

1 **A semi-synthetic glycosaminoglycan analogue inhibits and reverses *Plasmodium***
2 ***falciparum* cytoadherence.**

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23

24 **Abstract**

25 A feature of mature *Plasmodium falciparum* parasitized red blood cells is their ability to bind
26 surface molecules of the microvascular endothelium via the parasite-derived surface protein
27 *Plasmodium falciparum* erythrocyte membrane protein 1 (PfEMP1). This ligand is associated
28 with the cytoadherence pathology observed in severe malaria. As pRBC treated with effective
29 anti-malarial drugs are still able to cytoadhere, there is therefore a need to find an adjunct
30 treatment that can inhibit and reverse the adhesion process. One semi-synthetic, sulfated
31 polysaccharide has been identified that is capable of inhibiting and reversing sequestration of
32 pRBC on endothelial cells *in vitro* under physiological flow conditions. Furthermore, it exhibits
33 low toxicity in the intrinsic (APTT assay) and extrinsic (PT assay) clotting pathways, as well as
34 exhibiting minimal effects on cell (HUVEC) viability (MTT proliferation assay). These findings
35 suggest that carbohydrate-based anti-adhesive candidates may provide potential leads for
36 therapeutics for severe malaria.

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38

39 Introduction

40 An important characteristic of the pathogenesis of severe malaria (SM) results from the
41 ability of parasitized red blood cells (pRBC) to sequester in the microvasculature, supported
42 by post-mortem studies of cerebral malaria (CM) that indicate high levels of pRBC bound in
43 brain microvessels [1, 2]. The involvement of sequestration in pathogenesis could be a result
44 of microvasculature occlusion, and/or downstream effects caused by interactions between
45 pRBC and the endothelium, including local inflammatory responses. [3].

46 The cytoadherence of pRBC to vascular endothelial cells occurs when PfEMP1, a
47 parasite derived molecule present on the surface of pRBC, binds to several distinct adhesion
48 molecules present on the surface of host endothelium. Previous studies have shown that
49 parasite isolates from children with SM bind to several receptors, suggesting that synergistic
50 effects between adhesion molecules may contribute to malaria pathophysiology. Yipp and
51 others indicated that, in some cases, multiple receptors may be involved in adhesion, and
52 recent data suggest that ICAM-1 and EPCR binding play a role in cerebral malaria [4-6]. The
53 role of cytoadherence in SM, coupled with splenic evasion, suggests that a compound capable
54 of reversing this adhesive phenotype would be desirable in terms of reducing clinical disease.
55 Previous work has concentrated on inhibiting cytoadherence, whereas for actual cases of
56 malaria it will be important that inhibitors of adhesion should also be able to reverse existing
57 adhesion.

58 It is preferable that antimalarial drug treatment is able to kill parasites in the non-
59 adhesive, ring stages to help prevent the next wave of pRBC from sequestering and thus
60 artemisinin is a good choice. This might explain the reduced mortality observed in field studies
61 from Thailand (SEQUAMAT) [7] and Africa (AQUAMAT) [8] in which artemisinin and quinine
62 (which kills exclusively mature pRBC) were compared. Despite this, there remains a high
63 mortality rate accounting for in excess of 50% of deaths during the first 48 hours following
64 hospital admission that is largely unaffected by the use of artemisinin-based combination

65 therapies (ACTs). This may be due to the pRBC already having been sequestered to the
66 endothelium. Consequently, there is a need for adjunct therapies to support the critically ill
67 patient, which can be used in combination with antimalarials such as artemisinin, to remove
68 the sequestered pRBC mass or reduce its effects on the host, whilst conventional drugs kill
69 the parasite effectively.

70 Polysaccharides, which are found throughout both the animal and plant kingdoms,
71 serve diverse functions in their tissues of origin and are frequently complex and heterogenous
72 in structure. In plants, they include acidic polysaccharides, usually as a result of the presence
73 of carboxylate groups (e.g. alginates and pectins) or O-sulfates groups (carrageenans), some
74 of which tend to form gels, often dependent on their association with divalent cation(s).
75 However, it is also possible to introduce other acidic groups (such as O-sulfates) by chemical
76 means and these modified plant polysaccharides exhibit a range of biological activities in
77 mammalian systems that arise from their ability to mimic the binding properties of the
78 mammalian glycosaminoglycan (GAG) class of extracellular polysaccharides [9], which
79 interact with many proteins.

80 Modified, semi-synthetic polysaccharides are capable of binding distinct proteins with
81 several levels of specificity. While highly acidic macromolecules could potentially interact in a
82 non-specific and non-physiologically relevant manner with proteins, several highly negatively
83 charged, sulfated polysaccharides, such as heparin (hep), heparan sulfate (HS), chondroitin
84 sulfate (CS), dextran sulfate, fucoidan, as well as the non-sulfated glycosaminoglycan
85 hyaluronic acid (HA), confer high affinity for particular proteins. Furthermore, anionic
86 carbohydrates have been reported to inhibit erythrocyte invasion by *Plasmodium* merozoites,
87 cytoadherence of pRBC to host cells, and to disrupt rosette formation between pRBC and
88 uninfected erythrocytes [10].

89 Polysaccharides with different levels and patterns of sulfation have been demonstrated
90 to inhibit *P. falciparum* growth and interfere with the adhesion of pRBC to the host endothelial

91 receptor CD36 [11-13]. Chemically O-sulfated cellulose was able to inhibit adhesion to CSA
92 expressed on both CHO cells and placental tissue [14]. Cellulose sulfate (ca. 50 kDa), with
93 significant levels of O-sulfate substitution at positions-2 (44 %) and -3 (36 %) of the glucose
94 repeating units, showed the most favourable inhibitory capacity and was able to reverse bound
95 pRBC to CHO cells and placental tissue [14]. In addition to being directly involved in adhesion,
96 it has been reported that sulfated CSA is also able to inhibit and reverse adhesion of CSA-
97 adherent pRBC *in vitro* [14] and in splenectomised monkeys *in vivo* [15]. Furthermore, altering
98 selected functional groups, especially sulfates on the saccharide branches, showed that it is
99 possible to reduce binding, and in some cases augment the attachment of the pRBC to
100 endothelium, all mediated by sulfated polysaccharides [16]. Other work using a modified
101 derivative of heparin (sevuparin) has demonstrated the ability of modified glycosaminoglycans
102 to inhibit cytoadherence and rosetting (the binding of infected erythrocytes to uninfected red
103 blood cells) [17-19]).

104 In this study, a variety of semi-synthetic, chemically modified polysaccharides with
105 various levels and patterns of sulfation, derived from important industrial polysaccharides,
106 were tested for their capacity to inhibit and reverse malaria cytoadherence, which may
107 contribute to the development of novel therapeutics capable of targeting adhesion of pRBC to
108 receptors such as ICAM-1 and CD36 on endothelial cells.

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111

112 **Materials & Methods**

113 **Endothelial cells**

114 Primary human umbilical vein vascular endothelial cells (HUVEC) and human dermal
115 microvascular endothelial cells (HDMEC) obtained from Promocell were cultured as per the
116 manufacturer's instructions. The Human Brain Endothelial Cell line (HBEC5i) was cultured as
117 described by Tripathi *et al.* [5]. Primary endothelial cells at passage 4-6 were used for all
118 experiments. Prior to use, cells were stimulated by addition of 1 ng.ml⁻¹ TNF for 18 h to allow
119 enhanced ICAM-1 expression on the surface of the endothelial cells.

120 **Parasite culture**

121 ItG (ITvar16) [20] and A4 (ITvar14) [21] laboratory parasite lines, which are well characterized
122 for their binding to ICAM-1 and CD36 [22], were cultured under standard conditions in RPMI
123 1640 medium supplemented with 37.5 mM HEPES, 7 mM D-Glucose, 25 ug.ml⁻¹ gentamycin
124 sulfate, 2 mM L-glutamine and 10% (v/v) pooled human serum at pH 7.2 in a gas mixture
125 comprising 96% nitrogen, 3% carbon dioxide, and 1% oxygen. Additionally, recently culture
126 adapted patient isolates: PO69, 8146 and 8026 were also included in the study [23].

127 **Plasmagel trophozoite enrichment**

128 Parasite culture at trophozoite stage was centrifuged (500 g, 5 min) and the pRBC pellet was
129 resuspended in a ratio of 2 volumes pellet to 3 volumes RPMI-based growth media without
130 human serum (incomplete medium) and 5 volumes Plasmion (Laboratoire Fresenius Kabi,
131 France), and allowed to settle for 20 - 30 min at 37 C. Trophozoite stage pRBC in the top layer
132 were washed three times in incomplete medium and the parasitaemia assessed by Giemsa
133 stained smear.

134

135 **Selection of pRBC on ICAM-1 purified protein**

136 To increase the homogeneity of the ItG parasite population that expresses a PfEMP-1
137 protein with high affinity for ICAM-1, the population was subjected to selection on ICAM-1
138 protein. 2.5 µg of ICAM-1 protein was coated on to 50 µl of protein A Dynabeads (Invitrogen)
139 in 200 µl of 1 % (v/v) Bovine Serum Albumin (BSA) in PBS and incubated for 1 hour at room
140 temperature with gentle rotation (15 rpm). Dynabeads were washed with 1% BSA/PBS and a
141 magnetic stand . 50 µl of synchronized and enriched ItG parasite culture using Plasmion were
142 incubated with the coated beads in 400 µl 1 % BSA/PBS for 45 min at room temperature by
143 gentle rotation. Bound pRBC were washed 3 times with 1 % BSA/PBS using the magnetic
144 stand. Beads were resuspended in 5 ml of complete RPMI media and transferred to culture in
145 a T25 culture flask with the addition of 100 µl of washed red blood cells.

146 **Chemical sulfation of polysaccharides**

147 Carbohydrate precursors were O-sulfated by a modified version of the chlorosulfonic acid
148 (CSA) sulphation method as described previously [9, 24]. Briefly, precursor carbohydrates
149 were dissolved in ice-cooled dimethylsulfoxide with pyridine, before chlorosulfonic acid was
150 added dropwise. The mixtures were held at 95 °C for 2 hr, cooled over ice and slowly
151 neutralized with sodium hydroxide. Ethanol precipitations were performed prior to extensive
152 dialyses (7 kDa cut-off membrane) against distilled water. Samples were lyophilized and stored
153 at 4 °C prior to use.

154

155 **Static inhibition adhesion assay screening of the chemically** 156 **modified, semi-synthetic anionic polysaccharides on** 157 **endothelial cells**

158 Initially, solutions of 44 chemically-modified semi-synthetic anionic polysaccharides
159 (see Supplementary Data) were diluted to 1 mg.ml⁻¹ in binding buffer (RPMI 1640 with 25 mM
160 HEPES, 11 mM D-Glucose, 2 mM L-glutamine pH 7.2) and screened for their anti-adhesive
161 properties using a static cell binding based assay as described by [22]. HUVEC, (2-6th
162 passage) were seeded onto 1% w/v gelatin coated 13 mm Thermanox coverslips (Nunc). Once
163 confluent, the cells were incubated overnight at 37 °C with 1 ng.ml⁻¹ TNF (Biosource
164 International). Cells were then washed with binding buffer prior to use. A suspension of 3%
165 pRBC and 1% HCT containing 1 mg.ml⁻¹ of compound was allowed to bind, for 1 hour with
166 mixing every 10 minutes, and following two dip washes, coverslips were placed in a gravity
167 wash for 30 min. Coverslips were then transferred to a second gravity wash for 10 min, fixed
168 in 1% v/v glutaraldehyde and stained with Giemsa. Control experiments lacking the addition of
169 the test compounds were also included. Coverslips were dried and mounted on slides using
170 DPX mountant (Sigma). Levels of adhesion were quantified by microscopy under 300x
171 magnification. The number of adherent pRBC per mm² was calculated.

172

173 **Flow adhesion assay inhibition by the chemically modified,** 174 **semi-synthetic anionic polysaccharides on endothelial cells**

175 This type of assay attempts to mimic the conditions seen in the post capillary venule by
176 allowing pRBC to flow over slides coated with endothelial cells. Permax chamber slides
177 (Nunc) were coated with 1% (w/v) gelatin for 1 hr at 37 °C, seeded with endothelial cells
178 (HUVEC; HDMEC; HBEC5i) and incubated until confluent. Confluent slides were then
179 incubated overnight at 37 C with 1 ng.ml⁻¹ TNF prior to use. PRBC suspensions (3%
180 parasitaemia and 1% HCT) with or without 1 mg.ml⁻¹ modified polysaccharides were flowed
181 over the endothelial cells for 5 min, followed by binding buffer (without the compounds) for 2
182 min to remove unbound cells. The flow rate yielded a wall shear stress of 0.05 Pa, used widely
183 to mimic wall shear stresses in the microvasculature. The number of adherent pRBC was

184 counted in six separate fields under 300x magnification and the density of parasitized red blood
185 cells per unit area (pRBC.mm⁻²) was calculated. All assays were carried out at 37 C and were
186 performed in duplicate or triplicate on three independent occasions.

187 **Flow adhesion assay reversal by the chemically modified,** 188 **semi-synthetic anionic polysaccharides on endothelial cells**

189 Reversal cell assays were carried out using a similar procedure to the flow inhibition adhesion
190 cell assay but with an initial phase in the absence of compounds. PRBC at 3% parasitaemia
191 and 1% HCT were flowed through the slide for 5 min to allow for pRBC adhesion. Flow was
192 continuous throughout the experiment at 0.05 Pa shear stress. Binding buffer was used to
193 remove unbound pRBC. Timing was started at the beginning of this wash, which continued for
194 2 min before the binding medium was swapped for medium containing the modified compound
195 being tested. The number of bound cells in six fields along the slide was counted at 0, 5, 10,
196 15 and 20 min. The number of adherent cells counted was used to calculate the pRBC bound
197 per mm².

198

199 **Prothrombin time (PT) coagulation assay**

200 Samples, controls and Thromborel S reagent (Siemens) were pre-warmed to 37 C prior to use.
201 Serially diluted, sulphated carbohydrate samples (50 µl) were incubated with normal human
202 citrated plasma (50 µl) for 1 min at 37 C prior to the addition of Thromborel S reagent (50 µl).
203 The time taken for clot formations to occur (an upper maximal of 2 min was observed) were
204 recorded using a Thrombotrak Solo coagulometer as per the manufacturer's instructions.
205 Water and sodium porcine mucosal heparin (203 IU/mg) were assayed as controls The EC₅₀
206 values of all semi-synthetic, sulphated carbohydrates were determined using a sigmoidal dose
207 response curved fitted post normalisation (with a 100% upper maximal at 2 mins; 0% lower
208 maximal represented by the time required for the water control to clot normal human citrated

209 plasma) with GraphPad Prism 6 software and compared to those obtained for the heparin
210 control.

211 **Activated partial thromboplastin time (aPTT) coagulation** 212 **assay**

213 Serially diluted, sulphated carbohydrate samples (25 μ l) were incubated with normal human
214 citrated plasma (50 μ l; NHSBT) and Pathromtin SL reagent (50 μ l; Siemens) for 2 min at 37 C
215 prior to the addition of calcium chloride (25 μ l; 50 mM). The time taken for clot formations to
216 occur (an upper maximal of 2 mins was observed) were recorded using a Thrombotrak Solo
217 coagulometer (Axis-Shield) as per the manufacturer's instructions. Water and sodium porcine
218 mucosal heparin (203 IU/mg; VWR) were assayed as controls. The EC₅₀ values of all semi-
219 synthetic, sulphated carbohydrates were determined using a sigmoidal dose response curved
220 fitted post normalisation (with a 100% upper maximal at 2 mins; 0% lower maximal represented
221 by the time required for the water control to clot normal human citrated plasma) with GraphPad
222 Prism 6 software and compared to those obtained for the heparin control.

223 **MTT cell proliferation assay**

224 Potential toxic effects of sulphated carbohydrates on endothelial cells were screened against
225 utilising the tetrazolium salt, 3-(4,5-dimethylthiazol-2-yl)-5-(3,4-diphenyl)-tetrazolium bromide (MTT)
226 proliferation assay. Briefly, a serial dilution of the test carbohydrate was prepared and added
227 to HUVEC cells (2×10^4 cells) in a multiwell plate (Greiner); a positive control, the Golgi disruptor
228 brefeldin A (10 μ l, 10 ng.ml⁻¹ in PBS), and PBS (10 μ l, as a negative control) were also included.
229 Post 48 hr incubation, MTT solution (10 μ l, 0.5% w/v in PBS) was added to all wells for 4 hr at
230 37 C. Finally, the supernatants were discarded, the cells washed (PBS) and treated with
231 dimethylsulfoxide (10 μ l). Cell proliferation levels were ascertained indirectly by
232 spectrophotometry at a λ of 540 nm.

233

234 **Statistical analyses**

235 Results shown are the mean of two independent experiments \pm Standard Deviation (SD). A
236 standard unpaired t-test was performed (Figs 1,2 and 3), ANOVA (Kruskal Wallis with post-
237 test) (Fig 4) and unpaired t-test (Fig 5) using GraphPad InStat3 software and considered
238 significant when $P < 0.05$.

239

240 Chemical characterisation methods for GSII are presented as supplementary information in S1
241 File.

242

243 **Results**

244 Both static and flow based assays were used for this work, partly to allow comparison with
245 other studies, which usually rely on static assays, but also to demonstrate the variation that
246 can be observed in these two different formats.

247 **Inhibition of pRBC binding to endothelial cells by chemically** 248 **modified, semi-synthetic anionic polysaccharides**

249 A number of chemically modified, semi-synthetic anionic polysaccharides were
250 screened for their anti-adhesion properties with a static based endothelial cell-binding assay
251 against two well-characterised parasite lines, ItG and A4, for their binding to TNF-activated
252 human primary endothelial cells HUVEC (human umbilical vein endothelial cells – large vessel
253 endothelium) and HDMEC (human dermal microvascular endothelial cells). TNF activation was
254 used to mimic the pro-inflammatory environment in a human host during malaria infection, and
255 upregulates the expression of a number of cytoadherence receptors. One of the key
256 differences between HUVEC and HDMEC, is that the latter expresses CD36, but HUVEC does
257 not (or at very low levels that do not support CD36-based adhesion). Of the 44 compounds
258 screened in a cell-based static assay (S1 Table ; S1 Fig), 10 showed potential adhesion
259 inhibition by producing 50% reduction in binding compared to the control receiving no
260 treatment (Table 1). From the flow-based screening we identified only two
261 compounds: glycogen type 2 sulfate from Oyster (MS34, GSII) and phenoxyacetylcellulose
262 sulfate (MS40, PACS)) that showed a significant adhesion inhibitory effect; they reduced the
263 binding of the A4 strain up to 70% and achieved 10-40% reduction in binding of the ItG strain
264 to TNF activated HDMEC in comparison to the un-treated control (S2 Fig).

265

266 **Table 1 – Compounds identified in primary screen.**

Compound	Description	Carbohydrate composition
1	Glycogen sulfate (type II) (GS)	$\alpha(1\rightarrow4)$, $\alpha(1\rightarrow6)$, polyglucan
2	Phenoxyacetyl cellulose sulfate (PACS)	$\beta(1\rightarrow4)$ phenoxyacetylated polyglucan
3	Ethyl cellulose sulfate	$\beta(1\rightarrow4)$ acetylated polyglucan
4	Gum Arabic sulfate	$\beta(1\rightarrow3)$, $\beta(1\rightarrow3)$ polygalactan core
5	Starch sulfate	$\alpha(1\rightarrow4)$ polyglucan
6	Poly-D-methylgalacturonic acid sulfate	$\alpha(1\rightarrow4)$ methylated polygalacturonic acid
7	Poly-D-galacturonic acid sulfate	$\alpha(1\rightarrow4)$ polygalacturonic acid
8	Tragacanth sulfate	$\alpha(1\rightarrow4)$ polygalacturonic acid, $\beta(1\rightarrow3)$ xylan
9	Hydroxypropyl methyl cellulose sulfate	$\beta(1\rightarrow4)$ hydroxypropylated polyglucan
10	Paramylon sulfate	$\beta(1\rightarrow3)$ polyglucan

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269

270 **Reversal effect of chemically modified, semi-synthetic** 271 **anionic polysaccharides on endothelial cells**

272 Having identified GSII and PACS as capable of inhibiting parasite adhesion, we further
273 screened these compounds in terms of their ability to reverse the binding of two-laboratory
274 parasite strains ItG (Fig 1) and A4 (Fig 2) to distinct endothelial cell lines (HUVEC, HDMEC
275 and HBEC5i) using a cell-based flow adhesion assay. HBEC5i is an immortalised human brain
276 microvascular cell line with a similar profile of receptor expression to HUVEC. For this
277 secondary screen, the comparisons for inhibitory effects were made to the situation at 0 min,
278 but it is known that there is a low-level loss of binding of PRBC during flow adhesion assays,
279 and this was incorporated into later experiments by running a 'no compound/ PBS' control for
280 the same period. GSII disrupted binding with 40% to 60% reduction of pRBC on all these cell
281 types, comparing 0 min with 20 min exposure to the compounds (Figs 1A and 2A), whereas
282 PACS showed smaller, variably significant reversal effects across all the endothelial cells (Figs
283 1B and 2B) (Table 2).

284 **Table 2 - p-values for cytoadherence assays: (unpaired t-test: Figs 1 and 2).** Comparison
 285 is between t=20 min and t=0 min and not a 'no compound' control at 20 min.

	ItG		A4	
	GSII	PACS	GSII	PACS
HUVEC	0.0079	0.0611	0.0037	0.018
HDMEC	0.0128	0.2326	0.0053	0.024
HBEC5i	0.002	0.0668	0.0098	0.0661

286

287

288 **Fig 1:** Reversal of ItG pRBC binding to different TNF-stimulated endothelial cells (HUVEC,
 289 HDMEC and HBEC5i) by 1 mg.ml⁻¹ of (A) GSII and (B) PACS for 20 mins under flow conditions.
 290 pRBC bound were calculated every 5 mins and expressed as a percentage (%) bound
 291 pRBC.mm⁻² compared to 0 min time point. Data for Figs 1 ~~and 2~~ are provided in supporting
 292 information in S1 ~~and S2~~ Data.

293 **Fig 2:** Reversal effect on A4 pRBC binding to different TNF-stimulated endothelial cells
 294 (HUVEC, HDMEC and HBEC) after flowing through 1mg.ml⁻¹ of (A) GSII and (B) PACS for 20
 295 mins under flow conditions. pRBC bound were calculated every 5 mins and expressed as a
 296 percentage (%) bound pRBC.mm⁻² compared to 0 min time point. [Data for Fig 2 are provided](#)
 297 [in supporting information in S2 Data.](#)

298

299 **Reversal effect of GSII and PACS on other parasite isolates**

300 To examine the effect of both GSII and PACS using a broader range of parasite
 301 variants, three recently laboratory-adapted, ICAM-1-selected patient isolates (P069, 8146 and
 302 8026) were used in addition to ItG [23]. Initially the ability of the compounds to disrupt pRBC
 303 already bound to ICAM-1 protein under flow conditions was tested. The reversal was similar
 304 for each parasite isolate, albeit with some variation, and around 20-30% pRBC were removed

305 with GSII being more effective than PACS (Fig 3 and Table 3), but only P069/ GSII was
 306 statistically significant. The reduction in binding was significant when using TNF activated
 307 HUVEC cells with a 40-50% reduction under static conditions that was significant with GSII
 308 (Fig 4 and Table 4), but lower and variably significant with PACS. Under flow conditions, GSII
 309 showed consistent reductions in binding of all isolates tested again reaching significant or near
 310 significant levels, unlike PACS, which showed no significant reductions in binding when
 311 compared to 'no-compound' control (Fig 5 and Table 5).

312 **Table 3 - p-values for cytoadherence assays: (unpaired t-test: Fig 3).** Comparison is
 313 between control (no compound) at 20 min with the relevant compound treatment at 20 min.

	ItG	P069	8146	8026
GSII	0.1094	0.0134	0.1328	0.0702
PACS	0.224	0.7762	0.4295	0.9271

314

315 **Table 4 - p-values for cytoadherence assays: (ANOVA-Kruskal-Wallis: Fig 4).**
 316 Comparison is with the relevant control for each isolate.

	ItG	P069	8146	8026
GSII	< 0.001	< 0.01	< 0.001	< 0.001
PACS	< 0.05	> 0.05	> 0.05	< 0.01

317

318 **Table 5 - p-values for cytoadherence assays: (unpaired t-test: Fig 5).** Comparison is
 319 between control (no compound) at 20 min with the relevant compound treatment at 20 min.

	ItG	P069	8146	8026
GSII	0.0418	0.0522	0.0673	0.0361
PACS	0.171	0.2095	0.407	0.5997

320

321

322 **Fig 3:** Reversal effect of 1 mg.ml⁻¹ GSII and PACS on binding of lab-adapted patient isolates
 323 A) ItG; B) P069; C) 8146; D) 8026 to ICAM-1 under flow conditions. pRBC bound were

324 calculated for every 5 mins and expressed as percentage (%) bound pRBC.mm⁻² compared to
325 0 mins. Control is no compound (PBS only). X-axis is time in minutes after addition of GSII,
326 PACS or PBS. Data for Fig 3 are provided in supporting information in S3 Data.

327 **Fig 4:** Reversal effect of 1 mg.ml⁻¹ GSII and PACS on binding to HUVEC using lab-adapted
328 patient isolates (P069, 8146 and 8026) under static assay conditions, with ItG used for
329 comparison. The remaining bound pRBC were counted and expressed as bound pRBC.mm⁻²
330 mean ± standard deviation. Control is no compound (PBS only). Data for Fig 4 are provided in
331 supporting information in S4 Data.

332 **Fig 5:** Testing 1 mg.ml⁻¹ GSII and PACS for their ability to reverse existing binding under flow
333 conditions. A) ItG; B) P069; C) 8146; D) 8026 reversal of binding on TNF-activated HUVEC.
334 pRBC bound were determined every 5 minutes and expressed as percentage (%) bound
335 pRBC. mm⁻² compared to 0 mins. Control is no compound (PBS only). X-axis is time in minutes
336 after addition of GSII, PACS or PBS. Data for Fig 5 are provided in supporting information in
337 S5 Data.

338

339 **Anticoagulant potential of GSII**

340 In light of the reversal efficacy of GSII, the ability of this sulfated carbohydrate to perturb
341 coagulation within pooled human plasma was determined. The prothrombin time (PT) and
342 activated partial thromboplastin time (aPTT) were measured for GSII, thereby determining the
343 overall effect on the extrinsic and intrinsic coagulation pathways respectively (both assays also
344 include the common coagulation pathway). The sulfated carbohydrate porcine mucosal
345 heparin (sodium), an approved clinical anticoagulant of known activity (201 IU.mg⁻¹) was
346 employed for comparison as a relevant control. The anticoagulant potential of GSII is highly
347 attenuated when compared to that of the sodium heparin control, in both the PT (EC₅₀ of 5.44
348 x 10⁴ µg.mL⁻¹ versus EC₅₀ of 2.7 x 10⁻¹ , respectively; Fig 6A) and aPTT (EC₅₀ of 1.53 x 10⁴
349 versus 6.74 x 10⁻¹ µg.mL⁻¹, respectively; Fig 6B) coagulation assays.

350

351 **Fig 6:** A) Prothrombin time (PT) assay to determine EC_{50} for the compounds for coagulation.
352 100% represents the inhibition of clotting, determined as a PT of >120 seconds, 0% represents
353 a normal PT clotting time for pooled human plasma ($\approx 13 - 14$ seconds). EC_{50} Porcine Mucosal
354 Heparin (PMH, the clinically used antithrombotic agent) = $2.74 \times 10^{-1} \mu\text{g.mL}^{-1}$; EC_{50} GSII = 5.44
355 $\times 10^4 \mu\text{g.mL}^{-1}$. B) Activated partial thromboplastin time (aPTT) assay. 100% represents the
356 inhibition of clotting, determined as a aPTT of >120 seconds, 0% represents a normal PT
357 clotting time for pooled human plasma (≈ 35 seconds). EC_{50} Porcine Mucosal Heparin (PMH)
358 = $6.74 \times 10^{-1} \mu\text{g.mL}^{-1}$; EC_{50} PACS = $1.53 \times 10^4 \mu\text{g.mL}^{-1}$. Data for Fig 6 are provided in supporting
359 information in S6 Data.

360

361 **Potential cytotoxicity of GSII**

362 Potential cytotoxic effects of GSII were screened for using the widely adopted MTT
363 assay, which detects the chromogenic change that occurs upon the mitochondrial reduction
364 of the tetrazol dye MTT to yield formazan in living cells. This reduction does not occur in
365 deceased cells, thereby acting as an indirect measure of toxicity, through the reduced levels
366 of cell proliferation that would be observed when a cytotoxic agent is present, compared to
367 that of the normally proliferating cell population. The incubation of HUVEC endothelial cells in
368 the presence of GSII at increasing concentrations up to 10 mg.mL^{-1} showed no apparent
369 evidence of cytotoxicity when compared to the PBS control (Fig 7).

370

371 **Fig 7:** Cell viability measured indirectly using the MTT cell proliferation assay after
372 incubation for 48 hours with varying concentrations of GSII on HUVEC cells. All results are
373 expressed as a percentage relative to a non-toxic control (PBS). Brefeldin A (a known Golgi
374 disruptor) at 10 ng.ml^{-1} was used as a toxicity control (data not shown). The data plotted is
375 the mean \pm S.D. of triplicate values. Statistical analysis was performed using an unpaired t-

376 test. All concentrations of GSII assayed were statistically insignificant from the PBS control
377 (i.e. $P > 0.05$). Data for Fig 7 are provided in supporting information in S7 Data.

378 Discussion

379 Erythrocytes infected with *P. falciparum* can bind to endothelial receptors, leading, in part, to
380 the clinical manifestations associated with SM. Consequently, molecules that can inhibit or
381 interrupt these interactions may have a role in improving understanding of host-parasite
382 biology, as well as in developing new therapies for severe disease. Highly sulfated
383 polysaccharides not only inhibit binding of pRBC to CSA [12-14], but can also reverse pRBC
384 adhesion in the placenta during pregnancy [25, 26]. Interactions between pRBC and host cell
385 membranes involve multiple interactions between several distinct ligands and receptors. The
386 cell surface receptors, including ICAM-1, CD36, EPCR, complement receptor 1 and
387 chondroitin sulfate-A, bind to different regions of the PfEMP-1 protein (the cysteine rich inter-
388 domain region (CIDR) and Duffy binding-like domains (DBL) [27, 28]. The extent of
389 involvement of these receptors varies between parasite variants and may correspond to
390 pathology [29]. For example, chondroitin sulfate A (repeating -4 GlcA β (1-3) GalNAc4S β (1-
391 disaccharide units) is implicated in placental malaria [25].

392

393 The interactions between pRBCs and host cells almost certainly involve manifold
394 protein-carbohydrate contacts across the interacting surfaces, in addition to HS. These lead to
395 strong interactions between cells that can be much more difficult to dislodge than the simple
396 sum of the individual components. This multi- or polyvalent effect is why small molecules are
397 unable to reverse binding interactions at reasonable concentrations. In contrast, if larger
398 molecules are employed, then effective inhibition becomes possible as long as the appropriate
399 geometric and stereochemical requirements to make effective bonds with receptors are
400 present [30].

401 Receptor-ligand binding is a dynamic process and this is especially relevant in the case
402 of multivalent interactions. In cell-to-cell interactions, multivalency exists at two distinct levels.
403 In the first, there is multivalency across the surface of the interacting cells, formed by many
404 individual molecular contacts. In the second, at the more detailed molecular level, there are

405 multiple bonds formed between individual interacting molecules, e.g. carbohydrate-protein or
406 protein-protein, which could comprise, for example, hydrogen bonds at several locations. The
407 dynamic aspects of both situations are essentially similar in nature, while being very different
408 in scale. Polyvalent interactions can be viewed as a series of discrete interactions, spread out
409 in space, but dynamic in nature and in a constant state of change. Each-interacting pair spends
410 some time bound and some unbound, but this lacks overall synchronisation. A small molecule
411 inhibitor can only bind one, or a few, of the available binding sites as and when they become
412 available, but cannot achieve efficient competition because, without resorting to extremely high
413 concentrations, it can never occupy sufficient sites locally to dislodge the original binding
414 molecules. If, on the other hand, a larger inhibitor with the appropriate spatial and
415 stereochemical characteristics to enable it to make first one, two and then several interactions
416 is introduced, then this can effectively compete with the ligand-receptor system and eventually
417 dislodge the original binding partners to replace one of them with itself. Such effective
418 competition can only be achieved by larger molecules, including polysaccharides.

419 The inhibitory interactions described here are not simply charge-driven. There is an
420 element of complementarity, supported by studies showing that low-sulfated structures can
421 inhibit pRBC binding [16]. GSII must possess specific dimensions and stereochemical
422 characteristics that enable it to act as an effective multivalent inhibitor at the level of both
423 molecular and cellular interactions. This requirement for particular geometric and charge
424 distribution is seen in that simply increasing charge does not necessarily improve inhibition
425 [see compound 36, supplementary data]. Additionally, polysaccharide inhibitors are not
426 necessarily acting only via a single type of ligand – receptor interaction, so polysaccharide
427 heterogeneity may be an advantage.

428 Ten potential inhibitory polysaccharides were identified (Table 1) from a library of 44
429 modified semi-synthetic anionic polysaccharides using a simple static binding assay. These
430 exhibited significant inhibition with two lab-adapted *P. falciparum* lines, A4 and ItG under static
431 conditions (S1 Fig). We then used the more complex, but physiologically relevant, flow-based

432 binding assay to investigate the ability of the ten polysaccharides to inhibit binding of these two
433 laboratory-adapted strains. GSII and PACS gave approximately 60-70% reduction in binding
434 when using A4, but not ItG (S2 Fig). The interaction between pRBC and endothelial cells varies
435 with the composition of the variant surface protein expressed on the different pRBC lines and
436 the repertoire of endothelial receptors on the host cells. Comparing binding efficiencies is
437 complicated by the different assay platforms used, however, under static conditions ItG
438 supports higher binding to ICAM-1 and CD36 than A4, which is also seen under static
439 conditions for HUVEC and HDMEC [22]. Under flow conditions, binding to endothelial cells for
440 these two parasite lines is different; A4 showing higher binding to TNF-induced HUVEC and
441 both lines showing essentially equivalent levels of binding to HDMEC. Therefore, the variability
442 seen in the level of inhibition between the different parasite lines may be due to differential
443 inhibition of a range of interactions, and no preference for any specific host receptor can be
444 discerned. This is un-surprising since the polysaccharide library was not designed to target
445 specific host-parasite interactions for cytoadherence.

446 Polysaccharides such as dextran sulfate and fucoidan can inhibit adhesion of *P.*
447 *falciparum* to host receptors such as CSA and CD36 [12-14] and regioselectively modified
448 polysaccharides, including modified carrageenans inhibit binding of pRBC to CD36 [31, 32],
449 as well as modified heparin structures [18]. GSII can now be added to this list but, the greater
450 challenge related to whether GSII or PACS could reverse bound pRBC, which is the situation
451 found clinically in SM. To answer this, three different EC (HUVEC, HDMEC and HBEC5i) were
452 used in combination with A4 and ItG to provide a further screen for activity prior to more
453 detailed analysis. Brain EC is potentially important as it may indicate whether GSII and PACS
454 could reduce sequestration in this tissue, as detected in cerebral malaria.

455 GSII gave a better response than PACS on different EC (HUVEC, HDMEC and
456 HBEC5i) and was more effective in reversal of A4 binding to HUVEC and HDMEC cells
457 compared to human brain microvascular endothelial cells (HBMEC). PACS did not show any
458 ability to reverse binding with ItG but gave slightly better results on reversing A4 binding to

459 HDMEC and HUVEC. Although GSII and PACS showed different effects on reversing
460 adhesion of both lab-adapted strains, neither had a significant effect on HBEC5i, which is a
461 concern in terms of developing either compound as an adjunct therapy for CM. However, the
462 HBEC5i used here is an immortalized rather than a primary cell line and this may have
463 influenced the level of binding and inhibition. Differential binding of parasite lines to HBEC5i
464 and primary HBMEC has been reported [33]. Further work on primary brain endothelium and
465 tissue sections will be needed, as will understanding the impact of releasing many rigid pRBC
466 into the circulation on the health of the patient.

467 GSII and PACS were further investigated by testing a panel of recently laboratory-
468 adapted parasite isolates (8146, 8026 and P069). Isolate 8146 is a strong ICAM-1 binder, with
469 8206 and P069 showing slightly lower levels of adhesion to this receptor [23]. Each parasite
470 strain showed roughly equivalent reversal responses under static and flow conditions, with up
471 to 80% reduction under flow (Figs 4 and 5) suggesting that GSII could have broad application
472 in terms of pRBC binding inhibition. GSII contains low sulfate (and charge) and the results
473 support the work of McCormick *et al.*, who showed that low sulfated glycoconjugates were able
474 to modulate binding of pRBC to different receptors [16].

475 Significant anticoagulant activity is a well-known off-target effect that has previously
476 hampered the application of some, but not all, sulphated molecules as potential therapeutic
477 agents. The PT and aPTT coagulation times of GSII suggest that it possesses negligible
478 anticoagulant potency (10^{-6}) compared to pharmaceutical heparin. Furthermore, GSII does not
479 significantly perturb endothelial cell proliferation by tetrazolium based dye, MTT, suggesting
480 that it possesses favourable bioactivity, is minimally anticoagulant and non-toxic.

481 This work has demonstrated the potential of chemically modified, semi-synthetic
482 anionic polysaccharides to both inhibit and reverse cytoadherence in malaria and offers
483 potential for the future development of pharmaceutical agents based on these materials.

484

485

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488

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615

616 **Supporting Information**

617 **S1 Fig. Screening compounds under static assay conditions.** Binding response of ItG
618 (top) and A4 (bottom) to modified polysaccharide compound at 1 mg/ml on TNF-stimulated
619 HUVEC and HDMEC under static condition (single screening). pRBC binding (3%,
620 parasitemia; 1% HCT) observed after polysaccharide treatment for one hour. The remaining
621 bound pRBC after gravity wash were counted and expressed as % bound pRBC/ mm² (N=1)
622 compared to control, without polysaccharide.

623

624 **S2 Fig. Screening compounds under flow assay conditions.** Binding response of ItG and
625 A4 to modified polysaccharides compounds at 1 mg/ml on TNF-stimulated HDMEC under
626 flow conditions. The remaining bound parasites after 20 mins wash was counted and
627 expressed as % bound pRBC/ mm² ± standard deviation compared to control without
628 polysaccharide. ND; not done. MS34 (ItG and A4) & MS40 (A4), *P* < 0.05 (compared to
629 control).

630

631 **S3 Fig. Characterisation of GSII.** ¹H NMR spectrum of GSII at 400 MHz, with 128 scans, 2s
632 delay. Inset - Comparison of ¹H NMR spectra of GSII and its unsulfated precursor, glycogen
633 type II. This is provided as background information and is not cited in the paper.

634

635

636 **S1 Table. Sulfated carbohydrates assayed.** A list of all the compounds screened in this
637 paper.

638 **S1 File. Supplementary methods.** Description of methods associated with chemical
639 characterisation of GSII.

640 **S1 ~~and~~ Data: [Primary data supporting Fig 1.](#)**

641 **S2 Data: Primary data supporting [Figs 1 and 2.](#)**

642 **S3 Data: Primary data supporting Fig 3.**

643 **S4 Data: Primary data supporting Fig 4.**

644 **S5 Data: Primary data supporting Fig 5.**

645 **S6 Data: Primary data supporting Fig 6.**

646 **S7 Data: Primary data supporting Fig 7.**

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