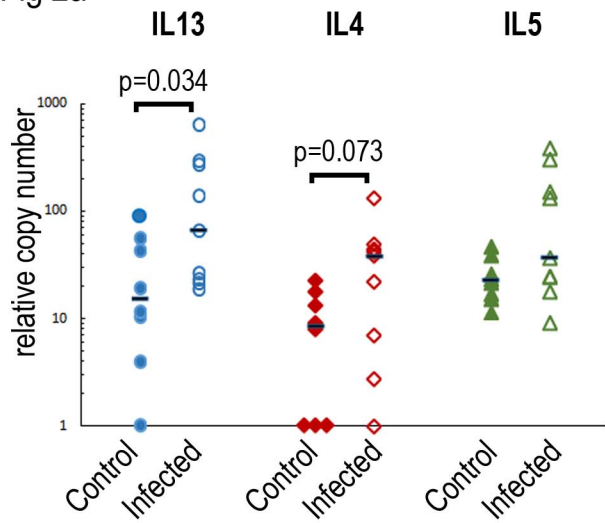


Fig 2b

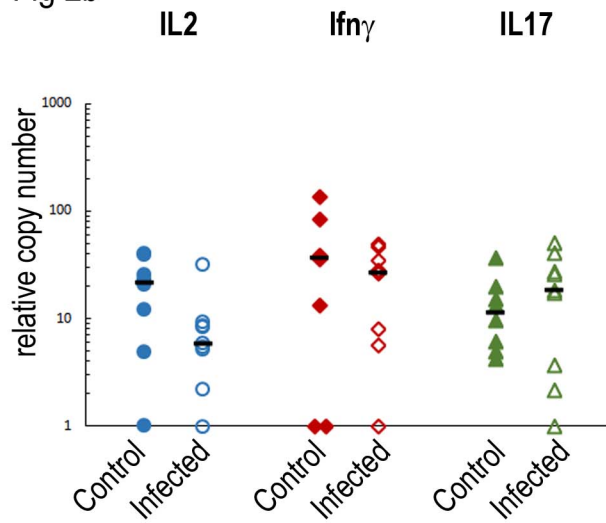


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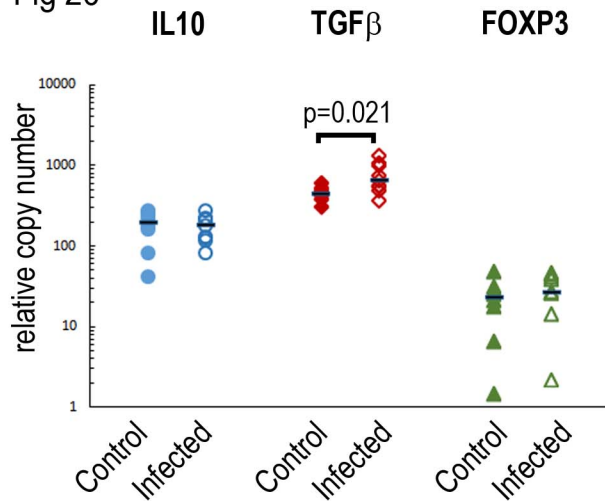


Fig 2d

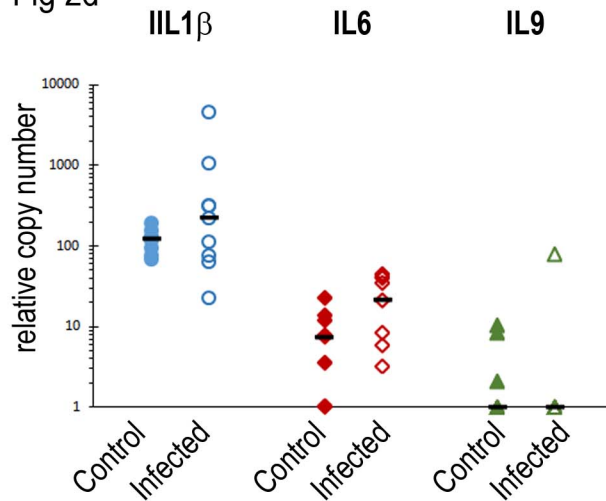


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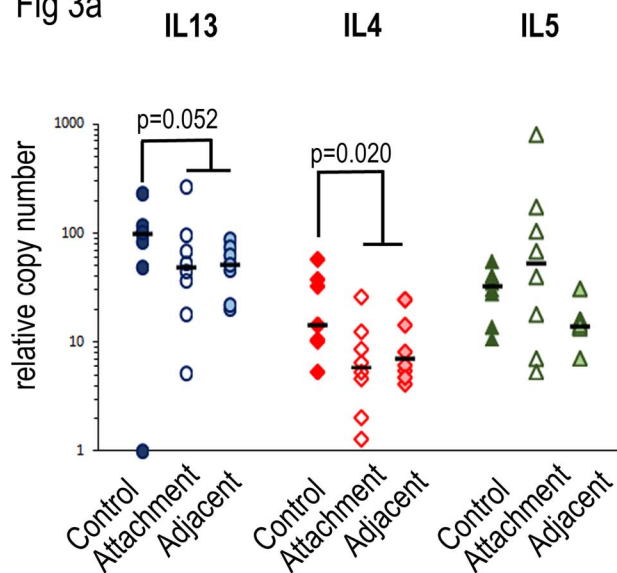


Fig 3b

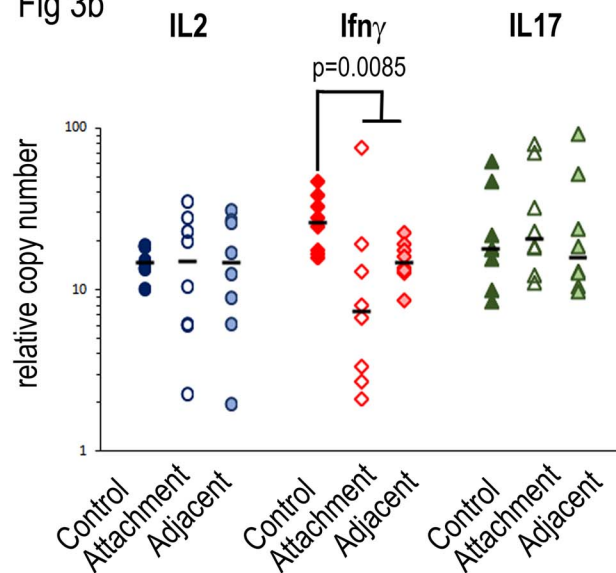


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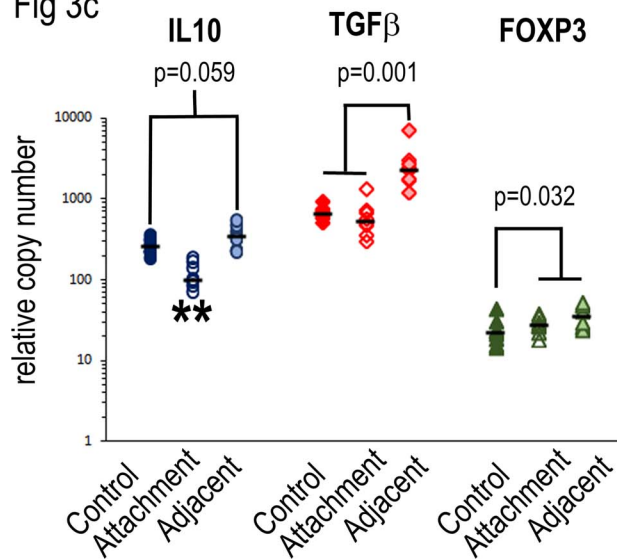


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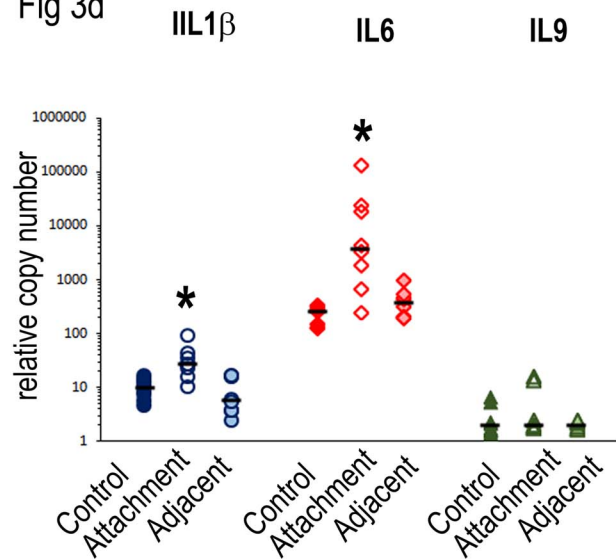


fig 4a

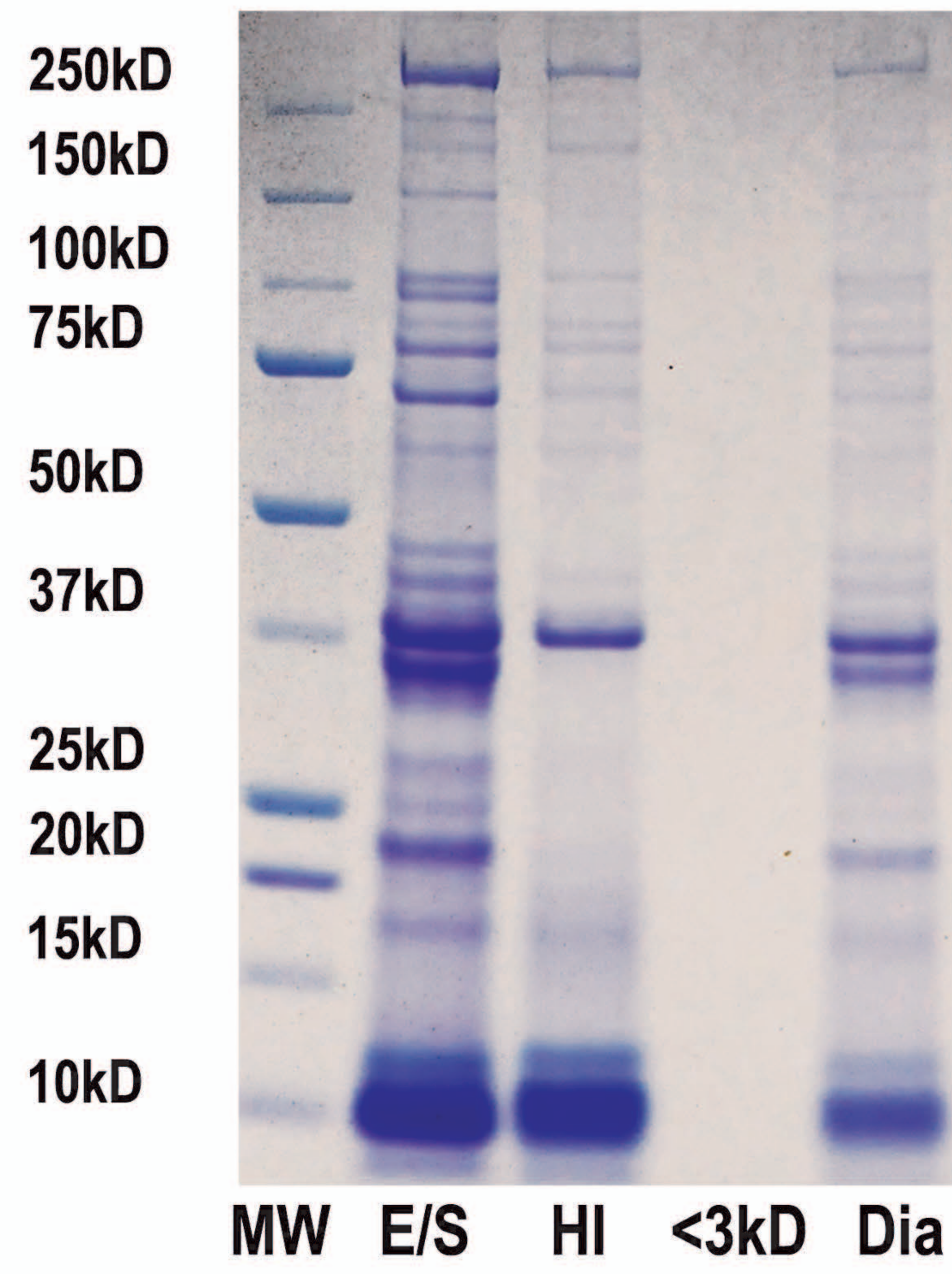


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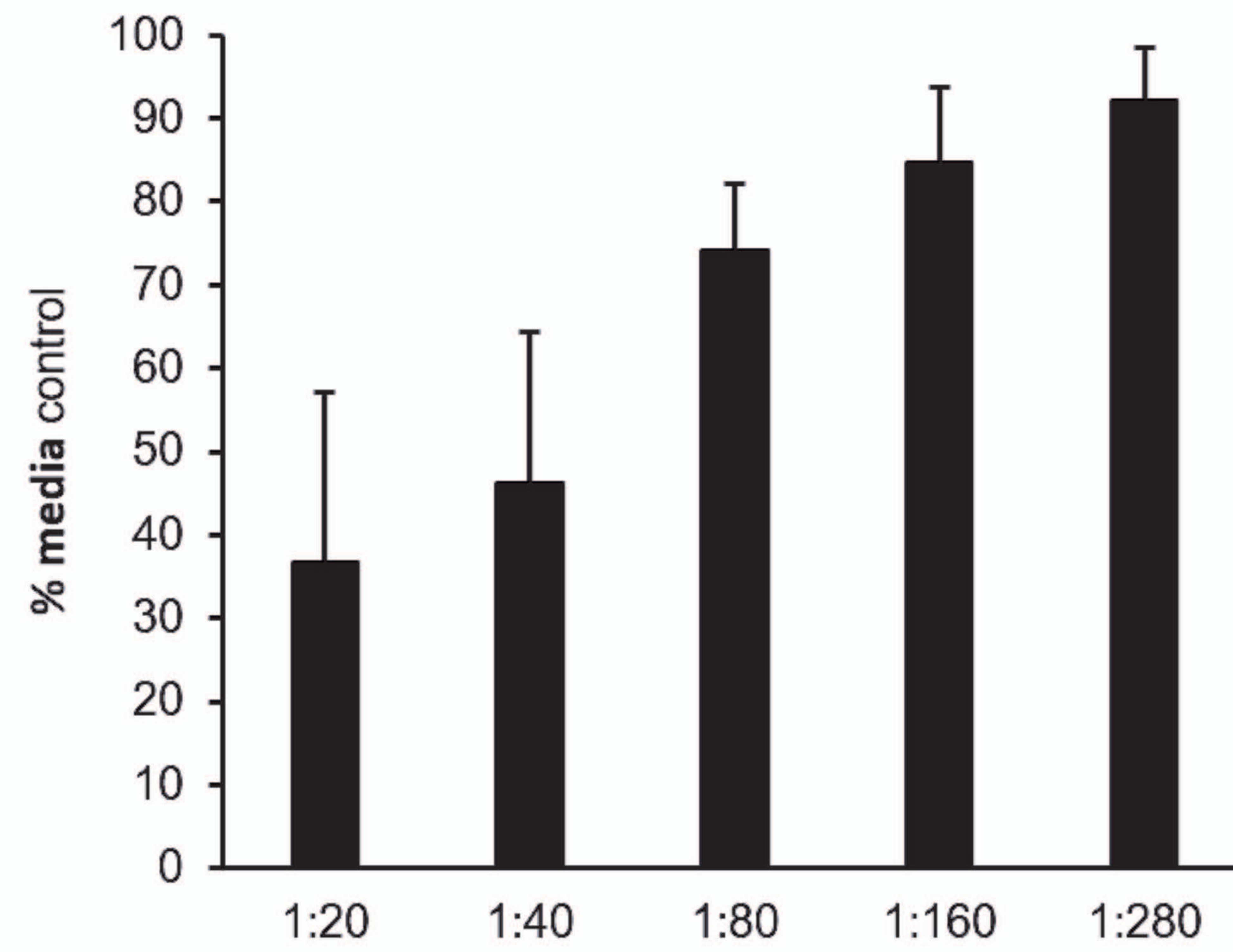


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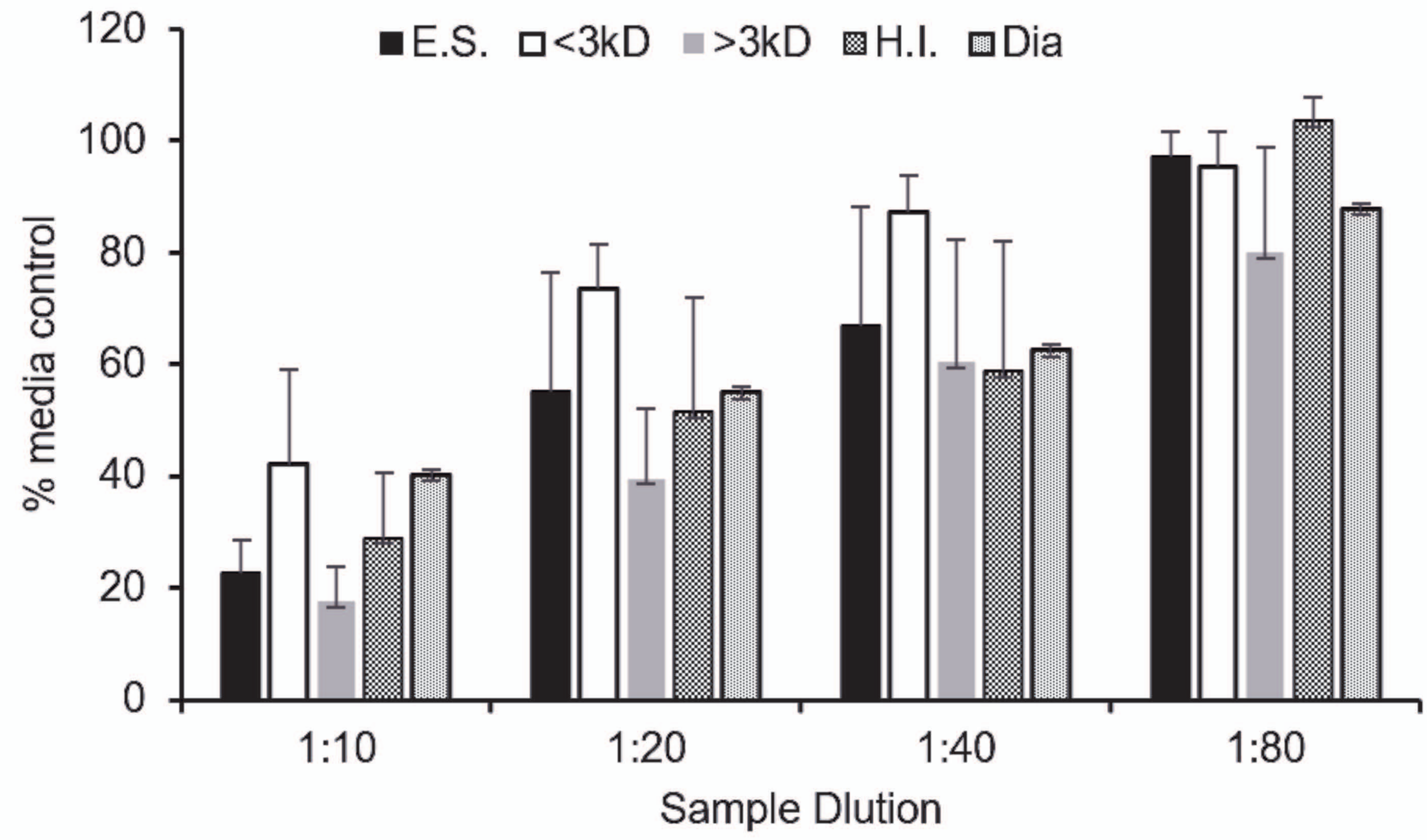


fig 4d

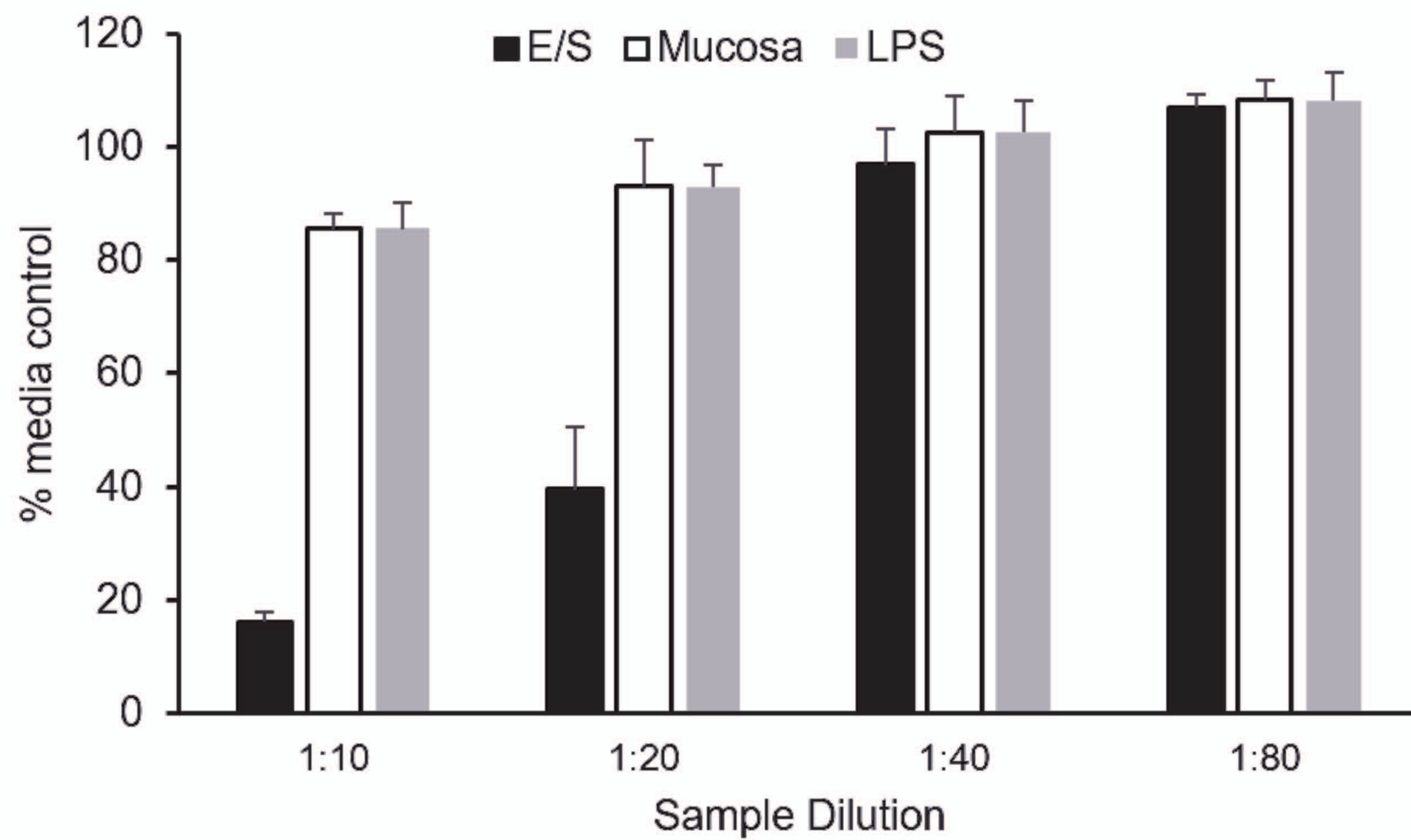
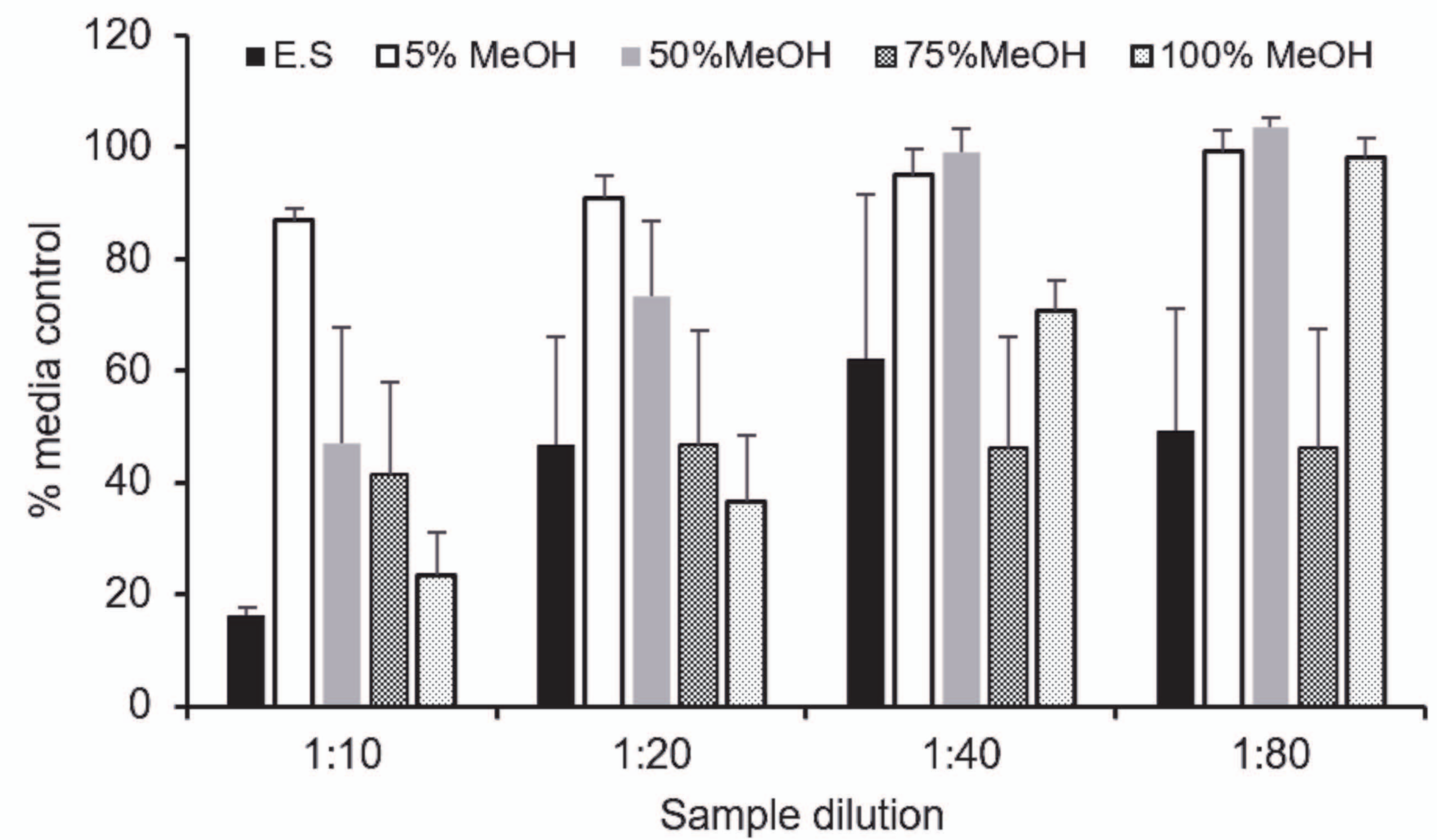
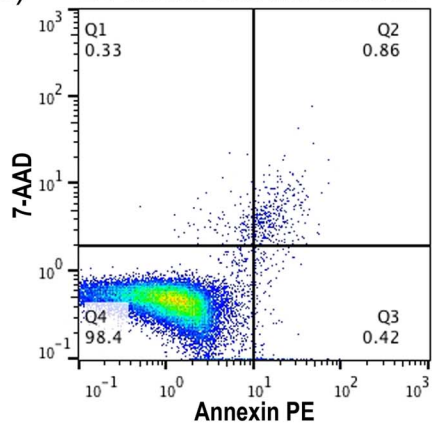


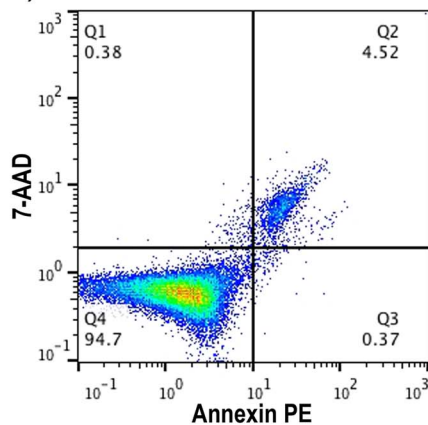
fig 4e



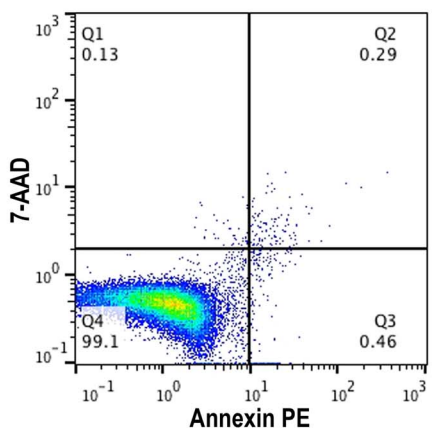
a) 24 hours media control



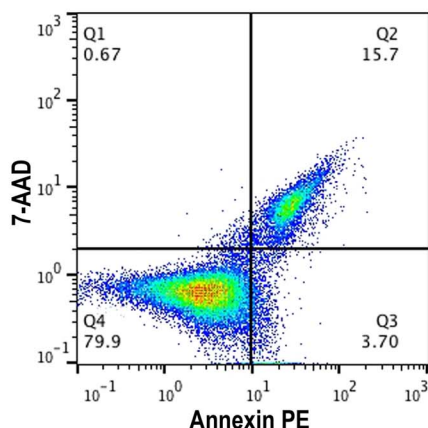
b) 24 hours ES treated



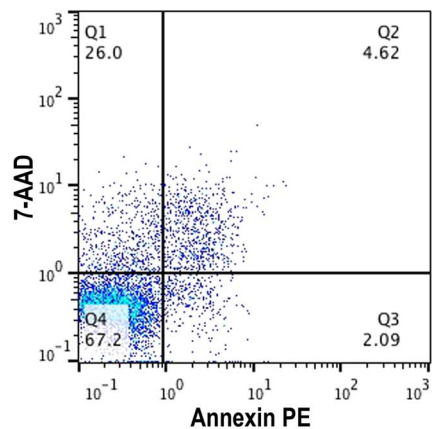
c) 48 hours media control



d) 48 hours ES treated



e) 72 hours media control



f) 72 hours ES treated

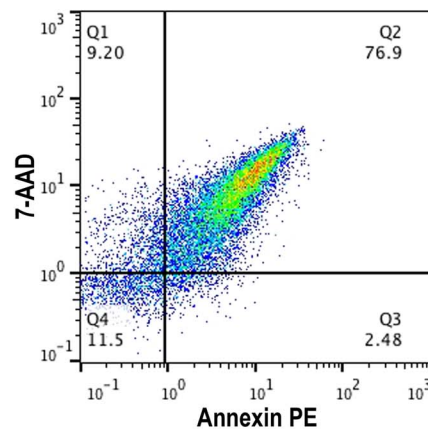


Fig 6a

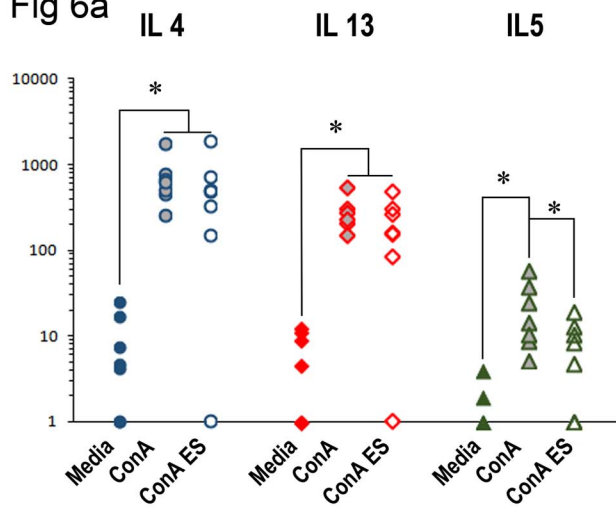


Fig 6b

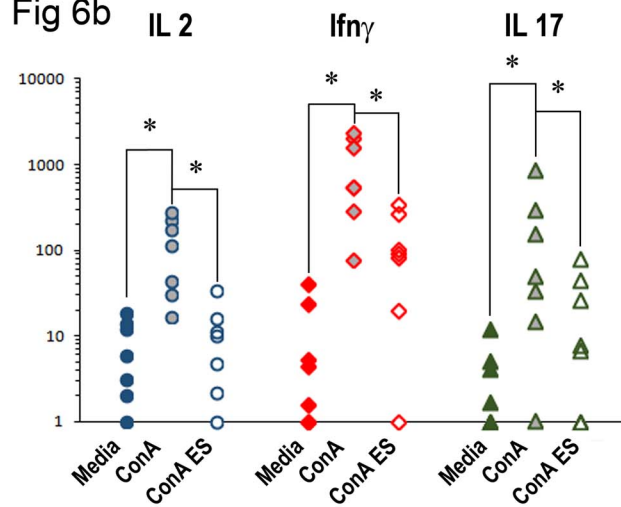


Fig 6c

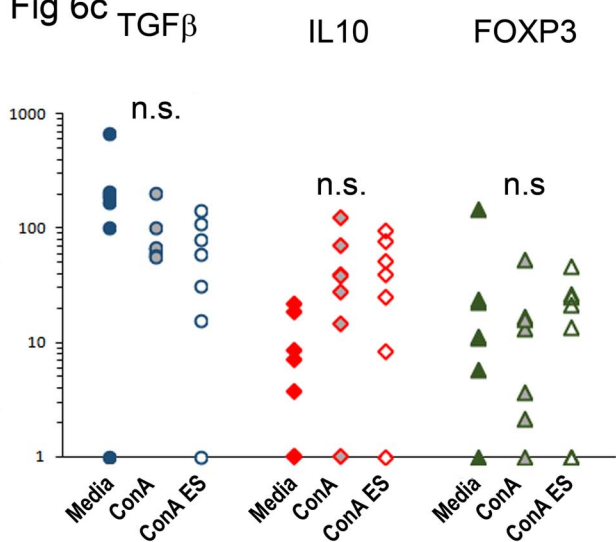


Fig 6d

