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# ▶ To cite this version:

Euripide Avokpaho, Tania C. D'almeida, Ibrahim Sadissou, Léonidas Tokplonou, Rafiou Adamou, et al.. HLA-G expression during hookworm infection in pregnant women. Acta Tropica, 2019, 196, pp.52 - 59. 10.1016/j.actatropica.2019.04.026 . hal-03485711

# HAL Id: hal-03485711 https://hal.science/hal-03485711v1

Submitted on 20 Dec 2021

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Version of Record: https://www.sciencedirect.com/science/article/pii/S0001706X18303279 Manuscript\_aace7a2689ce2e303fe58b4cf6e264aa

# 1 HLA-G expression during hookworm infection in pregnant women

Euripide Avokpaho<sup>1</sup>, Tania C. d'Almeida<sup>1</sup>, Ibrahim Sadissou<sup>1,2,3</sup>, Léonidas Tokplonou<sup>1,2</sup>,
Rafiou Adamou<sup>1,2</sup>, Paulin Sonon<sup>1,2,3</sup>, Jacqueline Milet<sup>1</sup>, Gilles Cottrell<sup>1</sup>, Amandine Mondière<sup>1</sup>,
Achille Massougbodji<sup>2</sup>, Kabirou Moutairou<sup>2</sup>, Eduardo A. Donadi<sup>3</sup>, Celso Teixeira Mendes
Junior<sup>4</sup>, Benoit Favier<sup>5,6</sup>, Edgardo Carosella<sup>5,6</sup>, Philippe Moreau<sup>5,6\*</sup>, Nathalie Rouas-Freiss<sup>5,6\*</sup>,
André Garcia<sup>1\*</sup> and David Courtin<sup>1\*</sup>

- 7 <sup>1</sup> MERIT IRD, Université Paris 5, Sorbonne Paris Cité, Paris
- 8 <sup>2</sup> Université d'Abomey-Calavi, Cotonou, Bénin

<sup>3</sup> Division of Clinical Immunology, School of Medicine of Ribeirão Preto, University of São
Paulo, Brazil

<sup>4</sup> Department of Chemistry, Faculty of Philosophy, Sciences and Letters of Ribeirão Preto,

12 University of São Paulo, Brazil

- <sup>5</sup> CEA, Institut des Maladies Emergentes et des Thérapies Innovantes (IMETI), Service de
- 14 Recherche en Hémato-Immunologie (SRHI), Hôpital Saint-Louis, IUH, Paris, France ;
- <sup>6</sup> Université Paris Diderot, Sorbonne Paris Cite, IUH, Hôpital Saint-Louis, UMRE5, IUH,
   Paris, France
- 17 \* Joint senior authors

Introduction: HLA-G plays a key role on immune tolerance. Pathogens can induce soluble HLA-G (sHLA-G) production to down-regulate the host immune response, creating a tolerogenic environment favorable for their dissemination. To our knowledge, no study has yet been conducted to assess the relationship between sHLA-G and geohelminth infections.

Methods: The study was conducted in Allada, Southeastern Benin, from 2011-2014. The study population encompassed 400 pregnant women, included before the end of the 28<sup>th</sup> week of gestation and followed-up until delivery. At two antenatal care visits and at delivery, stool and blood samples were collected. Helminths were diagnosed by means of the Kato-Katz concentration technique. We used quantile regression to analyze the association between helminth infections and sHLA-G levels during pregnancy.

**Results:** sHLA-G levels gradually increased during pregnancy and reached maximal levels at delivery. Prevalence of helminth infections was low, with a majority of hookworm infections. We found significantly more hookworm-infected women above the 80<sup>th</sup> quantile (Q80) of the distribution of the mean sHLA-G level (p<0.03, multivariate quantile regression). Considering only women above the Q80 percentile, the mean sHLA-G level was significantly higher in hookworm-infected compared to uninfected women (p=0.04).

**Conclusion:** High levels of sHLA-G were associated with hookworm infection in pregnant women. This result is consistent with the potential involvement of sHLA-G in immune tolerance induced by helminths during pregnancy.

38

# 39 **1. Introduction**

Helminths, and particularly geohelminths, infect 1.45 billion people worldwide (Pullan et al., 40 41 2014). The prevalence of geohelminth infection during pregnancy in endemic African areas could reach more than 70% (van Eijk et al., 2009). Children born to infected mothers are 42 reported to be more susceptible to helminth infection and to display weaker responses to 43 vaccines (Cooper et al., 1998; Elias et al., 2005; Malhotra et al., 1999; Malhotra et al., 2003; 44 45 Mehta et al., 2012; Sabin et al., 1996). These last observations could be explained by the phenomenon of immune tolerance. The latter was also suspected in children exposed to 46 malaria in utero (Malhotra et al., 2009). Lymphocyte responses to parasite antigens and 47 detection of specific immunoglobulins in cord blood, consistent with in utero sensitization, 48 have been described for several helminth species including ascariasis and hookworm 49 (Guadalupe et al., 2009; Larocque et al., 2006; Pit et al., 2000). Considering that the human 50 leukocyte antigen-G (HLA-G) is a well-recognized immune checkpoint molecule, the study of 51 the potential involvement of this molecule in geohelminth infections is of interest. 52

Human Leukocyte Antigen-G (HLA-G) is a non-classical HLA class I antigen that differs from 53 classical class I molecules by its restricted tissue distribution, diversity of protein isoforms 54 and limited gene polymorphism (Donadi et al., 2011). Seven isoforms of the protein are 55 56 generated by alternative splicing, four are membrane bound (HLA-G1-4) and three are soluble (HLA-G5-7 of which the shed HLA-G1 and the soluble HLA-G5 are usually measured 57 in plasma samples). HLA-G further differs from classical class I molecules through its 58 immune regulatory functions, which is of paramount importance in maternal-fetal immune 59 tolerance by inhibiting the cytotoxic effect of maternal NK cells (Moreau et al., 1998; Rouas-60 Freiss et al., 1997). A lack or a failure of HLA-G expression may lead to a breach of fetal-61 maternal tolerance and consequent fetus rejection (Cecati et al., 2011; Papamitsou et al., 62 2014). Modulation of HLA-G expression has also been described in tumor cell development, 63 64 allogenic organ engraftment (Carosella et al., 2015) and viral infections (Amiot et al., 2014b). It has been suggested that high HLA-G levels induce tolerance and contribute to immune 65

evasion by tumor cells (Morandi et al., 2016; Pistoia et al., 2007) or viruses (Amiot et al.,
2014a; Amiot et al., 2014b).

In tropical areas, parasitic infections during pregnancy modify the immune development of 68 69 the fetus (Dauby et al., 2012). Such infections affect the susceptibility of young children to infectious pathogens, their response to vaccines, and the development of allergic disorders 70 such as asthma and rhinitis (Thornton et al., 2010). Placental Plasmodium falciparum 71 infection is associated with susceptibility to malaria during infancy (Bouaziz et al., 2017; Le 72 73 Hesran et al., 1997; Le Port et al., 2011; Mutabingwa et al., 2005; Schwarz et al., 2008) and it has been hypothesized that immune tolerance may be responsible for the shortening of time 74 75 to the first occurrence of malaria infection, but a complete explanation of the physiopathology involved has yet to be established (Moya-Alvarez et al., 2014). Furthermore, placental 76 77 malaria is also associated with non-malarial infections, suggesting that immune tolerance may go beyond malaria-specific immunity (Rachas et al., 2012). The association of HLA-G 78 79 with the susceptibility to malaria during infancy and with low birth weight could reflect the potential involvement of sHLA-G in immune tolerance mediated by placental malaria 80 (d'Almeida et al., 2016; d'Almeida et al., 2017; Sadissou et al., 2014). 81

Considering that the expression of HLA-G may modulate viral and parasitic infections (Sabbagh et al., 2018), and considering that HLA-G plays an important role on pregnancy, in the present study we investigated the potential association between soluble HLA-G levels and hookworm infections during pregnancy.

#### 86 2. Material and methods

# 87 2.1. Study area, study design and follow-up

Our study took place in Allada, a town located 56 km North Cotonou, the economic capital of Benin. The study participants were residents of 21 villages surrounding two primary health facilities: Sékou and Attogon. This study is part of a larger one encompassing 1,183 human immunodeficiency virus (HIV)-negative pregnant women participating in the randomized

92 Malaria in Pregnancy Preventive Alternative Drugs (http://clinicaltrials.gov/ct2/show/ NCT00811421) trial, designed to compare intermittent preventive malarial treatment (IPTp) 93 94 with either sulfadoxine-pyrimethamine or mefloquine. The study protocol has been described 95 in detail elsewhere (Accrombessi et al., 2015; Ouedraogo et al., 2012). The first 400 children of these women were followed-up to investigate the impact of sociodemographic, economic, 96 environmental and biologic factors such as the parasitological status (geohelminth and P. 97 98 falciparum carriers) during pregnancy on infant health during the two years of life in southern Benin (Accrombessi et al., 2015; d'Almeida et al., 2017). The 400 mothers included in the 99 present study are the mothers of these 400 children. Women were included before a 100 gestational age (GA) of 28 weeks, and two doses of IPTp were administered at antenatal 101 visits (ANV). The second dose was given at least 1 month after the first one. During the first 102 103 ANV, a total dose of 600 mg of mebendazole was given to the women to be taken at home (100 mg two times per day for 3 days) according to the guidelines of the Beninese Ministry of 104 105 Health. Between antenatal visits, women were encouraged to attend the health facility for 106 any health complaint. At inclusion, a questionnaire was applied by the midwife to obtain data regarding socio-demographic characteristics, gynecological and obstetrical history. During 107 ANV1 and ANV2 (for IPTp1 and IPTp2, respectively) and at delivery, blood and stool 108 109 samples were collected before the intake of IPTp, to allow plasma sHLA-G measurements as 110 well as malaria and geohelminth diagnoses. At delivery, a placental blood smear was collected to detect placental malaria. Malarial infection was defined as the presence of 111 asexual P. falciparum parasites in a blood smear. All the medications prescribed during the 112 follow-up were free of charge. 113

114 Soluble HLA-G quantification

Soluble HLA-G (sHLA-G) was quantified using MEM-G/9 antibody, which recognizes the most abundant soluble isoforms (shed sHLA-G1 and intron 4 containing secreted HLA-G5), and an anti-human β2-microglobulin antibody, respectively, as capture and detection antibodies (Menier et al., 2003; Rebmann et al., 2005). Microtitration plates were coated with

10 µg/mL MEM-G/9 mouse-anti-human HLA-G mAb (Exbio, Praha, Czech Republic) and 119 incubated overnight at 4°C. Plates were saturated with 300 µL ready to use diluent buffer 120 121 (DAKO, Carpinteria, CA, USA) for 2 hours. Plasma samples were diluted (1/2) in diluent buffer, tested in duplicate and incubated for 2 hours. After incubation and washing, 122 monoclonal anti-human beta2-microglobulin antibody (DAKO, Glostrup, Denmark), which 123 recognizes the immobilized antibody sHLA-G complex, was added to the wells and incubated 124 125 for 1 hour. The plates were then incubated for 1 hour with 100 µL (1:200) envision buffer + system HRP (DAKO, Carpinteria, CA, USA) to obtain anti-β2-microglobulin-horseradish 126 peroxidase complex to improve the efficiency of the reaction. All incubation steps were 127 performed at room temperature and followed by four washes using washing buffer (H<sub>2</sub>O, 128 PBS 1X, 0.1% Tween 20). The plates were incubated for 30 min with substrate 129 (Tetramethylbenzidine super sensitive-TMB, Sigma Aldrich, Saint Louis, MO, USA) and 130 absorbance was measured at 450 nm after adding HCL (1 N). Total sHLA-G levels were 131 determined from a five-point standard calibration curve using dilutions (12.5- 200 ng/mL) of 132 133 HLA-G5 purified from M8-HLA-G5 cell line culture supernatants. Results are presented in 134 ng/mL.

# 135 Helminth identification

Women were invited to provide a sample of their fresh morning stool put in a 120 mL plastic 136 container that was distributed beforehand. Stool samples were transferred to the health 137 138 centre laboratory and were analysed at the same day, using standard 41.7 mg plastic templates, and thick smears were examined under a light microscope by experienced 139 laboratory technicians. Microscopical identification of Ascaris lumbricoides and Trichuris 140 trichiura eggs is easy recognizable using the Kato-Katz method; however, Ancylostoma 141 142 duodenale and Necator americanus eggs cannot be distinguished. For this reason, we included both Ancylostoma duodenale and Necator americanus species in a same hookworm 143 144 group.

#### 145 Ethics

The protocol received approval from the Comité Consultatif de Déontologie et d'Ethique of 146 147 the Institut de Recherche pour le Développement (IRD, France) and from the Ethics la 148 Committee of the Faculté des Sciences de Santé of Cotonou (N°43/11/2010/CE/FSS/UAC). Before inclusion, the protocol of the study was explained in the 149 relevant local languages and written consent was obtained. Women who could not read and 150 write provided thumbprints to confirm their agreement to participate in the study after a nurse 151 152 had explained the study.

# 153 **Definition and characteristics of the study variables**

## 154 Soluble HLA-G levels

155 The outcome variable is the level of sHLA-G, used as a quantitative variable after a 156 logarithmic transformation to take into account the skewness of the distribution.

#### 157 Variable of interest: Helminth infections

Both hookworms (*Ancylostoma duodenale, Necator americanus*) and round worms (*Ascaris lumbricoides* and *Trichuris trichiura*) were detected in faecal samples. However, due to the very low prevalence of both *Ascaris lumbricoides* and *Trichuris trichiura*, our analyses focused on hookworm infection during pregnancy, by creating a variable coded 1 <u>if at least</u> <u>one hookworm egg</u> was detected during follow-up.

## 163 Covariates

164 Information on the following variables was collected: gravidity, IPTp regimen, placental

165 malaria, inclusion heath center, ethnicity, marital status, occupation, education level, latrine

- use at home and "wealth score". The wealth score was computed using four socio-economic
- 167 items which were theoretically associated with social welfare: motorbike, television,
- electricity, and fridge. Each item could take the value 0 or 1 (yes = 1, no = 0), and the score
- 169 was obtained by summing the values of each item.

#### 170 Statistical analysis

Categorical variables were described using proportions and 95% confidence interval and the continuous variables were described by the median and interquartile range. To compare proportions, we used the Chi-square, Fisher T and Cuzick trend tests. Continuous variables were compared by means of Student T test, Wilcoxon-Mann-Witney rank-sum test and analysis of variance (ANOVA). We checked for normality of distribution using Skewness/Kurtosis normality test and compared the variances with Breusch-Pagan/Cook-Weisberg heteroscedasticity test. A p-value less than 5% was considered as significant.

Classical methods for repeated measurements (mixed models) were not used to analyse 178 sHLA-G level at antenatal visits because each antenatal visit had its particular characteristics 179 and cannot be considered as a repeated measurement of the same variable under the same 180 conditions. Measurements at the first antenatal visit were done before any intervention. 181 Medication was given at ANV1 and ANV2 (IPTp for malaria during the two first antenatal 182 visits and mebendazole during the first one). Furthermore, the third measurement occurred at 183 184 delivery. Consequently, to create a unique variable accounting for the level of sHLA-G during pregnancy, a mean sHLA-G (MsHLA-G) level was computed. To take into account the fact 185 that not all women were present at each measurement time, and that sHLA-G levels 186 187 increased significantly at delivery, the MsHLA-G was computed for each woman as the mean of her visit-adjusted sHLA-G level. The visit-adjusted sHLA-G levels were obtained by 188 subtracting the mean sHLA-G of the population at each visit. The association between 189 MsHLA-G and hookworm infection was analyzed by means of quantile regression that takes 190 into account the heteroscedasticity of the distribution (Koenker, 2005; Sims, 1996). Univariate 191 and multivariate analyses were done and variables with p-value less than 0.2 in univariate 192 193 analysis were considered in the multivariate model. The variable of interest was "at least one hookworm infection during the follow-up". Hookworm and "inclusion health facility" were 194 included in the multivariate analysis. The other covariates considered were: gravidity, 195 regimen of intermittent preventive treatment, placental malaria, age, ethnicity, marital status, 196 occupation, education, latrines use at home and wealth score. Starting from the initial model 197

we proceeded using a step-by-step backward regression. Only the variables exhibiting pvalue<0.05 were maintained in the final model. All analyses were performed using Stata12</li>
software (StatCorp LP, College Station, TX, USA).

201 Results

#### 202 Study population

Four hundred pregnant women were included in the study. Their median age was 25 years (range: 18-42; Interquartile range [IQR]: 21-30). The median gestational age (GA) at inclusion was 24 weeks (range: 13-28; IQR: 20-26). The median wealth score based on 4 variables (ownership of motorbike, electricity, television, and refrigerator) was 1, (range: 0-4, IQR: 1-2). The socio-demographic and economic characteristics of the participants are summarized in Table 1.

#### 209 Helminth infections

The results of parasitic coprology are shown in Figure 1. Stool samples were collected from 210 211 395 (98.75%) participants at ANV1, 387 (96.75%) at ANV2 and 355 (88.75%) at delivery, giving a total of 1137 stool samples collected. Among the 1137 stool samples, helminth eggs 212 were detected in 88 stool samples (7.7%). The sample positivity rate to helminth infections 213 was 9% at ANV1, 8.8% at ANV2 and 5% at delivery. Hookworm eggs (Ancylostoma 214 duodenale and Necator americanus) were found in 74 samples (27 infections at ANV1, 31 at 215 216 ANV2 and 16 at delivery), 10 samples contained eggs of Ascaris lumbricoides (1 at ANV1, 6 217 at ANV2 and 3 at delivery) and eggs of Trichuris trichiura were found in 3 stool samples (1 sample positive at ANV1, 1 at ANV2 and 1 at delivery, all from different participants). 218

The prevalence of helminth infection (at least one geohelminth in the stools during the whole
 follow-up) in the study population was 18.5% (74/400). The majority of the helminth infections
 detected in the stools were hookworm infections (86.5% (64/74)), while Ascaris lumbricoides
 and Trichuris trichiura eggs were respectively detected in 13.5% (10/74) and 4% (3/74) of the

223 positive stool samples. Two participants were co-infected with both hookworm and Ascaris
 224 *lumbricoides*, one diagnosed at ANV2 and the other diagnosed at delivery.

The distribution of hookworm infection according to the different characteristics of the study population is presented in Table 2. The prevalence of hookworm infection was significantly higher in Sekou health facility and in the Aizo ethnic group (p=0.02 in both cases, Chi<sup>2</sup>). The possession of latrines was associated with a significantly lower prevalence (p=0.001, Chi<sup>2</sup>) as was a high wealth score (p=0.024, Chi<sup>2</sup>).

#### 230 Soluble HLA-G level

Soluble HLA-G measurement was available on three occasions from 85% of mothers and 98% were sampled at least twice. sHLA-G levels were stable over the two first visits (mean sHLA-G level at ANV1 = 10.09 ng/ml and mean sHLA-G level at ANV2 = 10.57 ng/ $\mu$ l) but increased significantly at delivery (mean sHLA-G level = 17.26 ng/ $\mu$ l (p<10<sup>-3</sup>, ANOVA)).

#### 235 Relation between hookworm infection and adjusted mean of sHLA-G.

236 The distribution of adjusted means of sHLA-G (MsHLA-G) is presented in Figure 2. Despite 237 the log-transformation, both the hypotheses of normality (p<0.0001, Shapiro Wilk normality test) and of homoscedasticity (p=0.02; Breusch-Pagan/Cook-Weisberg heteroscedasticity 238 239 test) were significantly rejected. To deal with both the skewed distribution and heteroscedasticity, a quantile (5<sup>th</sup>, 10<sup>th</sup>, up to 95<sup>th</sup> quantile) regression (Chamberlain, 1994) 240 was used to analyze the effect of hookworm infection and of other covariates (inclusion 241 health facility, latrines use at home, wealth score, gravidity, regimen of intermittent preventive 242 treatment, placental malaria, age, ethnicity, marital status, occupation, education) on the 243 different quantiles of mean sHLA-G levels. 244

In univariate analysis, inclusion health facility had a significant effect on the distribution of sHLA-G with significantly fewer women above the Q50 of the distribution of the mean level of sHLA-G included at Sekou health center (Table 3). No significant effect of the others covariates was detected on the distribution of sHLAG. As hookworm infection is the variable

of interest, the multivariate model of regression included the inclusion health center and 249 hookworm infection. The effect of inclusion health facility on the distribution of sHLA-G 250 251 remained significant in multivariate analysis and a significant effect of hookworm infection was observed (p<0.03, multiple quantile regression). Significantly more hookworm-infected 252 women were found above the Q80 percentile of sHLA-G distribution (Table 3 and Figure 3). 253 Considering only women above the Q80 percentile, the geometric mean level of sHLA-G was 254 255 significantly higher in hookworm-infected than in uninfected women (1.57 vs 1.41 256 respectively; p=0.04, Student T test).

#### 257 Discussion

The main results of the present work show a relatively low frequency of helminth infection during pregnancy of Beninese women; however, we observed an association between the plasma levels of the tolerogenic molecule HLA-G and hookworm infection.

Prevalence of hookworm infections in our study decreased from 9% at ANV1 to 5% at delivery. <u>The detection of hookworms was performed with the inexpensive and field-</u> applicable Kato-Katz method however the low sensibility of the Kato-Katz method probably <u>underestimates the intestinal helminth infection prevalence in our study population</u>.

This finding could be explained by the intake of one dose of mebendazole (600 mg) during 265 antenatal visits after 16<sup>th</sup> week of gestation, as recommended by the Beninese National 266 Health Program. Antihelminthic treatment during pregnancy aims to fight against maternal 267 anemia (Ouedraogo et al., 2012), which has an important impact on neonatal and infant 268 health (Koura et al., 2012). Helminth infections during pregnancy are also associated with 269 270 increased susceptibility to geohelminth infections in the children (Mehta et al., 2012) and to weaker anti-vaccine responses (Labeaud et al., 2009). The mechanisms by which maternal 271 helminth infection can influence immune development of the fetus are partially understood 272 (Mpairwe et al., 2014). However, epidemiological evidence suggests that in utero 273 sensitization results in down-regulation of immune responses in offspring (Dauby et al., 274

275 2012), leading to a phenomenon of immune tolerance, in which HLA-G could play a crucial276 role.

277 In the multivariate model of regression, sHLA-G levels were associated with health center 278 and hookworm infection. Women included at Sekou health center had lower level of sHLA-G 279 than those from Attogon health center. Significant differences in term of ethnicity group 280 distribution and wealth score were observed between both health centers (p<0.001 and p<0.001 respectively, Chi-Square test). Women from Fon ethnicity group were more 281 282 represented at Attogon than at Sekou (57 % versus 9 %) while women from Aizo ethnicity group were more frequent at Sekou than Attogon (82 % versus 31 %). Differences in 283 behaviour and/or in genetic mechanism influencing regulation of HLA-G expression between 284 both ethnicity groups could partly explain the modulation of sHLA-G observed between 285 health centers. It is important to emphasize that Sekou is a semi-rural village while Attogon is 286 287 more rural therefore difference of HLA-G expression could also be related to environmental factors, such as the prevalence of other pathogens that could also influence the expression 288 289 of sHLA-G. The elevated risk of exposure to mosquito bites was previously associated with an increased expression of sHLA-G (Sadissou et al., 2014). In the present work, no 290 association was observed between peripheral or placental malaria infections during 291 292 pregnancy and a significant modulation of plasmatic sHLA-G in mothers (d'Almeida et al., 293 2017). Difference of sHLA-G level observed between the health centers seems not directly 294 related to malaria infection.

295 Our results show that there are more hookworm infected women above the Q80 and that 296 these women have significantly higher levels of sHLA-G than not infected women from the same quantiles. The use of Kato-Katz is a potential limitation in our study. Some intestinal 297 298 helminth infections could be failed in case of light-intensity hookworm infections and we have potentially included false negatives in the group of "hookworm uninfected women". Such 299 misclassification should not significantly affect our result because a higher level of sHLA-G is 300 301 expected in presence of false positives in the group of women with no hookworm egg. Such situation of misclassification could typically lead to a lack of statistical power, which could 302

303 reduce the chance to find an association between hookworm infection and sHLA-G levels.
 304 The use of more sensitive tools such as the PCR or coproculture methods could reinforce the
 305 result of association between hookworm infection and sHLA-G levels obtained in our study
 306 and give a better insight of the hookworm prevalence in the study areas.

It is known that hookworm infection results firstly in a T helper 2 cell type polarization, 307 characterized by the proliferation of T cells that produce IL-4, IL-5 and IL-13, activation of 308 309 basophils, eosinophils, mast cells and secretion of high levels of IgE. Then, a progressive 310 and generalized immunosuppression, characterized by induction of regulatory responses (regulatory T, B and dendritic cells) with increased expression of the regulatory cytokines IL-311 10 and TGF- $\beta$ , often occurs. The progressive emergence of an immunosuppressive state 312 could be a mechanism by which hookworms escape the effective anti-helminthic Th2 313 response (Borkow et al., 2000; Maizels et al., 2004). HLA-G can participate in this 314 immunoregulatory state (Carosella et al., 2015) that can lead to a tolerogenic environment. 315

HLA-G is a very complex molecule with well-recognized biological features, and its immune 316 317 functions have recently been reviewed (Carosella et al., 2015). HLA-G is known to inhibit the 318 function of T, NK and B cells through direct interaction with ILT2. Recently, Naji et al., (2014) showed that HLA-G inhibits B cell proliferation, differentiation, and immunoglobulin secretion 319 in both T cell-dependent and T cell-independent models of B cell activation (Naji et al., 320 321 2014). It has been recently shown that children born to mother with high sHLA-G levels are 322 predisposed to have high sHLA-G levels during their two first years of life (d'Almeida et al., 323 2016). We can then hypothesise that the high levels of sHLA-G found in hookworm infected women could be associated with high levels in their newborns that can be involved in the 324 immune tolerance phenomenon. 325

In tropical areas, immune tolerance is not restricted to helminth infections (Raberg, 2014). Placental malaria is classically described as a risk factor for malaria in early infancy (Le Hesran et al., 1997; Le Port et al., 2011; Malhotra et al., 2009; Mutabingwa et al., 2005) and although not totally understood, induction of a tolerant state has also been proposed to be an explanation to this susceptibility. However, we have recently demonstrated that children born

to mothers with placental malaria are not only more susceptible to malaria but also to non-331 malarial fever (Rachas et al., 2012). These findings underline the fact that immune tolerance 332 333 could impact on immunity in a more general manner rather than simply malaria-specific 334 immunity. The involvement of HLA-G in tropical diseases has been recently shown for both malaria and human African trypanosomiasis at both genetic and protein levels (Courtin et al., 335 2013; d'Almeida et al., 2016; Garcia et al., 2013; Gineau et al., 2016; Sadissou et al., 2014). 336 Some associations between antibody responses to malarial antigens and some 337 338 polymorphisms of HLA-G strengthen the concept of a potential role of HLA-G in the control of host immunity in tropical diseases (Sabbagh et al., 2013). 339

Our results also showed a significant increase of sHLA-G levels at delivery compared to the 340 sHLA-G levels quantified during antenatal visits. Similar result was previously observed in a 341 Swedish population (Rizzo et al., 2009). Rizzo and coll. found significantly higher plasmatic 342 sHLA-G level at delivery than during the 3rd trimester of pregnancy in allergic and non-343 allergic pregnant women. Moreover the progression of labour was associated with a 344 345 continuous and significant increase in the sHLA-G level (Knafel et al., 2009). In Africa, the 346 accessibility of health services is often difficult in rural area and most often pregnant women reach the maternity late (Thaddeus and Maine, 1994). In this context, our population study 347 could contain a greater proportion of women with advanced progression of labor. The 348 349 elevation of plasmatic sHLA-G could also be influence by others factors such as stress and 350 hormonal changes induced during delivery.

In conclusion, our results argue for the first time for the potential involvement of HLA-G in the complex phenomenon of helminth-mediated immune tolerance during pregnancy. Investigation of this immune tolerance might help in furthering our understanding of the development of infant immune response.

## 355 Acknowledgements

The authors thank Dr. Adrian Luty for helpful corrections to the manuscript. This study was supported by the Agence Nationale de la Recherche (ANR – TOLIMMUNPAL project) and by

Brazil-France Research Cooperation Program USP/COFECUB (Grant numbers Uc Me 16917). Tania C d'Almeida was partly supported by grants from « Fondation pour la Recherche
Médicale » and Institut de Recherche pour le Développement (IRD). PhD scholarships were
awarded by IRD and Ambassade de France, Cotonou to Ibrahim Sadissou, Rafiou Adamou
and Léonidas Tokplonou.

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Figure 1: Percentage of stool samples with helminth eggs during the pregnancy follow-up

A positive sample was defined as follow: stool sample with detection of helminth eggs by Kato-Katz method. Respectively 395, 387 and 355 stool samples were collected and examined at antenatal visit (AV)1, AV2 and at delivery.

Figure 2: Distribution of the adjusted mean of sHLA-G according to hookworm infection status of the women in the study





Figure 3: Effect of hookworm infection on the adjusted mean of sHLAG controlled for inclusion center (Adjusted coefficients of quantile regression).

\*Quantiles with significant difference between coefficients of infected or uninfected women.

The curves show the regression coefficients on 20 quantiles of sHLA-G distribution during the followup, in women infected or non-infected with hookworms, controlled for the inclusion health facility. <u>The graphic represents the effect of helminth infection on the quantiles of log HLA-G distribution.</u> <u>Significantly more hookworm-infected women were found above the Q80 percentile of sHLA-G</u> <u>distribution.</u>

Variables		N (%)
Inclusion health facility	Attogon Sékou	96 (24) 304 (76)
Marital status	Single Married	9 (2.3) 391 (97.7)
Occupation	Not working Working	223 (55.8) 177 (29.2)
Education	Not educated Educated	283 (70.8) 117 (29.2)
Ethnicity	Fon Aizo Others	83 (20.8) 278 (69.5) 39 (9.7)
Gravidity	Multigravidae Primigravidae	337 (84.3) 63 (15.7)
Placental malaria	No Yes Unknown	353 (88.3) 43 (10.7) 4 (1)
IPTp regimen	SP MQ single dose MQ standard dose	138 (34.5) 142 (35.5) 120 (30)
Household Latrines	No Yes	231 (57.8) 169 (42.2)
Electricity	No Yes	262 (65.5) 138 (34.5)
Refrigerator	No Yes	384 (96) 16 (4)
Television	No Yes	267 (66.7) 133 (33.2)
Motorbike	No Yes	90 (22.5) 310 (77.5)

Table 1: Socio-demographic and economic characteristics of the participants

<u>Table 2</u>: Distribution of hookworm infections according to socio-demographic and economic characteristics of the study population

Variable		Hookworm infection*		
Inclusion		N(%)	p-value	
Health facility	Attogon Sékou	8(8.3) 56(18.4)	0.02	
Marital status	Single Married	2(22.2) 62(15.9)	0.61	
Occupation	Not working Working	41(18.4) 23(13)	0.14	
Education	Not educated Educated	48(17) 16(13.7)	0.41	
Ethnicity	Fon Aizo Others	10(12.1) 53(19.1) 1	0.02	
Gravidity	Multigravidae Primigravidae	54(16) 10(15.9)	0.98	
Placental malaria	No Yes	54(15.3) 8(18.6)	0.57	
IPTp regimen	SP MQ single dose MQ standard dose	17(12.3) 24(16.9) 23(19.2)	0.30	
Household Latrines	No Yes	49(21.2) 15(8.9)	0.001	
Wealth score	<1 ≥1	48(19.2) 16(10.7)	0.024	

\* At least one hookworm infection during the follow-up

<u>Table 3:</u> Effect of hookworm infection on 20 quantiles of sHLAG distribution during the follow-up. Crude coefficients and adjusted coefficients (controlled for inclusion center) of simultaneous quantile regression.

Quantiles	Variable	Crude $\beta^1$	95% CI	P-value	adjusted $\beta^2$	95% CI	P-value
Q10	Hookworm infection	0	-0.01; 0.01	1	0	-0.07; 0.07	1
	Sekou health facility#	0	-0.19; 0.19	1	0	-0.21; 0.21	1
Q15	Hookworm infection	-0.02	-0.31; 0.26	0.87	0	-0.18; 0.18	1
	Sekou health facility	-0.21	-0.54; 0.12	0.21	-0.21	-0.53; 0.11	0.19
Q20	Hookworm infection	-0.36	-0.76; 0.05	0.08	-0.34	-0.73; 0.05	0.09
	Sekou health facility	-0.27	-0.85; 0.31	0.36	-0.21	-0.71; 0.29	0.42
Q25	Hookworm infection	-0.55	-1.22; 0.11	0.1	-0.55	-1.15; 0.05	0.07
	Sekou health facility	-0.3	-0.91; 0.31	0.34	-0.29	-0.79; 0.22	0.27
Q30	Hookworm infection	-0.3	-1.02; 0.42	0.41	-0.2	-0.99; 0.60	0.63
	Sekou health facility	-0.4	-1.14; 0.33	0.28	-0.39	-1.07; 0.29	0.26
Q35	Hookworm infection	-0.18	-1.02; 0.66	0.67	-0.13	-0.97; 0.71	0.76
	Sekou health facility	-0.66	-1.43; 0.11	0.09	-0.69	-1.42; 0.03	0.06
Q40	Hookworm infection	-0.04	-0.93; 0.85	0.93	-0.14	-1.10; 0.81	0.77
	Sekou health facility	-0.72	-1.54; 0.11	0.09	-0.77	-1.54; 0.01	0.05
Q45	Hookworm infection	0.004	-0.87; 0.88	0.99	-0.06	-1.14; 1.01	0.91
	Sekou health facility	-0.62	-1.31; 0.08	0.08	-0.57	-1.30; 0.17	0.13
Q50	Hookworm infection	-0.1	-0.82; 0.62	0.78	-0.06	-1.03; 0.90	0.9
	Sekou health facility	-0.65	-1.28; -0.03	0.04	-0.63	-1.22; -0.04	0.04
Q55	Hookworm infection	-0.09	-0.75; 0.57	0.78	0.06	-0.81; 0.93	0.89
	Sekou health facility	-0.42	-0.83; -0.01	0.05	-0.42	-0.87; 0.02	0.06
Q60	Hookworm infection	-0.003	-0.66; 0.47	0.74	0.09	-0.56; 0.74	0.79
	Sekou health facility	-0.37	-0.68; -0.06	0.02	-0.39	-0.72; -0.06	0.02
Q65	Hookworm infection	0.2	-0.49; 0.47	0.99	0.12	-0.38; 0.62	0.64
	Sekou health facility	-0.3	-0.57; -0.03	0.03	-0.32	-0.62; -0.03	0.03
Q70	Hookworm infection	0.2	-0.22; 0.61	0.35	0.27	-0.13; 0.66	0.19
	Sekou health facility	-0.25	-0.47; -0.03	0.02	-0.26	-0.49; -0.03	0.03

Q75	Hookworm infection	0.19	-0.18; 0.56	0.31	0.26	-0.10; 0.62	0.15
	Sekou health facility	-0.17	-0.42; 0.07	0.17	-0.21	-0.47; 0.04	0.10
Q80	Hookworm infection	0.22	-0.10; 0.54	0.18	0.27	-0.03; 0.58	0.08
	Sekou health facility	-0.23	-0.45; -0.01	0.04	-0.27	-0.51; -0.03	0.03
Q85	Hookworm infection	0.24	-0.02; 0.50	0.07	0.35	0.06; 0.64	0.02
	Sekou health facility	-0.22	-0.43; -0.01	0.04	-0.3	-0.52; -0.08	0.01
Q90	Hookworm infection	0.15	-0.08; 0.38	0.21	0.27	0.01; 0.53	0.04
	Sekou health facility	-0.23	-0.41; -0.06	0.01	-0.31	-0.51; -0.11	0.01
Q95	Hookworm infection	0.17	-0.11; 0.44	0.24	0.31	0.04; 0.58	0.02
	Sekou health facility	-0.13	-0.33; 0.07	0.21	-0.23	-0.41; -0.06	0.01

<sup>1</sup>: From univariate quantile regression

<sup>2</sup>: From multivariate quantile regression

#: Reference: Attogon health facility