

Reviewed Preprint

v1 • April 20, 2026

Not revised

✉ For correspondence:

Matt.Berriman@glasgow.ac.ukgabriel.rinaldi@biology.ox.ac.uk

* co-first authors

Competing interests: No

competing interests declared

Funding: See [page 18](#)

Reviewing editor: Dominique

Soldati-Favre, University of Geneva,
Switzerland

© 2026, Pichon et al. This article is distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use and redistribution provided that the original author and source are credited.

In vitro sexual dimorphism establishment in schistosomes

Remi Pichon^{1,2,*}, Magda E Lotkowska^{3,*}, Jude LD Bulathsinghalage², Madeleine McMath², Mary Evans², Benjamin J Hulme², Kirsty Ambridge³, Geetha Sankaranarayanan³, Simon Kershenbaum¹, Sarah D Davey², Josephine E Forde-Thomas², Karl F Hoffmann², Matthew Berriman^{3,4}✉, Gabriel Rinaldi^{1,2,3}✉

¹Department of Biology, University of Oxford, Oxford, United Kingdom • ²Department of Life Sciences, Aberystwyth University, Aberystwyth, United Kingdom • ³Wellcome Sanger Institute, Hinxton, United Kingdom • ⁴School of Infection and Immunity, College of Medical, Veterinary & Life Sciences, University of Glasgow, Glasgow, United Kingdom

eLife Assessment

This **useful** study presents an improved protocol for long-term *in vitro* culture of *Schistosoma mansoni* that enables progression toward sexually dimorphic stages, representing a meaningful advance for studying parasite development and reducing reliance on animal models. The findings show that host-specific culture conditions support essential developmental and metabolic functions required for parasite maturation, although development remains delayed compared to *in vivo* conditions. The evidence is **solid** overall, but limited pairing efficiency and the absence of egg production indicate that the system does not yet fully recapitulate complete reproductive development.

<https://doi.org/10.7554/eLife.111066.1.sa4>

Abstract

Schistosomes are parasitic flatworms that cause Schistosomiasis, a major Neglected Tropical Disease (NTD) that affects more than 250 million people worldwide. With two morphologically distinct sexes, a heterogametic female (ZW) and a homogametic male (ZZ), schistosomes are an exception among flatworms, which are largely hermaphroditic. Sexual dimorphism in schistosomes only becomes apparent by adulthood within the mammalian host. The cellular and molecular mechanisms underlying the sexual differentiation of these parasites are poorly understood, partly due to intrinsic challenges in assessing parasite development *in vivo*. Therefore, robust and reproducible approaches for maintaining and developing parasites *in vitro* may help to overcome these difficulties. However, to date, only a few studies have focused on protocols that allow cultured parasites to reach sexual dimorphic stages, and none have been reproduced, limiting the ability to understand the unique sexual biology of this major human parasite. Here, we refine a protocol for long-term culture of newly transformed cercariae that developed *in vitro* into sexually dimorphic forms. We assessed the effect of two different sera, Foetal Bovine Serum (FBS) and Human Serum (HS), added to the culture medium supplemented with human red blood cells. We found that in contrast to FBS-cultured worms, HS-cultured parasites digested red blood cells, a crucial step for long term parasite culture. Additionally, while most FBS-cultured parasites did not progress beyond an early liver stage, sexual dimorphism was clearly established in the HS-cultured worms, albeit delayed compared to *in vivo* development. Moreover, EdU pulse-chase experiments revealed a continuous proliferation of parasite cells over time in HS-cultured parasites, while a significantly lower number of proliferating cells were detected in FBS-cultured worms. This protocol paves the way to study parasite development *in vitro*, positively impacting the principles of the 3Rs (Replacement, Reduction and Refinement) for animal research, and allowing for in-depth studies of sexual dimorphism establishment as well as *in vitro* screening for novel control strategies across the life cycle of these major human parasites.

Introduction

Schistosomiasis, a major Neglected Tropical Disease (NTD) affecting >250 million people worldwide, is caused by the infection with blood flukes (class Trematoda) in the genus *Schistosoma*¹. The pathology associated with schistosomiasis is largely driven by egg trapping in different tissues depending on the species. The infection with *Schistosoma mansoni* leads to eggs lodged mainly in the liver inducing inflammation, fibrosis and granuloma formation². This suggests that interfering with the sexual development of these parasites could potentially restrict human pathology. Currently, the complete reliance on a single drug (Praziquantel), to treat the infection and its use in drug mass administration programs in endemic areas threatens the development of drug resistance³. Therefore, identifying novel targets for alternative, more efficient control strategies is a priority that could be addressed by elucidating mechanisms involved in schistosome development, including the establishment of sexual dimorphism.

While most of the parasitic flatworms are hermaphrodites, schistosomes are dioecious with genetically determined female (2n=16, ZW) and male (2n=16, ZZ) individuals. Whilst male and female parasites are morphologically indistinguishable across most developmental stages, sexual dimorphism becomes apparent by adulthood within the mammalian host⁴. Male and female worms undergo separate but concurrent sexual differentiation of their gonads and somatic tissues that eventually allows intersexual pairing, a critical step for female maturation, egg production and life cycle propagation^{5,6}. Transcriptomic studies, at both the bulk^{7–11} and single cell^{12–14} levels, have revealed key pathways and molecular mediators underlying this critical process of female sexual maturation induced by pairing. However, the cellular and molecular mechanisms driving sexual dimorphism prior to pairing remain poorly characterised, partly due to intrinsic challenges in assessing the development of the parasite *in vivo*¹¹.

Approaches for better understand the biology of parasites at large have been developed and optimised using *in vitro* or *ex vivo* refined culture systems^{15,16}. Combinations of culture media components, sera from different sources and additives have been tested^{17,18}. In addition, *organ-on-a-chip* systems and organoid-based 2D and 3D culture platforms^{19,20} have first been implemented for protozoa parasites such as *Plasmodium*, *Toxoplasma*, and *Cryptosporidium* species. These methods have facilitated the molecular dissection of processes underlying single-cell parasite development and interaction with the host^{21–23}. Recently, these technologies have started to be transferred to metazoan parasites to study host-parasite interaction and immune response in the context of helminth infections^{24–28}. *In vitro* cultivation of both larval tapeworms (cestodes) and stem cells isolated from cestodes, coupled with ‘omic’ and imaging analyses have unveiled the role of stem cells in parasite development^{29,30}. However, despite these significant developments in the field, reproducing the natural development of trematodes under controlled *in vitro* conditions remains extremely challenging¹⁶.

Significant progress was achieved in culturing schistosomes when, more than forty years ago, Paul F. Basch developed and optimised comprehensive culture protocols for intra-mammalian developmental stages of schistosomes^{31–33}. Basch reported for the first time the development of male and female parasites from cercariae, with worm pairing and production of infertile eggs entirely *in vitro*³¹. Remarkably, to the best of our knowledge, these results have not been replicated and no further attempt to obtain egg-laying worm pairs developed *in vitro* from cercariae has been reported since Basch’s seminal studies¹⁶. More recently, *in vitro* culture protocols for schistosomes have been refined either to maintain *ex vivo* parasites collected from infected mice to study parasite reproduction biology^{34,35}, or to culture juvenile parasites from cercariae and develop drug screening approaches^{36–39}. However, no long-term culture conditions to specifically assess schistosome sexual differentiation have been reported since 1981³². Despite one research group referring to the approach⁴⁰, there has been no detailed description and no widespread adoption. In fact, published culturing studies have continued to utilise FBS^{34,41}.

Novel functional approaches applied to culture systems that allow reliable and reproducible *in vitro* establishment of sexual dimorphism will shine new light into molecular mechanisms underlying schistosome intra-mammalian development. Here, we refined a protocol for long-term

culture of newly transformed cercariae by assessing the effect of two different sera, Foetal Bovine Serum (FBS) and Human Serum (HS), added to the medium supplemented with human Red Blood Cells (hRBCs). Striking differences were evident between parasites maintained in either FBS or HS. First, hRBC were digested and hemozoin became apparent in the intestines of HS-cultured parasites within 5 days in contrast to FBS-cultured parasites that mostly lacked visible hemozoin. Second, while most of the FBS-cultured parasites did not progress beyond lung and early liver stage, HS-cultured parasites reached sexually dimorphic stages, albeit at a slightly delayed rate compared to *in vivo* development. Furthermore, phenotypic differences between FBS- and HS-cultured parasites became evident as early as 48 hours in culture, with HS-cultured parasites exhibiting higher rates of cell proliferation. Taken together, this protocol allows early sexual development of schistosomes to be studied *in vitro*, provides a method for high throughput drug screening targeting parasite development, and reduces the reliance on mammalian hosts in research.

Results

Sexually dimorphic schistosomes developed entirely *in vitro* from cercariae

Searching for culture media able to cultivate parasites from cercariae to maturity, we decided to test Human Serum (HS) compared to the commonly used Foetal Bovine Serum (FBS). This rationale is based both on the fact that humans are the definitive host of schistosomes, and that two previous reports in 1981 showed that HS was capable of producing mature schistosomes *in vitro*^{31,32}, but they were never replicated. Therefore, to ascertain the *in vitro* sexual dimorphism establishment of schistosomes entirely developed from cercariae, we compared the effect of two different sources of blood serum: FBS and HS. The development of schistosomula derived from mechanically transformed cercariae was assessed in at least 15 independent experiments, five of which were maintained over a period of at least 10 weeks (Figure 1A [↗](#); Supplementary Table S1 [↗](#)).

No morphological differences were observed between parasites cultured either in FBS or HS within the first week in culture; in both conditions most parasites (76% in FBS and 72% in HS in average) were classified as early schistosomula (Supplementary Figure S1 [↗](#)) with few lung and early liver schistosomula (Figure 1B [↗](#), Week 1). The mean mortality at Week 1 was slightly higher, but not statistically significant ($P = 0.8$), in worms cultured in HS ($11.3 \pm 2.7\%$) compared to the mortality registered in FBS-cultured parasites ($5 \pm 2.3\%$), consistent with previous findings³⁸.

From Week 2 onwards, differences in parasite development between the two conditions became apparent (Figure 1B [↗](#)). Parasites cultured in FBS were no longer able to develop beyond early liver stage (14.8% in average), with the vast majority of worms (79% in average) still stunted and classified as early and lung schistosomula. The mortality rate of FBS-cultured parasites continued to increase, reaching an average of $\sim 80\%$ by Week 10, after which the experiments under this condition were stopped as most parasites were dead (Supplementary Figure S2 [↗](#)). On the other hand, parasites cultured in HS further developed over time throughout all categories, achieving marked sexual dimorphism by Week 6 (Figure 1B [↗](#); Supplementary Figure S3A [↗](#)), confirmed by PCR (Supplementary Figure S3B [↗](#); Supplementary Table S2 [↗](#)). As previously described for the *in vivo* development of schistosomes¹¹, the *in vitro* cultured parasite showed developmental asynchrony in agreement with Basch's observations³²; however, by Week 10 the majority of the worms ($73.7 \pm 25.4\%$) have acquired sexual dimorphism (Figure 1B [↗](#)).

The differences in parasite development between the two tested conditions were also confirmed by measuring worm areas at different time points in culture (Figure 2 [↗](#); Supplementary Table S3 [↗](#)). Parasites cultured in the presence of HS grew exponentially over time to reach $\sim 0.1 \text{ mm}^2$, ~ 24 -fold larger than parasites cultured in FBS for 10 weeks, which only grew slightly, plateauing at an average 0.005 mm^2 , similar to the size reached by parasites after only two weeks in HS (Figures 1 [↗](#) and 2 [↗](#)).

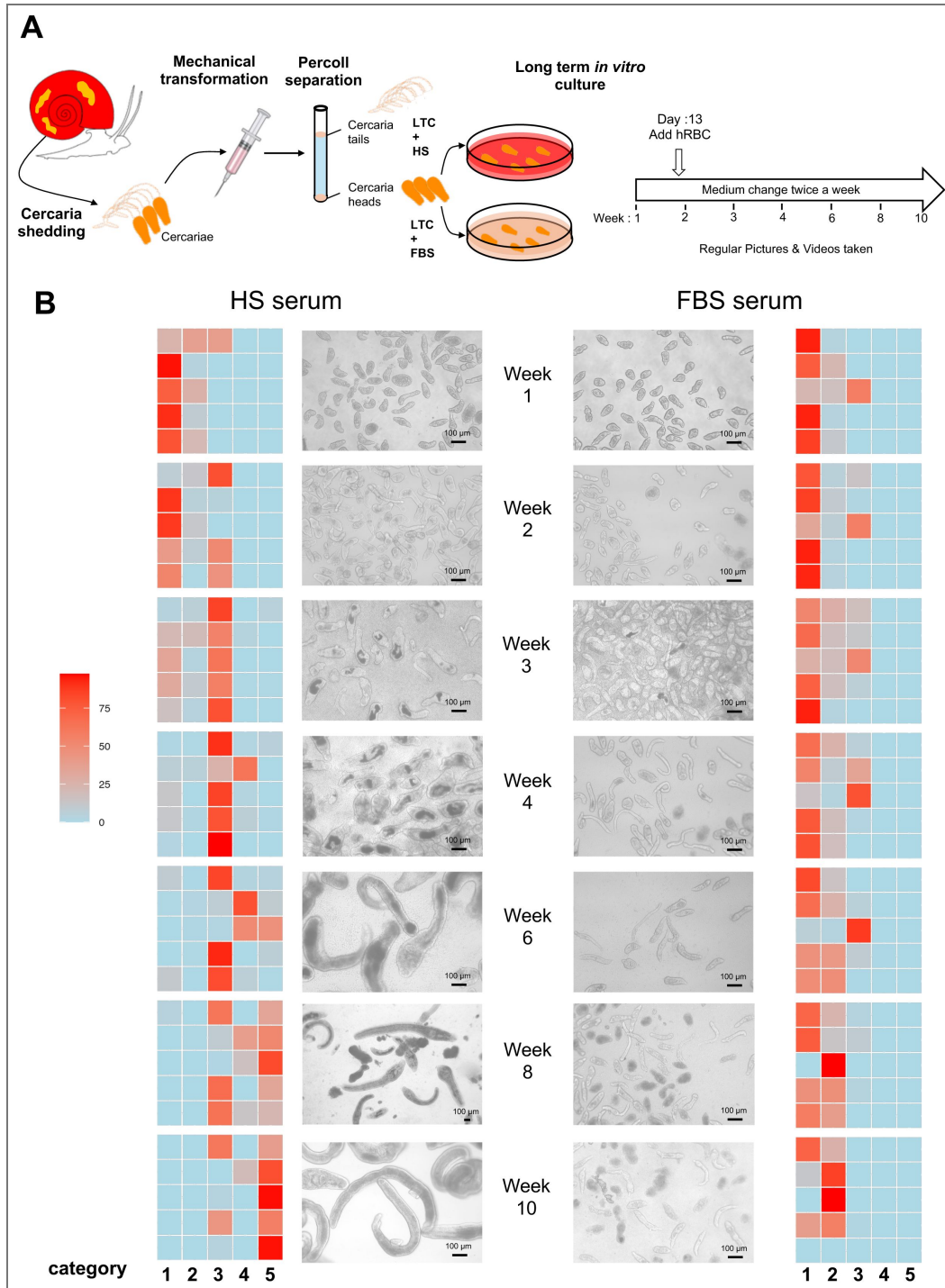


Figure 1. Dimorphic female and male schistosomes entirely developed *in vitro* from cercariae.

A. Schematic representation of the collection and mechanical transformation of cercariae into schistosomula for long-term *in vitro* culture. **B.** Morphological scoring of cultured *S. mansoni* schistosomula at the indicated time points after *in vitro* transformation (week 1 to week 10) for worms in Long Term Culture (LTC) medium supplemented either with Human Serum (HS - left) or Foetal Bovine Serum (FBS - right). Heatmap columns represent five distinct morphological categories, and rows indicate 5 independent culture experiments, i.e. parasites obtained from different batches of infected snails. Heatmap colours represent the percentage of worms for each replicate in each morphological category. Middle panel: Representative images of *in vitro* schistosomula cultured in either HS or FBS as indicated. Scale bars: 100 μm. Category 1: Early schistosomula; Category 2: Lung schistosomula; Category 3: Early liver schistosomula; Category 4: Late liver schistosomula; Category 5: Dimorphic schistosomula.

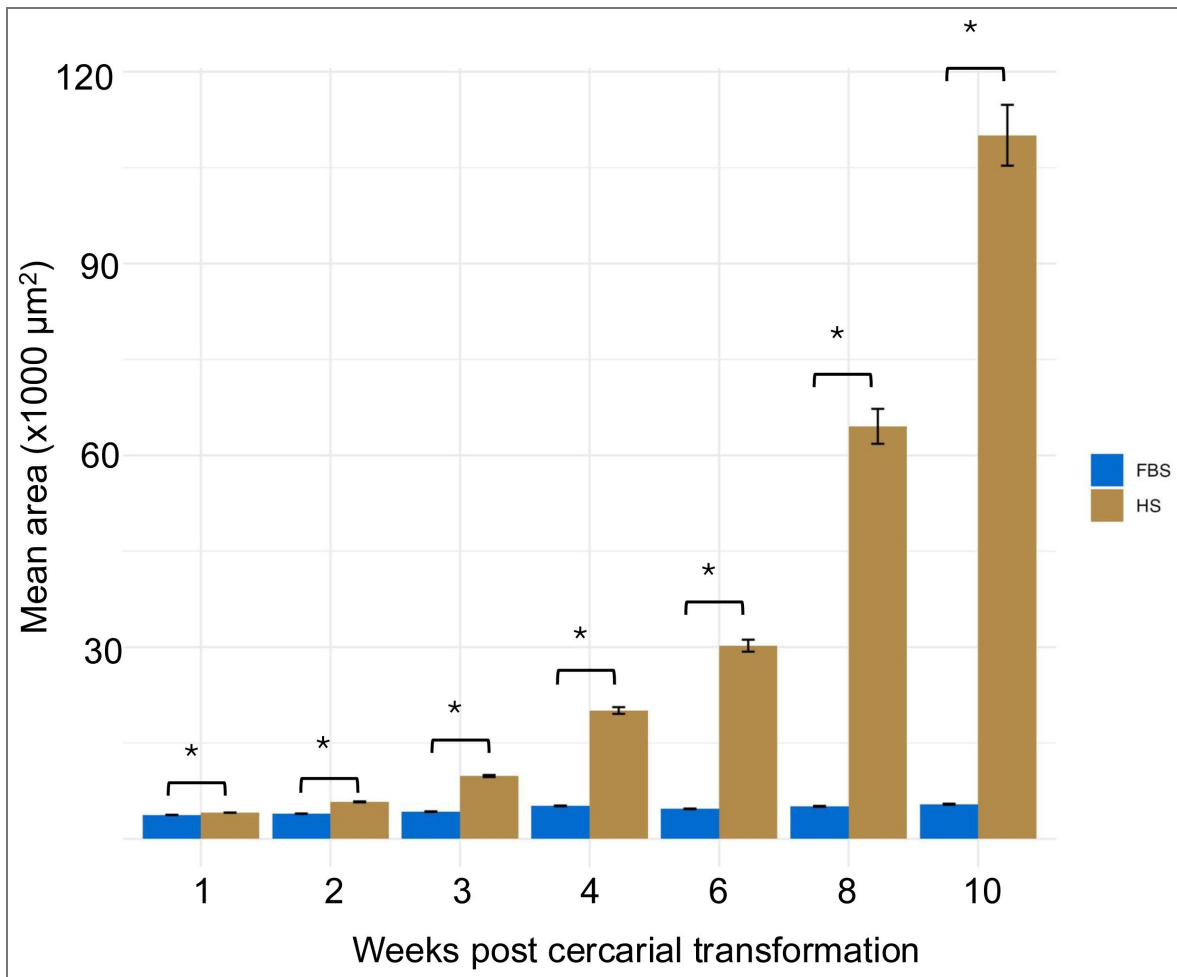


Figure 2. Parasites cultured in Human Serum grew in size unlike those cultured in Foetal Bovine Serum.

Bar plot representing area measurements of schistosomula developed *in vitro* and cultured in media complemented either in Foetal Bovine Serum (FBS) or Human Serum (HS) at indicated weeks after cercarial transformation. Bars indicate mean area (μm^2) \pm SEM. HS: Human Serum (light brown); FBS: Foetal Bovine Serum (blue). *: P-value < 0.05 in indicated pairwise comparisons.

Parasites developed in human serum readily digest red blood cells

It is widely accepted that within the mammalian host schistosomes begin to feed on blood cells through their digestive tract at 10 days post-infection; at this time the majority of the parasites have left the lungs and started to colonise the portal system veins in the liver⁴². On the 13th day of culture, human Red Blood Cells (hRBCs) were added to the parasites to allow them to feed and thus continue their development. At this point, they begin to capture and degrade erythrocytes producing hemozoin, a black pigment derived from host haemoglobin degradation and visible in the worms' intestines. Even though few parasites in FBS reached the early liver stage within the first week in culture, a minority of them were able to digest hRBCs ($3.6 \pm 4.7\%$), indicated by displaying black guts (BG+, Figure 3 [↗](#)). In contrast, a significant proportion of HS-cultured parasites ($36.2 \pm 33.6\%$) had already reached the early liver stage by the second week in 3 out of 5 experiments (Figure 1B [↗](#), Week 2). Moreover, parasites cultured in HS displayed a functional digestive system capable of assimilating hRBCs; more than half of the HS-cultured parasites ($65 \pm 6\%$, P -value < 0.05) were BG+ in comparison to FBS-cultured parasites ($3.7 \pm 4.7\%$) (Figure 3 [↗](#); Supplementary Table S4 [↗](#)).

Development of parasites in human serum may be driven by stem cell proliferation

The growth and development of an organism is driven by a finely regulated combination of cell hyperplasia, hypertrophy, and apoptosis across different tissues⁴³. In schistosomes, a complex stem cell system comprised of both somatic and germline stem cells has been described by leveraging recent single cell transcriptomics⁴⁴. Therefore, we decided to investigate whether the differences in development, growth and feeding capacity between parasites cultured in either FBS- or HS-complemented medium were associated with distinct cellular proliferation rates. EdU pulse-chase experiments revealed notably lower cell proliferation in FBS-cultured parasites as early as 2 days post transformation (Figure 4 [↗](#)). It has previously been demonstrated that a group of 5 well-defined somatic stem cells is the only set of cells that actively proliferate in 2-day schistosomula⁴⁵. Hence, the proliferating cells observed and quantified in our study were most likely stem cells (Supplementary Figure S4 [↗](#); Supplementary Video S1 [↗](#)). The difference between the number of proliferating stem cells in parasites cultured in FBS or HS increased significantly over time. HS-cultured schistosomula showed higher numbers of proliferating stem cells, with an average of >60 EdU+ cells per worm (Figure 4 [↗](#)), whereas most FBS-cultured parasites remained stunted in the lung or early liver stages. These stunted worms displayed no more than an average of 20 EdU+ cells per worm (Figure 4 [↗](#)). hRBCs were added at day 13 post-cercarial transformation to both FBS- and HS-complemented culture media. Worms kept in FBS or HS in the absence of hRBCs were included as controls. Regardless of the serum employed, no significant differences in the numbers of proliferating cells were observed between worms cultured in the presence or absence of hRBCs (Supplementary Tables S5 [↗](#) and S6 [↗](#)).

In vitro cultured schistosomes display sexual dimorphism, developing reproductive systems and pairing capacity

In vitro-developed schistosomes began to show sexually dimorphic features from day ~42 onwards in HS-complemented culture medium (Figure 1 [↗](#), Week 6). Confocal microscopy of DAPI- and Phalloidin-stained *in vitro* cultured male worms confirmed the presence of gynaecophoric canal, developing 3 to 5 testis lobes containing sperm cells, and cirrus (Figure 5A-D [↗](#); Supplementary Video S2 [↗](#)). Likewise, we confirmed the presence of primordial ovaries, oviduct, ootype and uterus in female parasites entirely developed *in vitro* (Figure 5E-H [↗](#); Supplementary Video S3 [↗](#)). In some experiments sexually dimorphic parasites cultured in HS medium were kept alive for more than 150 days (Supplementary Figure S5 [↗](#)). While the establishment of sexual dimorphism was robust and reproducible across more than 15 independent experiments, pairing between male and female parasites was rare. Pairing was observed in ~7% of the experiments, and usually after day ~80 in culture (Figures 6A [↗](#) and 6B [↗](#); Supplementary Video S4 [↗](#)).

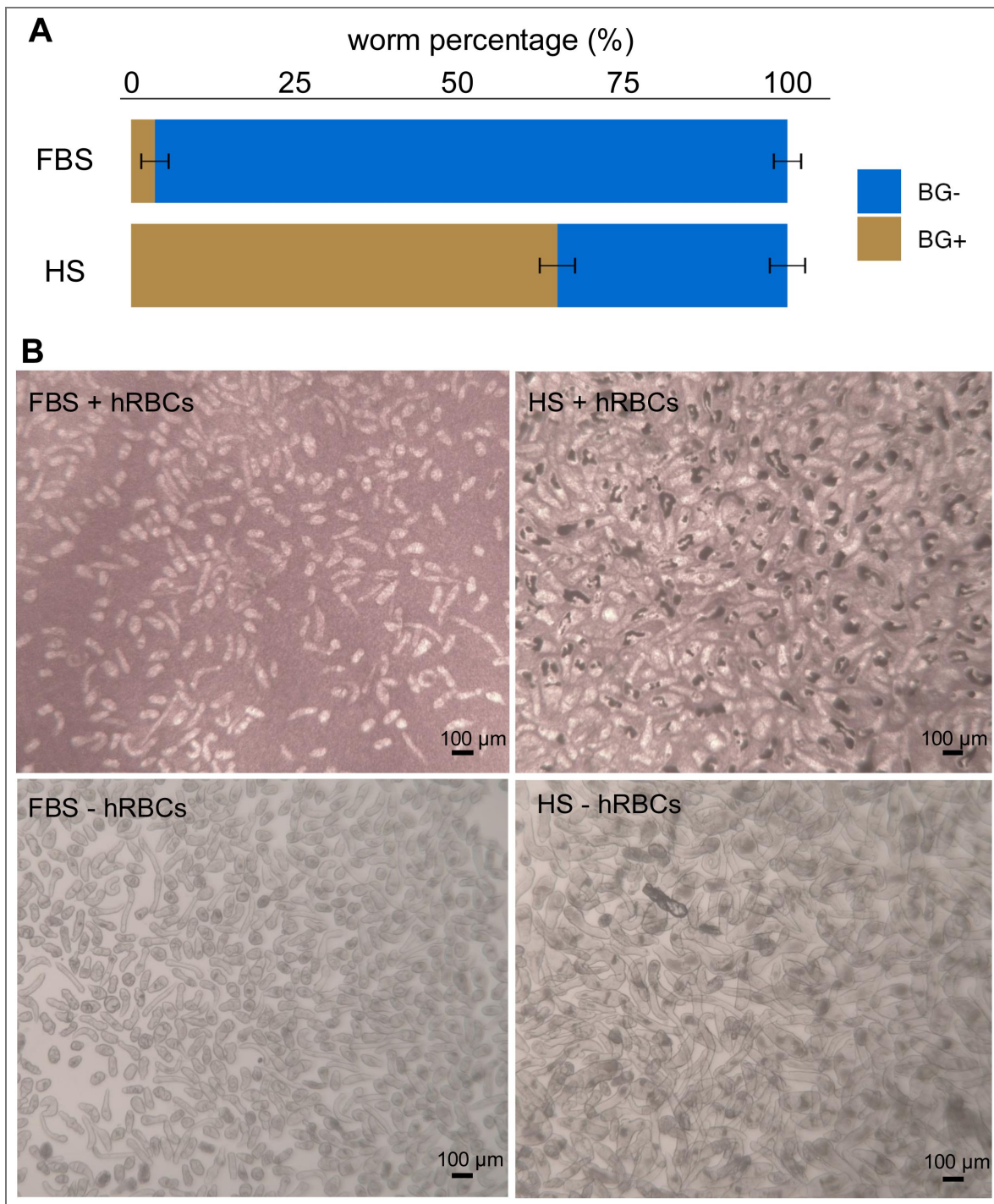


Figure 3. Parasites developed in Human Serum readily digest red blood cells.

A. Bar Plot representing the percentage of Human Serum (HS)- or Foetal Bovine Serum (FBS)-cultured schistosomula with (BG+, light brown) or without (BG-, blue) black guts due to the presence of intestinal hemozoin. Washed human Red Blood Cells (hRBCs) were added into the media at day 13 post-transformation and images captured one or two days later. Error Bars = SEM. **B.** Representative images of *in vitro* developed schistosomula cultured in FBS or HS one day after adding hRBC (+ RBC) and controls without RBC (- RBC). Scale bars: 100 µm.

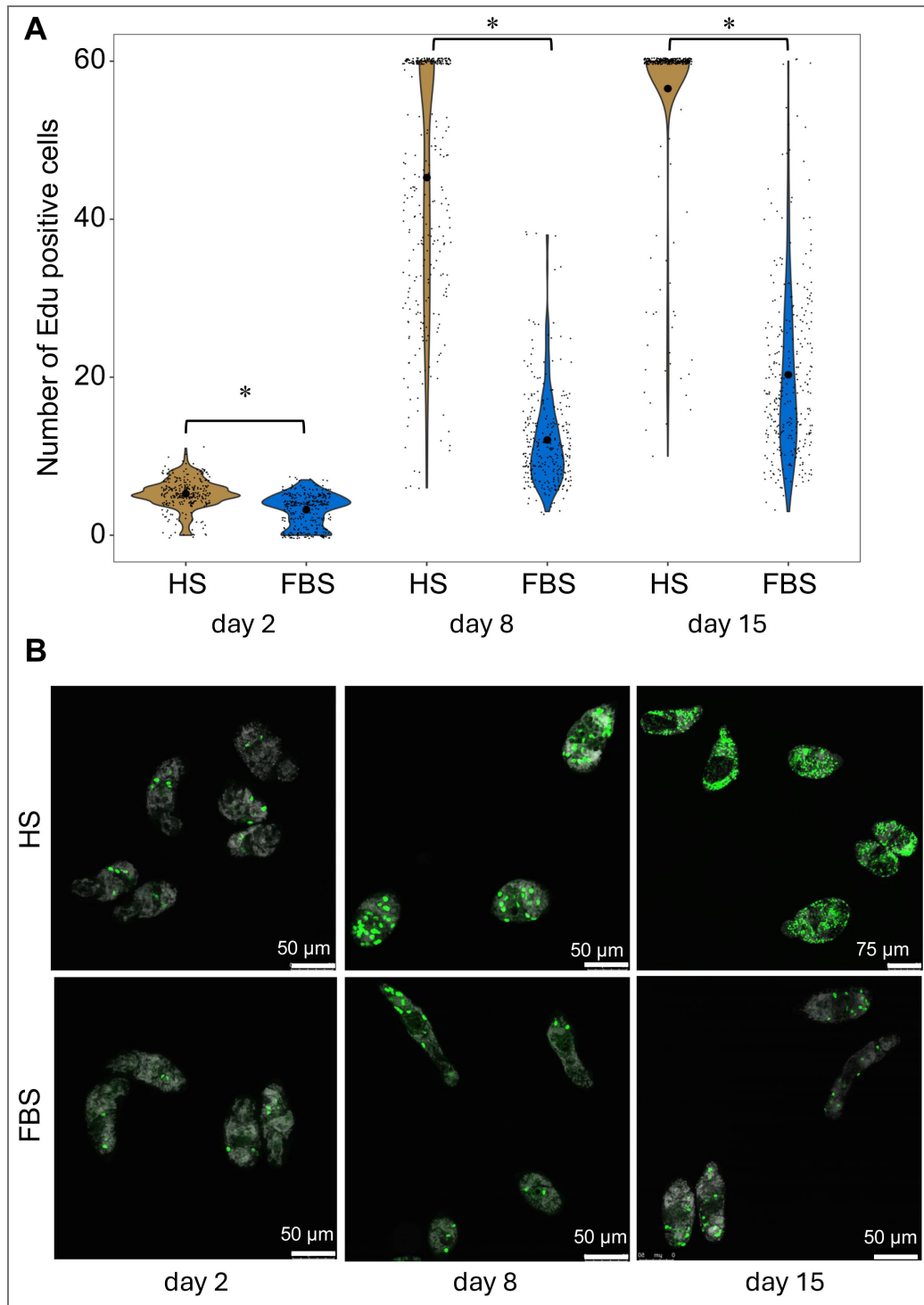


Figure 4. Development of parasites in human serum may be driven by stem cell proliferation.

A. Violin plots showing the number of Edu+ cells per worm at indicated time points (2, 8, and 15 days post-cercarial transformation) in parasites cultured either in Foetal Bovine Serum (FBS, blue) or Human Serum (HS, light brown). Human Red Blood Cells (hRBCs) were added in the culture at day 13 post-cercarial transformation. The small black dots indicate individual worms, and the big black point indicates the mean of Edu+ cells per worm. Statistical analysis was performed by Kruskal-Wallis test with Dunn multiple comparison post-hoc test, with $P \leq 0.05$ (*) considered significant. **B.** Representative images of parasites displaying Edu+ cells at each indicated time point and culture condition. Edu+ cells and nuclei were labelled with Alexa fluor 488 (green) and DAPI (white/grey), respectively. Scale bars: 50 μm or 75 μm as indicated.

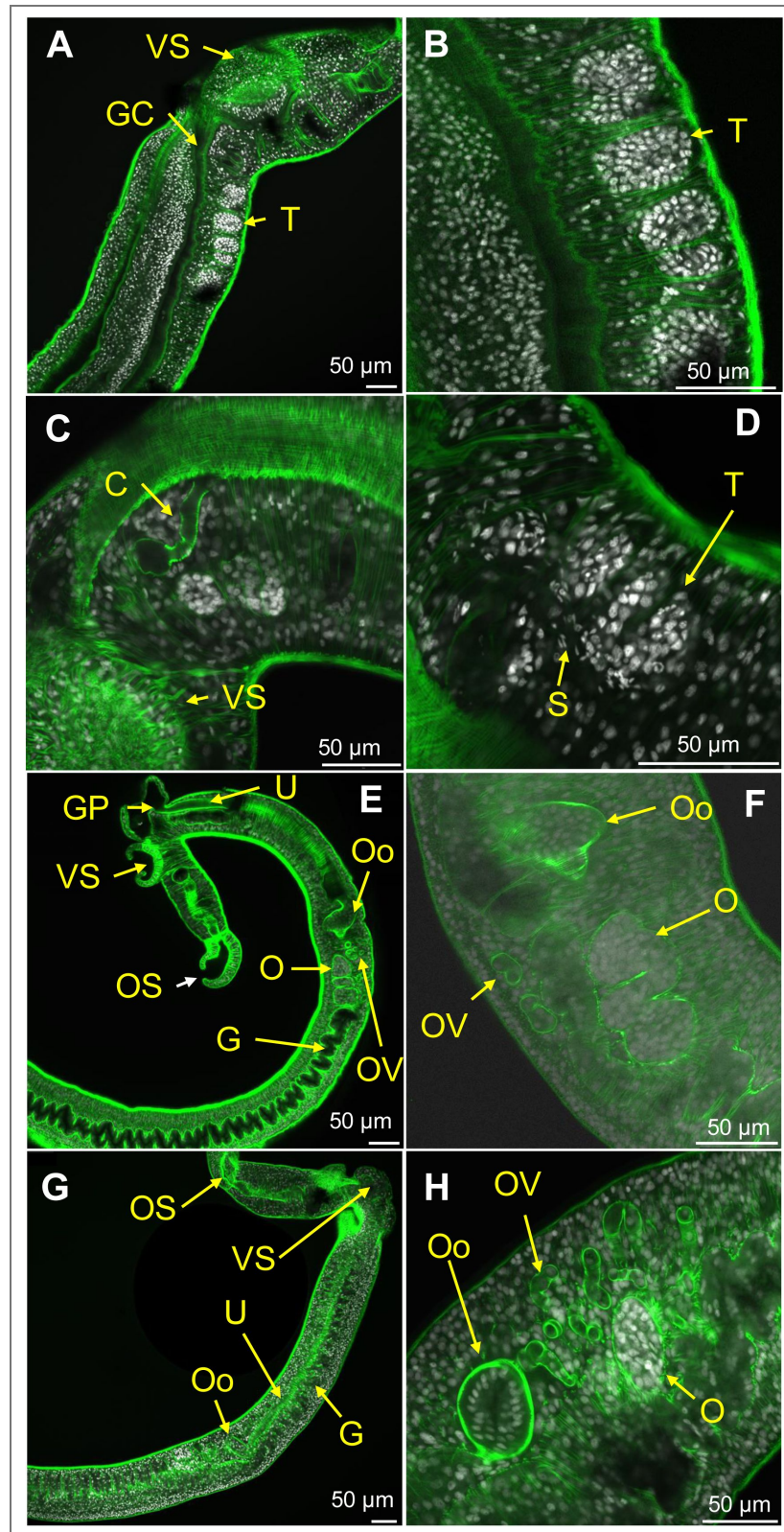


Figure 5. *In vitro* cultured schistosomes display sexual dimorphism and developing reproductive systems.

Representative confocal microscopy images of *in vitro* developed male (A-D) and female (E-H) at day 60 in culture. Worms shown in panels G and H are different individuals. CellMask Green Actin Tracking Stain: green, DAPI: grey (A-E, G), cyan (F, H). C: Cirrus. T: Testis, OS: Oral sucker. VS: Ventral sucker. GC: Gynaecophoric canal. G: Gut. GP: Genital pore. U: Uterus. O: Ovary. Oo: Ootype. OV: Oviduc. S: sperm. Scale bar: 50 μm.

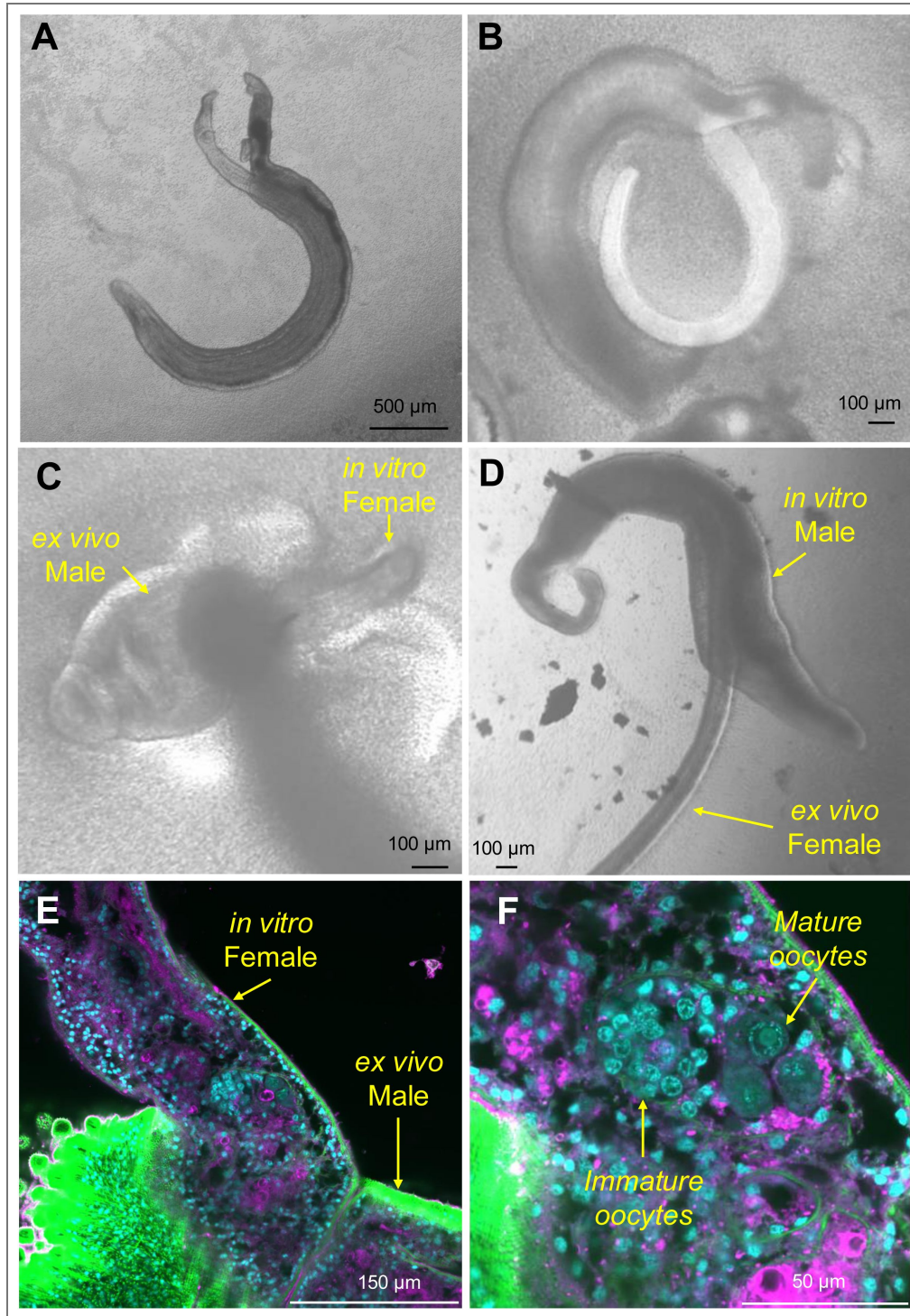


Figure 6. *In vitro* cultured schistosomes are capable of pairing.

A, B: Representative bright-field images of pairs of schistosomes developed entirely *in vitro* after 80 (A) and 150 (B) days of culture in culture medium supplemented with HS. Scale bars: 500 μm (A), 100 μm (B). **C, D:** Representative bright-field images of a worm pairs between *ex vivo*-collected males and *in vitro*-developed females (C) and *ex vivo*-collected females and *in vitro*-developed males (D) within 24 hours after placing the worms in the same well to facilitate pairing. Scale bars: 100 μm. **E, F:** Confocal microscopy image of a schistosome pair (*ex vivo*-collected male and *in vitro*-developed female) *in copula* (E), and magnification of the ovarian area, highlighting maturing oocytes (F). CellMask Green Actin Tracking Stain: green, DAPI: grey, CellMask Deep Red Plasma Membrane Stain: magenta. Scale bar: 150 μm (E), 50 μm (F).

Considering the rarity of the pairing between *in vitro* developed worms, we investigated whether these parasites display the capacity of pairing with *in vivo* developed worms collected from experimentally infected mice. Male and female adult worms developed *in vivo* and recovered from mice by portal perfusion on day 42 post-infection, were sorted by sex and placed in the same tissue culture plate well with worms of the opposite sex developed *in vitro* (>70 days). Within 24 hours after co-culturing the *in vitro* developed with *ex vivo* collected worms, couples were observed (Figure 6C, D [↗](#); Supplementary Video S5 [↗](#)). These findings suggested that the *in vitro* developed and sexually dimorphic parasites are capable of intersexual pairing. Moreover, *in vitro* developed females coupled with *ex vivo* collected mature males displayed signs of primordial ovaries maturation with larger oocytes towards the posterior region of the ovary (Figure 6E, F [↗](#); Supplementary Video S6 [↗](#)). Remarkably, in more than 30 independent *in vitro* culture experiments, where male and female parasites developed sexual dimorphism and eventually paired up, no eggs were produced or laid. This indicates that further refinements in the culture protocol are needed to accelerate the parasite development (the *in vitro* development is delayed compared to the *in vivo* development), increase the likelihood of pairing, and facilitate the production of eggs.

Discussion

Recent progress in culture systems and functional genomic tools is allowing researchers to address long standing questions in helminth biology that were previously inaccessible to experimentation^{46–50}. Among these enduring questions, stand those related to the unique sexual biology of the trematode family Schistosomatidae. Schistosomes are dioecious with genetically determined female and male individuals, which is unusual for flatworms^{4,51}. However, the sexual dimorphism of male and female schistosome worms only becomes established within the mammalian host, and represents a critical step towards worm maturation, intersexual pairing, and egg production⁵². A better understanding of the molecular and cellular basis underlying sexual dimorphism establishment in schistosomes would lead to approaches to block parasite development and life cycle propagation. Studying the sexual development of schistosomes by performing controlled experiments in which parasites can be co-cultured, genetically manipulated or treated with different compounds requires robust and reproducible protocols for long-term *in vitro* culture.

Culture protocols have recently been developed to maintain *ex vivo* schistosomes collected from infected mice^{34,35}, or to obtain juvenile parasites from cercariae for drug screening experiments^{36–39}. However, no long-term culture conditions have been successfully refined to experimentally assess sexual differentiation and dimorphism establishment in schistosomes. Here, aiming to refine a culture medium formulation that supports *in vitro* parasite development and establishment of sexual dimorphism, we decided to compare the effect of modified Basch's media¹⁸, supplemented with human Red Blood Cells (hRBCs) and 20% of either Fetal Bovine Serum (FBS) or Human Serum (HS). While initial parasite survival and development appeared comparable in both conditions during the first week in culture, striking morphological differences emerged from Week 2 onwards. Parasites cultured in HS, progressed through all developmental categories, and acquired sexual dimorphism by Week 6. On the other hand, parasites maintained in FBS were stunted at early stages (mainly lung stage). These experimental outcomes were consistent with the findings reported by Paul F. Basch³¹ in 1980. Probably, from the 1980s onwards a combination of factors that include ethical concerns, variability associated with human-derived products and safety considerations related to blood borne viruses such as HIV, and Hepatitis B and C, determined that parasitologists favoured the use of FBS over HS for helminth culture. FBS has been extensively used in cell culture media, refined and adapted to diverse human and animal cell types since 1958⁵³ providing a reliable source of amino acids, carbohydrates, hormones, lipids, proteins, vitamins, and growth factors⁵⁴. Nevertheless, the scientific community is increasingly advocating for the replacement of FBS as a supplement in tissue culture⁵⁵. This trend is driven by

limitations associated with FBS, including the presence of undefined tentatively harmful factors for the cells in culture, lack of reproducibility, lack of transparency in its production, and critically ethical concerns related to animal welfare⁵⁶.

Our protocol, that supports a fully *in vitro* development of schistosomes, would also positively impact the 3Rs (i.e., Reduction, Replacement, Refinement) for minimising the use of animals for research and serum production^{57,58}.

Parasites cultured in the presence of HS not only developed into sexually dimorphic male and female worms, but strikingly, were able to digest hRBC and process haemoglobin within one day after the addition of hRBC. These findings may reflect differences in the gastrodermis development; worms cultured in the presence of HS may display a differentiated gastrodermis that enables haemoglobin digestion and accumulation of hemozoin within the intestines⁵⁹. On the other hand, most parasites cultured in FBS were unable to produce hemozoin in the presence of hRBCs. This may be due to impaired gastrodermis differentiation that ultimately would lead to stunted development and death. Hematophagous parasites including *Plasmodium* species and schistosomes, obtain key nutrients *via* the proteolysis of host hemoglobin. However, this process leads to the production of free-heme groups which in turn generate highly toxic oxygen free radicals and lipid peroxidation⁶⁰. These toxic free-heme derivatives become inactive when aggregated into an inert crystalline polymer, named hemozoin, observed as dark pigment within the intestines of schistosome intra-mammalian stages⁵⁹. In addition, increasing evidence shows that hemozoin may play critical roles during parasite development by supplying iron for egg production⁵⁹, and interaction with the mammalian host by an immune modulatory role⁶¹. In the mouse model, schistosomes begin to feed on blood once they have left the lungs and reached the portal system 9 to 11 days post infection^{42,62}. Moreover, schistosomes collected from experimentally infected mice at day 13 post infection already show hemozoin in their guts¹¹. Similarly, HS-developed schistosomes were able to feed on blood *in vitro* within a comparable time window; hRBCs were added in the medium on day 13 in culture, and within 24 hours most of the parasites' gut contained hemozoin. These findings suggest that our *in vitro* culture system successfully recapitulates *in vivo* gastrodermis development of schistosomes. A better understanding of the *in vitro* development of the parasite gastrodermis and its role in the production of hemozoin will expose tentative novel targets for control⁶³.

Although previous reports have shown that it is possible to cultivate parasites in cell-free environments^{38,64}, adding hRBC has proven essential for long-term maintenance of parasites and for establishing sexual dimorphism. EdU pulse-chase experiments suggested that schistosome *in vitro* development in the presence of HS may be driven by the proliferation and differentiation of stem cells. The stem cell system in schistosomes has been extensively studied since the first description of somatic stem cells in adult worms⁶⁵, followed by single cell transcriptomic identification and functional characterisation of three key stem cell populations in intra-snail stages⁴⁵. Two of these stem cell types, maintained in intra-mammalian stages, proliferate and differentiate into precursors of somatic and germ line cells throughout development¹³. Although we have not performed transcriptomic analyses in this study, the positive correlation between number of EdU+ cells, and growth (determined by increasing worms' area) and gross development in HS-developed parasites was evident already within the first 48 hours in culture. These findings indicate that stem cells may play a central role in driving organised growth, tissue differentiation and ultimately the establishment of the sexual dimorphism in HS-developed parasites. Conversely, FBS-cultured parasites displayed no more than 20 EdU+ cells per worm in average, reaching a plateau in the number of proliferating cells from day 8 in culture, which probably underlie the arrested development of these worms. Overall, the FBS-cultured parasites did not progress beyond the lung stage. Consistent with these findings, confocal microscopy imaging revealed in HS-developed parasites clear somatic and reproductive anatomical structures, including male gynaecophoric canals and testes, as well as female ovary primordia, oviducts, ootypes and uteri. These developmental features illustrate the capacity of our culture system to capture key biological transitions previously only accessible from experimentally infected animal models. That said, while our system was highly efficient in producing sexually dimorphic worms, spontaneous

pairing between male and female parasites was extremely rare, mainly in aged *in vitro* cultures (from 80 to 100 days in culture) indicating that other factors, e.g., cholesterol, may be missing³⁴. In any case, we decided to test the pairing capacity of the dimorphic worms by conducting pairing experiments between *in vitro*-developed females and *ex vivo*-collected males, and vice versa. Strikingly, couples were observed within 24 hours after co-cultivating *in vivo* and *ex vivo* opposite sex parasites, albeit no eggs were produced. However, we observed evidence of pairing-trigger oocyte maturation in *in vitro*-developed females paired with *ex vivo*-collected male worms. Our research group has recently been involved in the discovery and functional characterisation of a transcription factor of the retinoic acid receptor family, *SmRAR*, and related genes⁶⁶. *SmRAR* and associated genes may play a key role in oocyte differentiation triggered by pairing. Our pairing experiments and confocal imaging analyses suggested that *in vitro* developed female oocytes started to differentiate after pairing with an *in vivo* developed male worm, probably mediated the activation of the *SmRAR* pathway⁶⁶. Moreover, the involvement of a male-derived nonribosomal peptide pheromone, i.e., β -alanyl-tryptamine or 'BATT', in the pairing-driven female sexual maturation was demonstrated⁵. In addition, soluble factors produced by the worms with effect on the opposite sex cannot be ruled out⁶. It has been suggested that host-derived factors, such as the cytokine transforming growth factor β (TGF- β) may be critical for female reproductive development and embryogenesis⁶⁷. More recently, other host-derived factors that include cholesterol and ascorbic acid have been shown to be critical for maintaining *ex vivo* fully mature females collected from infected mice³⁴. Further research is needed to assess the role of these host- and parasite-derived cues in schistosome development.

In summary, we have demonstrated that the presence of HS is essential to support fully *in vitro* development of *S. mansoni* parasites from mechanically transformed cercariae to sexually dimorphic adults. Differences in the numbers of proliferating stem cells between worms developed in HS versus FBS were observed as early as 48 hours in culture. These findings may raise some concerns about studies that rely on *in vitro* culture protocols using FBS, including those focused on parasite developmental biology, interaction with the host, and drug screening. Human serum contains essential host-specific molecules that are absent or insufficient in FBS, and future efforts should aim to identify these factors. Optimising schistosome long-term culture systems will: (1) deepen our understanding of fundamental aspects of schistosome biology, development and host-parasite molecular crosstalk; (2) positively impact the 3Rs principles for animal research; and (3) reveal tentative targets for novel control strategies.

Material and Methods

Ethics statement

The complete life cycle of *Schistosoma mansoni* (NMRI strain) was maintained at the Wellcome Sanger Institute (WSI) and Aberystwyth University (AU), by infecting *Biomphalaria glabrata* snails and outbred mice (TO strain). All the animal regulated procedures were conducted under Home Office Project Licences P77E8A062 (WSI) and PP2955700 (AU), following the ARRIVE guidelines (<https://arriveguidelines.org>) and in accordance with guidelines and regulations stated by the UK Animals (Scientific Procedures) Act 1986 Amendment Regulations 2012. All the protocols were presented and approved by the Animal Welfare and Ethical Review Bodies (AWERB) of the WSI and AU. The AWERB is constituted as required by the UK Animals (Scientific Procedures) Act 1986 Amendment Regulations 2012. Human blood products, including serum and red blood cells were obtained from NHS Blood and Transplant (NHSBT), Non-Clinical Issue (NCI) services for research purposes. NHSBT-NCI provides donated material surplus to clinical requirements or unsuitable for therapeutic use that has been appropriately consented. The supply chain complies with all statutory and regulatory obligations including (but not limited to) the Human Tissue Act (2004) and associated Codes of Practice. The NHSBT-NCI products were acquired under the Study 'In vitro development of the human parasite *Schistosoma*', Integrated Research Application System (IRAS) project ID 319400, protocol number AU/DLS/011, Research Ethics Committee (REC) reference number 23/SS/0017.

Cercariae transformation and culture of schistosomula

Schistosoma mansoni (NMRI strain) schistosomula were obtained and cultured as previously described with minor modifications¹⁸. In brief, patent mix strain of *Biomphalaria glabrata* snails⁶⁸ experimentally infected *en masse* with ~20 miracidia per snail, were thoroughly rinsed, transferred to Lepple water (~100 ml) and exposed to light for 2 h at 26 °C to induce cercarial shedding. Cercariae were collected, passed through a 100 µm filter into 50 ml tubes to remove snail debris and faeces, placed 1 h on ice, and concentrated by centrifugation (300 g for 3 min at 4°C with half break deceleration). Cercariae were washed 3 times in 1X PBS supplemented with 200 U/ml penicillin, 200 µg/ml streptomycin, 500 ng/ml amphotericin B (Merck). During the washes, the cercarial pellets were successively combined and finally resuspended in 'Schistosomula wash medium' (Dulbecco's modified Eagle medium - DMEM, supplemented with 10 mM Hepes, 200 U/ml penicillin, 200 µg/ml streptomycin, 500 ng/ml amphotericin B). Cercariae were vortexed full speed for 30 sec and placed on ice for 1 min. Thereafter, the cercarial tails were sheared off by ~10 back and forth passes through a 19-gauge (19-G) needle, a ~3 µl aliquot of parasites inspected under microscope to confirm that >90% of cercariae had their tail removed, and the cercarial heads were separated from the tails by a Percoll (Merck) gradient (1.5:1 percoll:DMEM) and centrifugation (300g for 15 min at 4 °C, half break deceleration). The pellet containing purified cercarial heads, was collected and washed 3 times by centrifugation (300g for 3 min at 4°C) in 'Schistosomula wash medium'. Newly transformed schistosomula were counted in 12 5µl-aliquots under the microscope. To reduce parasite mortality due to a high density of worms, ~8,000 schistosomula were transferred to each well of a 6-well tissue culture plate (Fisher Scientific, Loughborough, UK) containing 5 ml of modified Basch's medium¹⁸ supplemented with additives and FBS or HS heat-inactivated FBS or HS serum, i.e., 'Long-term culture medium' (LTC) as indicated in Table 1 [↗](#). Parasites were cultured in a tissue culture incubator at 37°C under 5% CO₂ in air. Washed human red blood cells (hRBCs) were added into the culture medium at indicated time points. LTC medium was replaced twice a week and washed hRBCs added to a final concentration of 0.02% v/v at indicated days after transformation. All human blood products, obtained from NHSBT-NCI, were handled with universal precautions within a biosafety cabinet class II and suitable PPE. The HS was heat-inactivated at 56°C for 30 min in an incubator, aliquoted and stored at -20°C until use. The hRBC were washed in 'Schistosomula wash medium' and stored at 4°C until use as previously described¹⁸. Briefly, the hRBCs were transferred from the pack to 50 ml tubes and washed by centrifugation at 500g for 5 min at 4°C. Half the supernatant was removed, tubes' content combined, centrifuged as above and resuspended in ~25 ml of 'Schistosomula wash medium'. The washes were repeated 4 more times.

Phenotype scoring of developing schistosomula

To analyse the *in vitro* development of schistosomula, images and videos were taken at regular intervals of at least once a week in >10 independent culture experiments, using a digital Euromex camera connected to an Olympus CK2 optical microscope. Up to 18 pictures per experiment with an average of ~400 worms per picture were used for parasite staging from at least 5 independent experiments (Supplementary Table S1 [↗](#)). The investigators who assigned categories were blinded to experiment conditions. Based on a well-defined staging system^{11,32}, the parasites were classified into 6 developmental categories as described³⁸ (Supplementary Figure S1 [↗](#)); category 0: *dead parasites* showing granulation, rounded shape and degraded or broken tegument - the number of dead parasites may be underestimated due to the loss of some dead worms during the media change; category 1: *newly transformed schistosomula*, corresponding to the *in vivo* 'skin stage', these worms retain the cercarial head shape with no visible internal structures such as gut; category 2: *lung stage schistosomula*, elongated and slim worms, usually with a bulged end; category 3: *early liver stage*, bigger and wider worms compared to the previous stage, with hemozoin pigment (i.e., haemoglobin degradation product) and two ceca gut; category 4: *late liver stage*, in which the gut is further developed, the two intestinal ceca have fused behind the ventral sucker, and the parasites started to acquire an evident vermiform shape; category 5: *sexually dimorphic stage*, comprising large, vermiform male- and female-looking worms. The females are

Reagent name	Final Concentration	Reagent source
DMEM, high glucose, sodium pyruvate		Fisher Scientific (Cat. 13476146)
Lactalbumin hydrolysate	1 mg/ml	Merk Life Sciences (Cat. 61300-500G)
Hypoxanthine	500nM	Merk Life Sciences (Cat. H9636-1G)
Serotonin	1 μM	Merk Life Sciences (Cat. H9523)
Hydrocortisone	1 μM	Merk Life Sciences (Cat. H0888-1G)
Triiodothyronine	200nM	Merk Life Sciences (Cat. T6397-100MG)
MEM vitamins	1X	Merk Life Sciences (Cat. M6895-100mL)
Schneider's insect media	10%	Merk Life Sciences (Cat. 50146-500mL)
Hepes	20mM	Merk Life Sciences (Cat. H0887-100mL)
Heat inactivated Human serum	20%	HS: NHSBT serum (Cat. NC02)
Foetal Bovine Serum		FBS: Fisher Scientific (Cat. 11550356)
Antibiotic antimycotic solution	2X	Fisher Scientific (Cat. 15140-122)
Insulin solution	8 μg/ml	Merk Life Sciences (Cat. I9278-5mL)
L-glutamine	2mM	Fisher Scientific (Cat. 11539876)
Human red Blood Cells	0.02% v/v	NHSBT-NCI (Cat. NC15)

Table 1. Long term culture medium (LTC medium) composition.

longer and thinner than males, the ceca fusion localises closer to the anterior end of the worm (i.e., $\sim\frac{1}{3}$ of the worm length from the anterior end), and both suckers are smaller compared to the male suckers. The male worms are characterised by a larger and wider head, larger suckers, the ceca fusion closer to the posterior end of the animal (i.e., $\sim\frac{2}{3}$ of the worm length from the anterior end), and a clearly visible gynecophoral groove.^{11,32}

To quantify positive or negative parasites for the presence of hemozoin within their intestines, i.e., black guts + (BG+) or BG- parasites, respectively, hRBCs were added to the culture at day 13, and one or two days later bright field microscopy images were taken. Counts of BG+ and BG- parasites were collected from 5 independent *in vitro* culture experiments. Statistical analysis was performed by ANOVA followed by Tuckey's post-hoc multiple pairwise comparison with $P \leq 0.05$ considered significant (Supplementary Table S6 [↗](#)).

Measurement of parasite area

Images of *in vitro* developing parasites were taken at weeks 1, 2, 3, 4, 6, 8 and 10 after cercariae transformation with an Euromex camera fitted to an Olympus CK2 optical microscope at 4x and 10x magnification. For these magnifications images of scale bars were captured for pixel to area values (μm^2). Parasites from 5 independent long-term culture experiments were masked in COCO format from captured images using the CVAT (Computer Vision Annotation Tool) online server (<https://www.cvat.ai/> [↗](#)). Area metrics were then calculated for each parasite by leveraging the pycocotools package in a customised Python script. Raw and processed data is provided in Supplementary Table S3 [↗](#). Normal distribution was checked using Shapiro test and Mann Whitney tests performed with a significant value set up for $P \leq 0.05$ (Supplementary Table S6 [↗](#)).

PCR for sexing and confirmation of sexual dimorphism

To confirm the *in vitro* establishment of sexual dimorphism, parasites were individually collected from culture by day ~ 80 , for blind sex assignment by bright field microscopy and confirmation by sex-specific PCR⁶⁹ (Supplementary Figure S3B [↗](#)). Briefly, pictures from individual *in vitro* developed parasites were taken and sex assigned based on morphological features. Thereafter, genomic DNA from the sex-assigned individual parasites was extracted by lysing the worms at 95°C for 1h in 50mM NaOH, 0.4mM disodium ethylenediaminetetraacetic acid (EDTA), followed by the addition of neutralising buffer (80 mM Tris-HCl in water). Samples were stored at 4°C until use. PCR reactions were performed in a final volume of 10 μl consisting of 5 μl Platinum 2X hot start PCR master mix (Invitrogen, California, USA), 0.5 μl (10 μM stock) of forward and reverse primers targeting W1 repeat⁷⁰ and actin⁷¹ (Supplementary Table S2 [↗](#)), 1 μl of template DNA and 3 μl of water. The PCR program comprised an initial denature step at 94°C for 2 min followed by 30 cycles of denature at 98°C for 5 secs, and annealing/elongation at 60°C for 15 secs. The PCR products were resolved on a 1% agarose gel using 1% TAE buffer and 1kb GeneRuler DNA ladder (Thermo Scientific, UK).

Detection and quantification of proliferating cells

Parasites cultured for 15 days (D15) in media supplemented with either HS or FBS and hRBC added at D13 were harvested at days 2, 8 or 15 after cercariae transformation. Parasites cultured under the same conditions but with no hRBC were included as controls. The day before each indicated time point, a solution of 5-ethynyl-2'-deoxyuridine (EdU) was added to the culture medium at 10 μM final concentration. At each indicated time point, parasites from each group were collected and transferred to a 35 μm mesh basket (CEM Microwave Tech) in a well of a 24-well plate, washed in 1x PBS, fixed in 4% Paraformaldehyde (PFA) in 1x PBS for 30 min, washed in 1x PBS and stored at 4°C until use. The worms were incubated in PBST (1x PBS + 0.3% Triton x-100) containing 2 μl of Proteinase K (20 mg/ml) for 5 min at room temperature, washed three times in PBST, fixed in 4% PFA in 1x PBS for 10 min at room temperature and washed 5 times in PBST. EdU-positive cells were revealed by incubating the parasites for 30 min in the dark in a solution containing Azide fluor 488 (Sigma-Aldrich), diluted in a solution of 1 mM CuSo4, 0.025 mM Azide fluor 488, 95 mM L-Ascorbic Acid in 1x PBS. The worms were washed 5 times in PBST and incubated in mounting media

containing DAPI (Fluoromount-G™ Mounting Medium, with DAPI, Invitrogen) before mounting for confocal microscopy (Leica SP8 super resolution laser confocal microscope). EdU+ cells per parasite were counted for an average of 100 parasites across three independent experiments. Statistical analysis was performed by Kruskal-Wallis test with Dunn multiple comparison post-hoc test, with $P \leq 0.05$ considered significant (Supplementary Table S6 [↗](#)).

Pairing experiments

Schistosoma mansoni adult male and female worms were collected from experimentally infected mice at 47 days post infection¹⁸. Briefly, mice were euthanised by intraperitoneal injection of 200 μ l of 200 mg/ml pentobarbital supplemented with 100 U/ml heparin, and worms collected by portal perfusion (the hepatic portal vein was sectioned followed by intracardiac perfusion with phenol-red-free DMEM, containing 10 U/mL heparin) and washed in DMEM. In 24-well plates, *in vitro* developed female worms were cultured in the presence of *in vivo* developed male worms and vice versa at a ratio of 1 female every 2 males in the well containing culture medium (Table 1 [↗](#)) supplemented with 20% HS. The parasites were cultured for several days in an incubator at 37°C, 5 % CO₂ and checked daily under microscope for the presence of pairing.

Parasite staining and confocal imaging

Live parasites collected from culture at indicated time points were stained with 1:1000 dilutions of CellMask™ Green Actin Tracking Stain (Invitrogen) and CellMask™ Deep Red Plasma Membrane Stain (Invitrogen) to label polymerized/filamentous actin (F-actin) and cell membranes, respectively. Nuclei were stained by adding two drops per millilitre of NucBlue™ Live ReadyProbes™ Reagent (Invitrogen). After an overnight incubation at 37°C, worms were collected in 15 ml tubes, washed by gravity 3 times in 1x PBS, fixed in 4% PFA (in 1x PBS) overnight at 4°C, washed 3 times in 1x PBS and stored in 200 μ l of Fluoromount-G, with DAPI (Invitrogen) before mounting on slides for confocal microscopy. All the images were acquired using a Leica SP8 super resolution laser confocal microscope.

Data availability

All raw data and statistical analyses provided in Supplementary Tables.

Acknowledgements

The authors thank Julie Hirst for technical assistance with the *Schistosoma mansoni* life cycle maintenance at Aberystwyth University, and Anais Bordes for technical support with the *Schistosoma mansoni* life cycle maintenance at University of Oxford. The authors also gratefully acknowledge the Biology Dept Imaging Suite, University of Oxford for their support & assistance in this work (Funder: EPA Cephalosporin Fund and Department of Biology, Project Number CBR00830/REF CF 401). The authors acknowledge NHS Blood and Transplant (NHSBT), Non-Clinical Issue (NCI) for providing human blood products (detailed information in Methods). The study was partially funded by the Wellcome Trust (grants 098051 and 206194), and the European Union [Project 101080784 – WORMVACS2.0]. GR is supported by UKRI Future Leaders Fellowships [MR/W013568/2].

Additional information

Author contributions

Remi Pichon, Conceptualization, Methodology, Investigation, Data curation, Formal analysis, Visualization, Writing – original draft, Writing – review and editing; Magda E Lotkowska, Conceptualization, Methodology, Investigation, Data curation, Formal analysis, Visualization; Jude L. D. Bulathsinghalage, Methodology, Investigation, Data curation; Madeleine McMath, Methodology, Investigation, Data curation, Formal analysis, Visualization; Mary Evans, Methodology, Investigation; Benjamin J. Hulme, Methodology, Investigation; Kirsty Ambridge,

Methodology, Investigation, Visualization; Geetha Sankaranarayanan, Methodology, Investigation; Simon Kershenbaum, Formal analysis, Writing – original draft, Writing – review and editing; Sarah D. Davey, Methodology, Investigation, Data curation, Formal analysis, Visualization; Josephine E. Forde-Thomas, Methodology, Investigation, Visualization; Karl F. Hoffmann, Conceptualization, Supervision, Funding acquisition; Matthew Berriman, Supervision, Funding acquisition, Project administration; Gabriel Rinaldi, Conceptualization, Methodology, Investigation, Data curation, Formal analysis, Visualization, Supervision, Funding acquisition, Project administration, Writing – original draft, Writing – review and editing.

Funding

Funder	Grant reference number	Author
Wellcome Trust (WT)	https://doi.org/10.35802/098051	Matthew Berriman
Wellcome Trust (WT)	https://doi.org/10.35802/206194	Matthew Berriman
European Union	https://doi.org/10.3030/101080784	Karl F Hoffmann
UKRI- Future Leaders Fellowship	MR/W013568/2	Gabriel Rinaldi

Author ORCID iDs

Remi Pichon: <https://orcid.org/0000-0002-5821-9529>

Mary Evans: <https://orcid.org/0009-0001-9120-6017>

Benjamin J Hulme: <https://orcid.org/0000-0002-5638-6323>

Geetha Sankaranarayanan: <https://orcid.org/0000-0001-8303-2451>

Simon Kershenbaum: <https://orcid.org/0009-0000-1797-1282>

Sarah D Davey: <https://orcid.org/0000-0001-8676-7330>

Josephine E Forde-Thomas: <https://orcid.org/0000-0002-7202-3565>

Karl F Hoffmann: <https://orcid.org/0000-0002-3932-5502>

Matthew Berriman: <https://orcid.org/0000-0002-9581-0377>

Gabriel Rinaldi: <https://orcid.org/0000-0002-7767-4922>

Additional files

Supplementary Figures [Supplementary Figure S1](#). [Developmental staging scoring](#). Summary of morphological attributes for the five different categories of development of worms cultivated *in vitro* (HS), as well as representative bright field images of each of these morphological categories. Arrows in categories 2 and 4 indicate elongated lung schistosomula, and the fusion of the two ceca posterior to the ventral sucker in the late liver schistosomula, respectively. Scale bar: 100 μ m. [Supplementary Figure S2](#). [Representative FBS-cultured parasites by Week 10 in culture](#); most of the worms are dead (red arrows) and the few still alive are lung schistosomula (green arrows). Scale bar: 100 μ m. [Supplementary Figure S3](#). [A. Representative pictures of *in vitro* developed parasites by day ~80 in culture, male or female worms as indicated. Scale bar: 500 \$\mu\$ m B. Representative results of the PCR-based sex genotyping using primers to amplify a fragment of the control actin gene in both male and female \(upper band\), and primers to amplify a W-specific region only in female worms \(lower band\). A non-template control, and male and female positive controls were included as indicated. Assigned sexes for each of the groups of clonal cercariae emitted by individual snails infected with single miracidia. \[Supplementary Figure S4\]\(#\). \[Representative confocal high magnification images of individual parasites displaying Edu+ cells at each indicated time point and culture condition. Edu+ cells and nuclei were labelled with Alexa fluor 488 \\(green\\) and DAPI \\(white/grey\\), respectively. Scale bars: 50 \\$\mu\\$ m. \\[Supplementary Figure S5\\]\\(#\\). \\[Representative bright-light pictures parasites cultured in HS supplemented medium for more than 100 days, indicating male and female worms. A-F: parasites cultured for 103 days. G, H. Parasites cultured for 145 days. Scale bar: 100 \\\$\mu\\\$ m.\\]\\(#\\)\]\(#\)](#)

Supplementary Tables [Supplementary Table S1](#). [Raw counts of parasites for each developmental stage. \[Supplementary Table S2\]\(#\). \[Primers used in PCR for sexing parasites. \\[Supplementary Table S3\\]\\(#\\). \\[Area raw measurements of developing worms. \\\[Supplementary Table\\\]\\\(#\\\)\\]\\(#\\)\]\(#\)](#)

S4. [Raw counting of parasites with either black positive \(hemozoin\) or negative \(no hemozoin\) intestine. Supplementary Table S5.](#) [Raw counting of Edu positive cells per parasite. Supplementary Table S6.](#) [Summary of all statistical tests.](#)

Supplementary Videos [Supplementary Video S1.](#) [Representative Z-stack of EdU+ cells after 2 days of *in vitro* culture with human serum \(A\) or foetal bovine serum \(B\). EdU+ cells and nuclei were labelled with Alexa fluor 488 \(green\) and DAPI \(grey\), respectively. Scale bars: 50 \$\mu\$ m.](#) [Supplementary Video S2.](#) [Representative Z-stack of an *in vitro* developed male worm. CellMask Green Actin Tracking Stain \(green\), and DAPI-stained nuclei \(cyan\). Scale bar: 25 \$\mu\$ m.](#) [Supplementary Video S3.](#) [Representative Z-stack of an *in vitro* developed female worm. CellMask Green Actin Tracking Stain \(green\), and DAPI-stained nuclei \(cyan\). Scale bar: 10 \$\mu\$ m.](#) [Supplementary Video S4.](#) [Representative video of *in vitro* developed females and males *in copula* at ~80 days in culture. Scale bar: 100 \$\mu\$ m.](#) [Supplementary Video S5.](#) [Representative videos of *in vivo* developed male and *in vitro* developed female *in copula* \(A\), and *in vivo* developed female and *in vitro* developed male *in copula* \(B\) Scale bar: 100 \$\mu\$ m.](#) [Supplementary Video S6.](#) [Z-stack of a schistosome pair \(*in vivo*-developed male and *in vitro*-developed female\) *in copula* \(A\), and magnification of the ovarian area \(B\), highlighting maturing oocytes. CellMask Green Actin Tracking Stain: green, DAPI: grey, CellMask Deep Red Plasma Membrane Stain purple. Scale bars: 25 \$\mu\$ m \(top panel\) and 50 \$\mu\$ m \(bottom panel\).](#)

References

- 1 LoVerde P. T. (2024) Schistosomiasis. *Adv Exp Med Biol* **1454**:75-105 https://doi.org/10.1007/978-3-031-60121-7_3
- 2 Buonfrate D., Ferrari T. C. A., Adegnik A. A., Russell Stothard J., Gobbi F. G. (2025) Human schistosomiasis. *Lancet* **405**:658-670 [https://doi.org/10.1016/S0140-6736\(24\)02814-9](https://doi.org/10.1016/S0140-6736(24)02814-9)
- 3 Berger D. J., et al. (2024) Extensive transmission and variation in a functional receptor for praziquantel resistance in endemic. *bioRxiv* <https://doi.org/10.1101/2024.08.29.610291>
- 4 Loker E. S., Brant S. V (2006) Diversification, dioecy and dimorphism in schistosomes. *Trends Parasitol* **22**:521-528 <https://doi.org/10.1016/j.pt.2006.09.001>
- 5 Chen R., et al. (2022) A male-derived nonribosomal peptide pheromone controls female schistosome development. *Cell* **185**:1506-1520.e1517 <https://doi.org/10.1016/j.cell.2022.03.017>
- 6 Shakir E. M. N., Rinaldi G., Kirk R. S., Walker A. J (2023) Schistosoma mansoni excretory-secretory products induce protein kinase signalling, hyperkinesia, and stem cell proliferation in the opposite sex. *Commun Biol* **6** <https://doi.org/10.1038/s42003-023-05333-9>
- 7 Elkrewi M., Moldovan M. A., Picard M. A. L., Vicoso B (2021) Schistosome W-Linked Genes Inform Temporal Dynamics of Sex Chromosome Evolution and Suggest Candidate for Sex Determination. *Mol Biol Evol* **38**:5345-5358 <https://doi.org/10.1093/molbev/msab178>
- 8 Fitzpatrick J. M., Hoffmann K. F (2006) Dioecious Schistosoma mansoni express divergent gene repertoires regulated by pairing. *Int J Parasitol* **36**:1081-1089 <https://doi.org/10.1016/j.ijpara.2006.06.007>
- 9 Lu Z., et al. (2016) Schistosome sex matters: a deep view into gonad-specific and pairing-dependent transcriptomes reveals a complex gender interplay. *Sci Rep* **6** <https://doi.org/10.1038/srep31150>
- 10 Picard M. A. L., et al. (2016) Sex-Biased Transcriptome of Schistosoma mansoni: Host-Parasite Interaction, Genetic Determinants and Epigenetic Regulators Are Associated with Sexual Differentiation. *PLoS Negl Trop Dis* **10**:e0004930 <https://doi.org/10.1371/journal.pntd.0004930>
- 11 Wangwiwatsin A., et al. (2020) Transcriptome of the parasitic flatworm Schistosoma mansoni during intra-mammalian development. *PLoS Negl Trop Dis* **14**:e0007743 <https://doi.org/10.1371/journal.pntd.0007743>
- 12 Diaz Soria C. L., et al. (2020) Single-cell atlas of the first intra-mammalian developmental stage of the human parasite Schistosoma mansoni. *Nat Commun* **11**:6411 <https://doi.org/10.1038/s41467-020-20092-5>

- 13 Li P., et al. (2021) Single-cell analysis of *Schistosoma mansoni* identifies a conserved genetic program controlling germline stem cell fate. *Nat Commun* **12** <https://doi.org/10.1038/s41467-020-20794-w>
- 14 Wendt G., et al. (2020) A single-cell RNA-seq atlas of identifies a key regulator of blood feeding. *Science* **369**:1644-1649 <https://doi.org/10.1126/science.abb7709>
- 15 Pance A., Rinaldi G. (2024) Editorial: Stem cell technologies meet stem cell biology to shine new light into tropical infectious diseases. *Front Cell Infect Microbiol* **14**:1411728 <https://doi.org/10.3389/fcimb.2024.1411728>
- 16 Sutrave S., Richter M. H (2023) The Truman Show for Human Helminthic Parasites: A Review of Recent Advances in In Vitro Cultivation Platforms. *Microorganisms* **11** <https://doi.org/10.3390/microorganisms11071708>
- 17 Ahmed N. H. (2014) Cultivation of parasites. *Trop Parasitol* **4**:80-89 <https://doi.org/10.4103/2229-5070.138534>
- 18 Mann V. H., Morales M. E., Rinaldi G., Brindley P. J (2010) Culture for genetic manipulation of developmental stages of *Schistosoma mansoni*. *Parasitology* **137**:451-462 <https://doi.org/10.1017/S0031182009991211>
- 19 Mellin R., Boddey J. A (2020) Organoids for Liver Stage Malaria Research. *Trends Parasitol* **36**:158-169 <https://doi.org/10.1016/j.pt.2019.12.003>
- 20 Zorrinho-Almeida M., de-Carvalho J., Bernabeu M., Silva Pereira S. (2025) Leveraging microphysiological systems to expedite understanding of host-parasite interactions. *PLoS Pathog* **21**:e1013088 <https://doi.org/10.1371/journal.ppat.1013088>
- 21 Faral-Tello P., Pagotto R., Bollati-Fogolín M., Francia M. E (2023) Modeling the human placental barrier to understand's vertical transmission. *Front Cell Infect Microbiol* **13** <https://doi.org/10.3389/fcimb.2023.1130901>
- 22 Korwin-Mihavics B. R., Dews E. A., di Genova B. M., Huston C. D (2023) Organoid-based systems to model infection in 2D and 3D. *bioRxiv* <https://doi.org/10.1101/2023.09.29.560165>
- 23 Zhou Y., et al. (2024) Full maturation of in vitro *Plasmodium falciparum* oocysts using the AlgiMatrix 3D culture system. *Malar J* **23** <https://doi.org/10.1186/s12936-024-05079-7>
- 24 Britton C., et al. (2023) New technologies to study helminth development and host-parasite interactions. *Int J Parasitol* **53**:393-403 <https://doi.org/10.1016/j.ijpara.2022.11.012>
- 25 Duque-Correa M. A., et al. (2022) Defining the early stages of intestinal colonisation by whipworms. *Nat Commun* **13**:1725 <https://doi.org/10.1038/s41467-022-29334-0>
- 26 Faber M. N., et al. (2022) Development of Bovine Gastric Organoids as a Novel Model to Study Host-Parasite Interactions in Gastrointestinal Nematode Infections. *Front Cell Infect Microbiol* **12**:904606 <https://doi.org/10.3389/fcimb.2022.904606>
- 27 Vitkauskaitė A., et al. (2025) In vitro co-culture of *Fasciola hepatica* newly excysted juveniles (NEJs) with 3D HepG2 spheroids permits novel investigation of host-parasite interactions. *Virulence* <https://doi.org/10.1080/21505594.2025.2482159>
- 28 White R., Blow F., Buck A. H., Duque-Correa M. A (2022) Organoids as tools to investigate gastrointestinal nematode development and host interactions. *Front Cell Infect Microbiol* **12**:976017 <https://doi.org/10.3389/fcimb.2022.976017>
- 29 Herz M., et al. (2024) Genome-wide transcriptome analysis of larvae and germinative cell cultures reveals genes involved in parasite stem cell function. *Front Cell Infect Microbiol* **14**:1335946 <https://doi.org/10.3389/fcimb.2024.1335946>
- 30 Koziol U., Brehm K (2015) Recent advances in *Echinococcus* genomics and stem cell research. *Vet Parasitol* **213**:92-102 <https://doi.org/10.1016/j.vetpar.2015.07.031>
- 31 Basch P. F (1981) Cultivation of *Schistosoma mansoni* in vitro. II. production of infertile eggs by worm pairs cultured from cercariae. *J Parasitol* **67**:186-190 [PubMed](https://pubmed.ncbi.nlm.nih.gov/67186190/)

- 32 Basch P. F (1981) Cultivation of *Schistosoma mansoni* in vitro. I. Establishment of cultures from cercariae and development until pairing. *J. Parasitol* **67**:179-185 [PubMed](#)
- 33 Basch P. F., Humbert R (1981) Cultivation of *Schistosoma mansoni* in vitro. III. implantation of cultured worms into mouse mesenteric veins. *J Parasitol* **67**:191-195 [PubMed](#)
- 34 Wang J., Chen R., Collins J. J (2019) Systematically improved in vitro culture conditions reveal new insights into the reproductive biology of the human parasite *Schistosoma mansoni*. *PLoS Biol* **17**:e3000254 <https://doi.org/10.1371/journal.pbio.3000254>
- 35 You Y., et al. (2024) An improved medium for in vitro studies of female reproduction and oviposition in *Schistosoma japonicum*. *Parasit Vectors* **17**:116 <https://doi.org/10.1186/s13071-024-06191-y>
- 36 Anisuzzaman, et al. (2021) Host-Specific Serum Factors Control the Development and Survival of. *Front Immunol* **12**:635622 <https://doi.org/10.3389/fimmu.2021.635622>
- 37 Buchter V., Schneeberger P. H. H., Keiser J (2021) Validation of a human-serum-based in vitro growth method for drug screening on juvenile development stages of *Schistosoma mansoni*. *PLoS Negl Trop Dis* **15**:e0009313 <https://doi.org/10.1371/journal.pntd.0009313>
- 38 Maharjan S., Kirk R. S., Lawton S. P., Walker A. J (2021) Further evaluation and validation of HybridoMed Diff 1000 and its comparison to Basch medium for the cell-free culture of *Schistosoma mansoni* juvenile worm stages. *Int. J. Parasitol* **51**:613-619 <https://doi.org/10.1016/j.ijpara.2020.12.009>
- 39 Reimers N., et al. (2015) Drug-induced exposure of *Schistosoma mansoni* antigens SmCD59a and SmKK7. *PLoS Negl Trop Dis* **9**:e0003593 <https://doi.org/10.1371/journal.pntd.0003593>
- 40 Abba N., et al. (2017) Evaluation of the pharmacokinetic-pharmacodynamic relationship of praziquantel in the *Schistosoma mansoni* mouse model. *PLoS Negl Trop Dis* **11**:e0005942 <https://doi.org/10.1371/journal.pntd.0005942>
- 41 Milligan J. N., Jolly E. R (2011) Cercarial transformation and in vitro cultivation of *Schistosoma mansoni* schistosomules. *J Vis Exp* <https://doi.org/10.3791/3191>
- 42 Nation C. S., Da'dara A. A., Marchant J. K., Skelly P. J (2020) Schistosome migration in the definitive host. *PLoS Negl Trop Dis* **14**:e0007951 <https://doi.org/10.1371/journal.pntd.0007951>
- 43 Canepa M., Ding A., Fechner C. (2024) Cellular Adaptation: Hypertrophy, Hyperplasia, Atrophy. *General Pathology Student Guide* 50-64 https://doi.org/10.1007/978-3-662-67962-3_3
- 44 Nanes Sarfati D., Li P., Tarashansky A. J., Wang B. (2021) Single-cell deconstruction of stem-cell-driven schistosome development. *Trends Parasitol* **37**:790-802 <https://doi.org/10.1016/j.pt.2021.03.005>
- 45 Wang B., et al. (2018) Stem cell heterogeneity drives the parasitic life cycle of. *eLife* **7** <https://doi.org/10.7554/eLife.35449>
- 46 Arunsan P., et al. (2019) Programmed knockout mutation of liver fluke granulin attenuates virulence of infection-induced hepatobiliary morbidity. *eLife* **8** <https://doi.org/10.7554/eLife.41463>
- 47 Kalinna B. H., Ross A. G., Walduck A. K (2024) Schistosome Transgenesis: The Long Road to Success. *Biology* **13** <https://doi.org/10.3390/biology13010048>
- 48 Quinzo M. J., Perteguer M. J., Brindley P. J., Loukas A., Sotillo J (2022) Transgenesis in parasitic helminths: a brief history and prospects for the future. *Parasit Vectors* **15** <https://doi.org/10.1186/s13071-022-05211-z>
- 49 Stephens D. R., et al. (2025) A genome-scale drug discovery pipeline uncovers therapeutic targets and a unique p97 allosteric binding site in. *Proc Natl Acad Sci U S A* **122**:e2505710122 <https://doi.org/10.1073/pnas.2505710122>
- 50 Xie Y., et al. (2025) RNAi screening of uncharacterized genes identifies promising druggable targets in *Schistosoma japonicum*. *PLoS Pathog* **21**:e1013014 <https://doi.org/10.1371/journal.ppat.1013014>
- 51 Loker E. S. (2025) Review: the origin of digenetic trematodes: did the story begin with the blood flukes?. *J Parasitol* **111**:787-814 <https://doi.org/10.1645/25-27>

- 52 Moné H., Boissier J (2004) Sexual biology of schistosomes. *Adv Parasitol* **57**:89-189 [https://doi.org/10.1016/S0065-308X\(04\)57002-1](https://doi.org/10.1016/S0065-308X(04)57002-1)
- 53 Puck T. T., Cieciora S. J., Robinson A (1958) Genetics of somatic mammalian cells. III. Long-term cultivation of euploid cells from human and animal subjects. *J Exp Med* **108**:945-956 <https://doi.org/10.1084/jem.108.6.945>
- 54 van der Valk J., et al. (2018) Fetal Bovine Serum (FBS): Past - Present - Future. *Altex* **35**:99-118 <https://doi.org/10.14573/altex.1705101>
- 55 Gstraunthaler G., Lindl T., van der Valk J (2013) A plea to reduce or replace fetal bovine serum in cell culture media. *Cytotechnology* **65**:791-793 <https://doi.org/10.1007/s10616-013-9633-8>
- 56 Weber T., et al. (2025) Fetal bovine serum: how to leave it behind in the pursuit of more reliable science. *Front Toxicol* **7** <https://doi.org/10.3389/ftox.2025.1612903>
- 57 Rosolowski J., Weber T., Malakpour-Permlid A., Oredsson S (2025) Revisiting 3Rs: rethinking replacement and new approach methodologies. *Front Toxicol* **7** <https://doi.org/10.3389/ftox.2025.1664209>
- 58 Subbiahanadar Chelladurai K., et al. (2021) Alternative to FBS in animal cell culture - An overview and future perspective. *Heliyon* **7**:e07686 <https://doi.org/10.1016/j.heliyon.2021.e07686>
- 59 Xiao S.-H., Sun J (2017) Schistosoma hemozoin and its possible roles. *Int J Parasitol* **47**:171-183 <https://doi.org/10.1016/j.ijpara.2016.10.005>
- 60 Sun J., et al. (2025) Hemozoin: a waste product after heme detoxification?. *Parasit Vectors* **18** <https://doi.org/10.1186/s13071-025-06699-x>
- 61 Truscott M., Evans D. A., Gunn M., Hoffmann K. F (2013) Schistosoma mansoni hemozoin modulates alternative activation of macrophages via specific suppression of Retnla expression and secretion. *Infect Immun* **81**:133-142 <https://doi.org/10.1128/IAI.00701-12>
- 62 Miller P., Wilson R. A (1980) Migration of the schistosomula of Schistosoma mansoni from the lungs to the hepatic portal system. *Parasitology* **80**:267-288 <https://doi.org/10.1017/s0031182000000743>
- 63 Corrêa Soares J. B. R., et al. (2009) Interference with hemozoin formation represents an important mechanism of schistosomicidal action of antimalarial quinoline methanols. *PLoS Negl Trop Dis* **3**:e477 <https://doi.org/10.1371/journal.pntd.0000477>
- 64 Frahm S., et al. (2019) A novel cell-free method to culture Schistosoma mansoni from cercariae to juvenile worm stages for in vitro drug testing. *PLoS Negl Trop Dis* **13**:e0006590 <https://doi.org/10.1371/journal.pntd.0006590>
- 65 Collins J. J., et al. (2013) Adult somatic stem cells in the human parasite Schistosoma mansoni. *Nature* **494**:476-479 <https://doi.org/10.1038/nature11924>
- 66 Moescheid M. F., et al. (2025) The retinoic acid family-like nuclear receptor SmRAR identified by single-cell transcriptomics of ovarian cells controls oocyte differentiation in Schistosoma mansoni. *Nucleic Acids Res* **53** <https://doi.org/10.1093/nar/gkae1228>
- 67 Doenhoff M. J., Modha J., Walker A. J (2019) Failure of in vitro-cultured schistosomes to produce eggs: how does the parasite meet its needs for host-derived cytokines such as TGF- β ?. *Int J Parasitol* **49**:747-757 <https://doi.org/10.1016/j.ijpara.2019.05.004>
- 68 Geyer K. K., et al. (2017) The Biomphalaria glabrata DNA methylation machinery displays spatial tissue expression, is differentially active in distinct snail populations and is modulated by interactions with Schistosoma mansoni. *PLoS Negl Trop Dis* **11**:e0005246 <https://doi.org/10.1371/journal.pntd.0005246>
- 69 Buddenborg S. K., Lu Z., Sankaranarayan G., Doyle S. R., Berriman M (2023) The stage- and sex-specific transcriptome of the human parasite Schistosoma mansoni. *Sci Data* **10** <https://doi.org/10.1038/s41597-023-02674-2>
- 70 Grevelding C. G (1999) Genomic instability in Schistosoma mansoni. *Mol. Biochem. Parasitol* **101**:207-216 [https://doi.org/10.1016/s0166-6851\(99\)00078-x](https://doi.org/10.1016/s0166-6851(99)00078-x)

- 71 Rinaldi G., et al. (2009) RNA interference targeting leucine aminopeptidase blocks hatching of *Schistosoma mansoni* eggs. *Mol. Biochem. Parasitol* **167**:118-126
<https://doi.org/10.1016/j.molbiopara.2009.05.002>

Peer reviews

Reviewer #1 (Public review):

Pichon, Rémi et al. describe an in vitro method for transforming *Schistosoma* cercariae into mature adult worms. The authors show that human serum (HS) supports parasite growth and differentiation more effectively than fetal bovine serum (FBS). They also observed differences in parasite growth and activity, with worms cultured in HS efficiently digesting human red blood cells (hRBC). Cultured worms were able to pair with ex vivo adult worms and produce eggs, indicating functional maturation suitable for downstream applications such as drug screening. While the experimental approach is comprehensive and supports the advantages of HS culture conditions, the pairing efficiency was low ($\approx 7\%$) and required long culture periods (70-80 days), highlighting limitations that may affect reproducibility.

A major strength of the study, in particular, is that the authors clearly differentiate the effects of FBS versus HS on developmental progression. The conversion rate observed in HS cultures is significant and consistent with previously published data.

While the study has several strengths, some aspects of the work are not fully explored. In particular, the role of hRBC supplementation requires further clarification. Although HS-cultured worms were shown to digest hRBC more readily, the implications of this observation remain unclear. Specifically, it would be useful to understand whether hRBC supplementation influences (1) long-term culture stability, (2) molecular pathways associated with development and differentiation, or (3) the pairing capacity of the worms. While addressing these questions may not be the main objective of the study, further discussion of these points would strengthen the manuscript.

The manuscript is clearly written and represents a valuable contribution to the field. Overall, the experimental approach is sound, and the results support a useful methodological framework for the in vitro culture of *Schistosoma* worms and the attainment of sexual maturity, particularly for adult male worms.

<https://doi.org/10.7554/eLife.111066.1.sa3>

Reviewer #2 (Public review):

Summary:

The authors perform confirmation studies of Paul Basch's seminal schistosome work from 1981, demonstrating the development of transformed schistosomules into sexually dimorphic adult parasites, albeit without successful egg production. In addition to the findings from Basch's earlier work, the authors add some new molecular data in the form of an analysis of proliferative cells in in-vitro-derived animals.

Strengths:

The authors successfully confirm experimental results from earlier schistosome researchers, providing a potential new tool for studying schistosome biology without the need for vertebrate hosts.

Weaknesses:

The display of data from the authors is sometimes difficult to follow/understand where it comes from. For example:

(1) Line 136: The authors claim that parasites in HS and FBS conditions have substantially different mortality rates (11.3 +/- 2.7 vs 5 +/- 2.3) but a quite high p-value (0.8). Analyzing the raw data myself, I obtained a mean of 8.2 +/- 1.7% vs 4.8% +/- 4.3% with a p-value of 0.15. Either the data are not clearly presented, and I did not follow them, or the data presented in the text do not match the raw data in the supplemental files.

(2) Line 187/Figure 4: Though it is not clearly stated, it appears that the authors treat their EdU counts as an ordinal data set of 61 steps (from 0 to >60) rather than a continuous measure of EdU+ cells per animal. In this author's opinion, the graph strongly suggests a continuous data set, and the fact that this reviewer had to dig through poorly-labeled raw data to discover the nature of the data is problematic. The authors should either switch to a continuous data set or make it explicit that the data shown are ordinal. If counting EdU+ cells is too arduous, the authors could consider comparing the amount of EdU+ area to the amount of DAPI+ area in maximum intensity projections of their confocal images, as this would roughly approximate the amount of proliferative cells in the animals.

There are some minor issues as well:

(1) Line 122: It is perhaps incorrect to refer to humans as "the" definitive host of schistosomes, as *S. japonicum* is primarily considered a zoonotic infection with water buffalo/cows being the primary definitive host.

(2) Line 185/298: The authors refer to EdU pulse-chase experiments, but the experiments described here are EdU pulse experiments.

<https://doi.org/10.7554/eLife.111066.1.sa2>

Reviewer #3 (Public review):

Summary:

This study is significant as it established a protocol for the long-term culture of *Schistosoma mansoni* newly transformed cercariae, which developed in vitro into sexually dimorphic forms. The impact of two different sera, Fetal Bovine Serum (FBS) and Human Serum (HS), added to the culture medium supplemented with human red blood cells was evaluated. The authors demonstrated that HS-cultured parasites were able to digest red blood cells, a critical step for long-term parasite development. Furthermore, while most FBS-cultured parasites did not progress beyond an early liver stage, sexual dimorphism was clearly evident in the HS-cultured worms, albeit delayed compared to in vivo development.

Strengths:

This study could contribute to further in vitro studies for a better understanding of the unique sexual biology of *Schistosoma mansoni* and for screening novel schistosomicidal compounds. By increasing parasite development in in vitro studies, this protocol could have a positive impact on the principles of the 3Rs (Replacement, Reduction and Refinement) for animal research.

Weaknesses:

As the authors mentioned, "pairing between male and female parasites was rare. Pairing was observed in approximately ~7% of the experiments, usually after day ~ 80 in culture. Egg production was also not achieved with this protocol.

<https://doi.org/10.7554/eLife.111066.1.sa1>

Author response:

eLife Assessment

This useful study presents an improved protocol for long-term in vitro culture of Schistosoma mansoni that enables progression toward sexually dimorphic stages, representing a meaningful advance for studying parasite development and reducing reliance on animal models. The findings show that host-specific culture conditions support essential developmental and metabolic functions required for parasite maturation, although development remains delayed compared to in vivo conditions. The evidence is solid overall, but limited pairing efficiency and the absence of egg production indicate that the system does not yet fully recapitulate complete reproductive development.

On behalf of the co-authors, we thank the three reviewers and the editors for their complimentary remarks as well as the major and minor comments/ concerns. Addressing these concerns have led to revisions that improved the manuscript. In particular, further analyses have generated an updated Figures 3 and 4, and Supplementary Tables S1, and S4-S6.

Public Reviews:

Reviewer #1 (Public review):

Pichon, Rémi et al. describe an in vitro method for transforming Schistosoma cercariae into mature adult worms. The authors show that human serum (HS) supports parasite growth and differentiation more effectively than fetal bovine serum (FBS). They also observed differences in parasite growth and activity, with worms cultured in HS efficiently digesting human red blood cells (hRBC). Cultured worms were able to pair with ex vivo adult worms and produce eggs, indicating functional maturation suitable for downstream applications such as drug screening. While the experimental approach is comprehensive and supports the advantage of HS culture conditions, the pairing efficiency was low ($\approx 7\%$) and required long culture periods (70-80 days), highlighting limitations that may affect reproducibility.

We acknowledge the reviewer for the positive highlights. Regarding the low *in vitro* pairing efficiency, we have now edited the manuscript to clarify a misleading statement related to 7%. We decided to remove the value of 7% — which corresponds to the percentage of experiments in which couples were observed, as it does not accurately represent the actual number of observed worm pairs and it is probably misleading. We have updated the text as follows:

Results, lines 230 ff.:

“While the establishment of sexual dimorphism was robust and reproducible across more than 15 independent experiments, pairing between male and female parasites was rare. Pairing was observed only in experiments lasting more than 80 days in which we were only able to observe a few couples. In addition, these pairings were temporary (Figures 6A, B; Supplementary Video S4).”

We also agree with the reviewer that the extended culture periods required to obtain fully sexually dimorphic parasites remain a limitation. As elaborated in Discussion (see below), key factors, probably derived from the host, are missing in the *in vitro* system explaining both the slow *in vitro* development and low rate of spontaneous pairing between *in vitro*

developed, sexually dimorphic male and female worms. This was discussed as follows (lines 340-343): “That said, while our system was highly efficient in producing sexually dimorphic worms, spontaneous pairing between male and female parasites was extremely rare, mainly in aged in vitro cultures (from 80 to 100 days in culture) indicating that other factors, e.g., cholesterol, may be missing[35].”

A major strength of the study, in particular, is that the authors clearly differentiate the effects of FBS versus HS on developmental progression. The conversion rate observed in HS cultures is significant and consistent with previously published data.

While the study has several strengths, some aspects of the work are not fully explored. In particular, the role of hRBC supplementation requires further clarification. Although HS-cultured worms were shown to digest hRBC more readily, the implications of this observation remain unclear. Specifically, it would be useful to understand whether hRBC supplementation influences (1) long-term culture stability, (2) molecular pathways associated with development and differentiation, or (3) the pairing capacity of the worms. While addressing these questions may not be the main objective of the study, further discussion of these points would strengthen the manuscript.

We agree that deciphering the role of the human Red Blood Cells (hRBCs) supplementation is critical. Regarding the influence of hRBCs on the long-term culture stability in parasite development it has been well established for more than four decades that schistosomes do need red blood cells to grow in culture [Basch, P. F. Cultivation of *Schistosoma mansoni* in vitro. II. production of infertile eggs by worm pairs cultured from cercariae. *J Parasitol* 67, 186-190 (1981); Basch, P. F. Cultivation of *Schistosoma mansoni* in vitro. I. Establishment of cultures from cercariae and development until pairing. *J. Parasitol.* 67, 179-185 (1981)]. The molecular pathways underlying development, sexual differentiation and pairing and modulated by hRBCs in culture is currently being investigated by our team. We decided not to include these data and analyses in the current manuscript, as they fall outside its scope.

*The manuscript is clearly written and represents a valuable contribution to the field. Overall, the experimental approach is sound, and the results support a useful methodological framework for the in vitro culture of *Schistosoma* worms and the attainment of sexual maturity, particularly for adult male worms.*

We thank the reviewer for highlighting the manuscript's strengths.

Reviewer #2 (Public review):

Summary:

The authors perform confirmation studies of Paul Basch's seminal schistosome work from 1981, demonstrating the development of transformed schistosomules into sexually dimorphic adult parasites, albeit without successful egg production. In addition to the findings from Basch's earlier work, the authors add some new molecular data in the form of an analysis of proliferative cells in in-vitro-derived animals.

Strengths:

The authors successfully confirm experimental results from earlier schistosome researchers, providing a potential new tool for studying schistosome biology without the need for vertebrate hosts.

We thank the reviewer for highlighting the manuscript's strengths.

Weaknesses:

The display of data from the authors is sometimes difficult to follow/understand where it comes from. For example:

(1) Line 136: The authors claim that parasites in HS and FBS conditions have substantially different mortality rates (11.3 +/- 2.7 vs 5 +/- 2.3) but a quite high p-value (0.8). Analyzing the raw data myself, I obtained a mean of 8.2 +/- 1.7% vs 4.8% +/- 4.3% with a p-value of 0.15. Either the data are not clearly presented, and I did not follow them, or the data presented in the text do not match the raw data in the supplemental files.

We thank the reviewer for pointing this out; we have now edited Supplementary Tables S1 and S6 by turning them into a long format for the sake of clarity. Accordingly, Results, Methods sections, and indicated supplementary tables were edited as follows:

Results, lines 142 ff.:

“No morphological differences were observed between parasites cultured either in FBS or HS within the first week in culture; in both conditions most parasites were classified as early schistosomula [category 1: 76% ± 30 (average ± SD) in FBS and 73% ± 29 (average ± SD) in HS] with few lung (category 2) and early liver schistosomula (category 3) (Figure 1B, week 1; Supplementary Figure S1). The mean mortality (category 0) at week 1 was slightly higher, but not statistically significant (P= 0.42), in worms cultured in HS [9.75% ± 2.76 (average ± SD)] compared to the mortality registered in FBS-cultured parasites [5.52% ± 5.18 (average ± SD), Supplementary Table S6], consistent with previous findings[39].”

Methods, lines 463-465:

“To evaluate differences in mortality between HS- and FBS-cultured parasites, data from 5 experiments were combined and analysed using a Shapiro-Wilk normality test to test normality of the data and a non-parametric Wilcoxon rank sum exact test (Supplementary Tables S1 and S6).”

Supplementary Tables:

Supplementary Table S1. “Raw counts of parasites within each developmental stage category. Each row corresponds to a picture of parasites in culture medium containing FBS or HS. Each column corresponds to the raw parasite counts at indicated stage development (categories 0 to 5), time in culture (Time in days - D), and experimental condition.”

Supplementary Table S6. “Summary of all statistical tests employed in this study. 1. Statistical tests of parasite mortality and the raw data table used for this test. 2. Statistical tests for worm size comparisons (correspond to Figure 2). 3. Statistical tests for worm black gut comparisons (correspond to Figure 3). BG: Black gut. 4. Statistical tests for EdU positive cells comparisons (correspond to Figure 4). Replicate code: E, M and L correspond to day 2, 8 and 15 respectively; R and W correspond to the presence (R) or absence (W) of RBCs added 13 days after transformation.”

For clarity, in Author response image 1 we provide the R script used to perform the statistical tests on the data shown in Supplementary Table S6 (column Raw count of parasite developmental category per image and experiment)

```

library(dplyr)
Table_S6 <- read.csv("/Users/RemiPichon/Desktop/test1/Table_S6.csv", header = TRUE, sep = ",")
# Step 1: Compute the total number of cells per row
Table_S6 <- Table_S6 %>% mutate(Total = Category.0 + Category.1 + Category.2 + Category.3 + Category.4 + Category.5)
# Step 2: Aggregate data by biological replicate (Experiment), condition, and week
exp_pct <- Table_S6 %>%
  group_by(Experiment, Condition, Week) %>%
  summarise(
    total_cells = sum(Total, na.rm = TRUE),
    cat0 = sum(Category.0, na.rm = TRUE),
    cat1 = sum(Category.1, na.rm = TRUE),
    cat2 = sum(Category.2, na.rm = TRUE),
    cat3 = sum(Category.3, na.rm = TRUE),
    cat4 = sum(Category.4, na.rm = TRUE),
    cat5 = sum(Category.5, na.rm = TRUE),
    .groups = "drop") %>% mutate(
    cat0_pct = 100 * cat0 / total_cells,
    cat1_pct = 100 * cat1 / total_cells,
    cat2_pct = 100 * cat2 / total_cells,
    cat3_pct = 100 * cat3 / total_cells,
    cat4_pct = 100 * cat4 / total_cells,
    cat5_pct = 100 * cat5 / total_cells)
# Step 3: Compute mean and standard deviation across experiments
summary_pct <- exp_pct %>%
  group_by(Condition, Week) %>%
  summarise(
    n_exp = n(),
    cat0_mean = mean(cat0_pct, na.rm = TRUE),
    cat0_sd = sd(cat0_pct, na.rm = TRUE),
    cat1_mean = mean(cat1_pct, na.rm = TRUE),
    cat1_sd = sd(cat1_pct, na.rm = TRUE),
    cat2_mean = mean(cat2_pct, na.rm = TRUE),
    cat2_sd = sd(cat2_pct, na.rm = TRUE),
    cat3_mean = mean(cat3_pct, na.rm = TRUE),
    cat3_sd = sd(cat3_pct, na.rm = TRUE),
    cat4_mean = mean(cat4_pct, na.rm = TRUE),
    cat4_sd = sd(cat4_pct, na.rm = TRUE),
    cat5_mean = mean(cat5_pct, na.rm = TRUE),
    cat5_sd = sd(cat5_pct, na.rm = TRUE),
    .groups = "drop")
print(summary_pct)
# Step 4: Export the summary table
#write.csv(summary_pct, "summary_pct.csv", row.names = FALSE)
# Step 5: Extract week 1 only for Category 0 statistical testing
week1_cat0 <- exp_pct %>% filter(Week == 1)
# Step 6: Test normality separately in each condition using Shapiro-Wilk
shapiro.test(week1_cat0$cat0_pct[week1_cat0$Condition == "FBS"])
shapiro.test(week1_cat0$cat0_pct[week1_cat0$Condition == "HS"])
# Step 7: Compare FBS vs HS using a Wilcoxon rank-sum test
#nit is a non-parametric test, suitable when normality is not guaranteed and when sample sizes are small.
wilcox.test(cat0_pct ~ Condition, data = week1_cat0)
week1_cat0 %>% select(Experiment, Condition, cat0_pct)

```

Author response image 1.

(2) Line 187/Figure 4: Though it is not clearly stated, it appears that the authors treat their EdU counts as an ordinal data set of 61 steps (from 0 to >60) rather than a continuous measure of EdU+ cells per animal. In this author's opinion, the graph strongly suggests a continuous data set, and the fact that this reviewer had to dig through poorly-labeled raw data to discover the nature of the data is problematic. The authors should either switch to a continuous data set or make it explicit that the data shown are ordinal. If counting EdU+ cells is too arduous, the authors could consider comparing the amount of EdU+ area to the amount of DAPI+ area in maximum intensity projections of their confocal images, as this would roughly approximate the amount of proliferative cells in the animals.

As the reviewer correctly pointed out, the data were treated as ordinal because counting worms with more than 60 Edu+ cells became extremely difficult and highly inaccurate.

Therefore, we decided to group in a single category, “60 EdU+ cells”, all worms showing more than 60 EdU+ cells. We have now updated Figure 4 where medians are shown instead of media values, Supplementary Table S5 to provide more comprehensive access to the raw counts, and Supplementary Table S6 to indicate the data for EdU+ cells per worm were considered ordinal. Accordingly, we have revised the corresponding sections as follows:

Results, lines 211 ff.:

“HS-cultured schistosomula showed higher numbers of proliferating stem cells, with a median of >48 and >60 EdU+ cells per worm at days 8 and 15, respectively (Figure 4). On the other hand, most FBS-cultured parasites displayed no more than an average of 20 EdU+ cells per worm (Figure 4).”

Methods, lines 520 ff.:

“EdU+ cells per parasite were counted for an average of 100 parasites across three independent experiments (Supplementary Table S5). Worms were grouped based on the number of cells per individual, but all those showing ≥ 60 EdU+ cells were counted in the same group named ‘60 EdU+ cells’. Therefore, the data were considered ordinal data. Statistical analysis was performed by Kruskal-Wallis test with Dunn multiple comparison post-hoc test, with $P \leq 0.05$ considered significant (Supplementary Table S6).”

Figure 4 legend, lines 830 ff.:

“A. Violin plots showing the number of Edu+ cells per worm at indicated time points (2, 8, and 15 days post cercarial transformation) in parasites cultured either in Foetal Bovine Serum (FBS, blue) or Human Serum (HS, light brown). Human Red Blood Cells (hRBCs) were added in the culture at day 13 post cercarial transformation. The small black dots indicate individual worms, and the big black point indicates the median of EdU+ cells per worm. All worms showing ≥ 60 EdU+ cells were counted and clustered together in the group named ‘60 EdU+ cells’. Hence, the data were treated as ordinal and statistical analysis performed by Kruskal-Wallis test with Dunn multiple comparison post-hoc test, with $P \leq 0.05$ (*) considered significant (Supplementary Tables S5 and S6).”

We thank the reviewer for the very interesting suggestion to quantify cell proliferation by calculating the ratio between EdU+ area to DAPI+ area in maximum intensity projections images. Measuring the fluorescence area for each worm in maximum projection is an excellent idea; however, due to the number of EdU+ cells present in some samples, we think this technique would not provide additional information or produce more detailed data compared with our analysis when the number of Edu+ cells exceeds 60 per worm. We will certainly consider this approximation for future studies.

There are some minor issues as well:

(1) Line 122: It is perhaps incorrect to refer to humans as "the" definitive host of schistosomes, as S. japonicum is primarily considered a zoonotic infection with water buffalo/cows being the primary definitive host.

We thank the reviewer for pointing this out; we have now replaced “schistosomes” with “*Schistosoma mansoni*” (current line 131)

(2) Line 185/298: The authors refer to EdU pulse-chase experiments, but the experiments described here are EdU pulse experiments.

This is a very good point, we thank the reviewer for bringing this up and have accordingly edited by replacing “EdU pulse-chase” with “EdU pulse” experiments in lines 37, 204, and 321.

Reviewer #3 (Public review):

Summary:

This study is significant as it established a protocol for the long-term culture of Schistosoma mansoni newly transformed cercariae, which developed in vitro into sexually dimorphic forms. The impact of two different sera, Fetal Bovine Serum (FBS) and Human Serum (HS), added to the culture medium supplemented with human red blood cells was evaluated. The authors demonstrated that HS-cultured parasites were able to digest red blood cells, a critical step for long-term parasite development. Furthermore, while most FBS-cultured parasites did not progress beyond an early liver stage, sexual dimorphism was clearly evident in the HS-cultured worms, albeit delayed compared to in vivo development.

Strengths:

This study could contribute to further in vitro studies for a better understanding of the unique sexual biology of Schistosoma mansoni and for screening novel schistosomicidal compounds. By increasing parasite development in in vitro studies, this protocol could have a positive impact on the principles of the 3Rs (Replacement, Reduction and Refinement) for animal research.

We thank the reviewer for highlighting the manuscript's strengths.

Weaknesses:

As the authors mentioned, "pairing between male and female parasites was rare. Pairing was observed in approximately ~7% of the experiments, usually after day ~ 80 in culture. Egg production was also not achieved with this protocol.

Following the reviewer's point and to clarify a misleading point, we have now decided to remove the value of 7% — which corresponds to the percentage of experiments in which couples were observed. However, this value does not accurately reflect the actual number of observed worm pairs, and it is probably misleading. We have updated the text as follows:

Results, lines 230 ff.:

“While the establishment of sexual dimorphism was robust and reproducible across more than 15 independent experiments, pairing between male and female parasites was rare. Pairing was observed only in experiments lasting more than 80 days in which we were only able to observe a few couples. In addition, these pairings were temporary (Figures 6A, B; Supplementary Video S4).”

<https://doi.org/10.7554/eLife.111066.1.sa0>