In vitro sexual dimorphism establishment in schistosomes

Rémi Pichon*[1,2], Magda E Lotkowska*[3], Jude L. D. Bulathsinghalage[2], Madeleine McMath[2], Mary Evans[2], Benjamin J. Hulme[2], Kirsty Ambridge[3], Geetha Sankaranarayanan[3], Simon Kershenbaum[1], Sarah D. Davey[2], Josephine E. Forde-Thomas[2], Karl F. Hoffmann[2], Matthew Berriman**[3,4], Gabriel Rinaldi**[1,2,3]

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

*co-first authors, **co-corresponding authors

Correspondence:
Matt Berriman: matt.berriman@glasgow.ac.uk
Gabriel Rinaldi: gabriel.rinaldi@biology.ox.ac.uk

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* are needed 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 development. 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 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.

Keywords. Schistosomiasis; Schistosoma spp; Long-term culture; Human Serum; *In vitro* schistosome development; Sexual dimorphism

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. Infection with *Schistosoma mansoni* leads to eggs lodged mainly in the liver, inducing inflation, fibrosis and granuloma formation[2]. 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 resistance[3]. 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 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 host[4]. 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 the 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 to 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 challenging[16].

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 studies[16]. 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 1981[32]. Despite one research group referring to the
approach[40], there has been no detailed description and no widespread adoption. In fact, published
culturing studies have continued to utilise FBS[34,41].
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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 effects 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 a day 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.
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Results

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118 Sexually dimorphic schistosomes developed entirely *in vitro* from cercariae
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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 fully replicated. More recently, culture media were complemented with HS in studies
focused on early stages of parasite development for drug testing[36-38]; however, these worms never
progressed beyond the liver stage. 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% on 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 findings[38].

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 ~80% by Week 10, after which the experiments under this
condition were stopped as most parasites were dead (**Supplementary Figure S2**). In contrast,
parasites cultured in HS further developed over time throughout all categories, achieving marked
sexual dimorphism by Week 6 (**Figure 1B; Supplementary Figure S3A**), as confirmed by PCR
(**Supplementary Figure S3B; Supplementary Table S2**). As previously described for the *in vivo*
development of schistosomes[11], *in vitro* cultured parasites showed developmental asynchrony in
agreement with Basch's observations[32]; however, by Week 10 the majority of the worms (73.7 $\pm$
25.4%) had 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 on average 110,069
$\mu m^2 \pm 72,959$ μm^2, ~20-fold larger than parasites cultured in FBS for 10 weeks, which only grew
slightly, plateauing at an average 5,417 $\mu m^2 \pm 1,322$ μm^2, similar to the size reached by parasites
after only two weeks in HS and always significantly smaller than worms in HS for all the weeks (P-
value < 0.01, Supplementary Table S6) (**Figures 1 and 2**).

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 liver[42]. 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 began 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 tissues[43]. In schistosomes, a complex stem cell system comprised of both somatic and germline stem cells has been described by leveraging recent single cell transcriptomics[44]. 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**) compared to HS-cultured parasites. 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 schistosomula[45]. 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 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-suplemented culture medium (**Figure 1,** Week 6). Confocal microscopy of DAPI- and Phalloidin- stained *in vitro* cultured male worms confirmed the presence of the 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**).

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 culture with worms of the opposite sex developed *in vitro* (>70 days). Within 24 hours of initiating co-culturing of *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 were capable of intersexual pairing. Moreover, *in vitro* developed females
coupled with *ex vivo* collected mature males displayed signs of primordial ovary 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 advance 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 production[52]. 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 compared the effect of modified Basch's media[18], 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.
Basch[31] 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 1958[53] providing a reliable source
of amino acids, carbohydrates, hormones, lipids, proteins, vitamins, and growth factors[54].
Nevertheless, the scientific community is increasingly advocating for the replacement of FBS as a
supplement in tissue culture[55]. 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 welfare[56].
Our protocol, that supports the 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 gastrodermis and oesophageal
gland development; worms cultured in the presence of HS displayed a differentiated gastrodermis
that enabled haemoglobin digestion and accumulation of hemozoin within the intestines[59]. In
contrast, 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 peroxidation[60]. 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 stages[59] . In addition, increasing evidence shows that hemozoin may
play critical roles during parasite development by supplying iron for egg production[59], and
interaction with the mammalian host by an immune modulatory role[61]. In the mouse model,
schistosomula 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, schistosomula collected from experimentally infected mice
at day 13 post infection already show hemozoin in their guts[11]. Similarly, HS-developed
schistosomula 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 schistosomula. A better understanding of the *in vitro* development of
the parasite oesophageal gland and gastrodermis and its role in the production of hemozoin will
expose tentative novel targets for control[63].

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 worms[65], followed by single cell transcriptomic identification and
functional characterisation of three key stem cell populations in intra-snail stages[45]. Two of these
stem cell types, maintained in intra-mammalian stages, proliferate and differentiate into precursors
of somatic and germ line cells throughout development[13]. 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 on 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 missing[34]. 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, *Sm*RAR, and related genes[66]. *Sm*RAR 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 *Sm*RAR pathway[66]. 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 demonstrated[5]. In addition, soluble factors produced by the worms with effect on the opposite sex cannot be ruled out[6]. 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 embryogenesis[67]. 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 mice[34]. 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 Sc*histosoma', 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 modifications[18]. In brief, patent mix strain of *Biomphalaria glabrata* snails[68] 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 g/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 medium[18] 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% CO_2 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
described[18]. 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 described[38] (**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 longer and thinner than males,
the ceca fusion localises closer to the anterior end of the worm (i.e., ~⅓ 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., ~⅔ 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≤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 (µm²). 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≤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 PCR[69] (**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 repeat[70] and actin[71] (**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) ccontaining 2µl 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

487 independent experiments. Statistical analysis was performed by Kruskal-Wallis test with Dunn
488 multiple comparison post-hoc test, with P≤0.05 considered significant (**Supplementary Table S6**).
489

Pairing experiments

Schistosoma mansoni adult male and female worms were collected from experimentally infected
mice at 47 days post infection[18]. Briefly, mice were euthanised by intraperitoneal injection of 200 μl
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
% CO_2 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^{TM} Green Actin Tracking Stain (Invitrogen) and CellMask^{TM} 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^{TM} Live
ReadyProbes^{TM} 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 μl 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.

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References

1   LoVerde, P. T. Schistosomiasis. *Adv Exp Med Biol* **1454**, 75-105 (2024).
https://doi.org/10.1007/978-3-031-60121-7_3

2 Buonfrate, D., Ferrari, T. C. A., Adegnika, A. A., Russell Stothard, J. & Gobbi, F. G.
Human schistosomiasis. *Lancet* **405**, 658-670 (2025). https://doi.org/10.1016/S0140-
6736(24)02814-9
3 Berger, D. J. *et al.* Extensive transmission and variation in a functional receptor for
praziquantel resistance in endemic. *bioRxiv* (2024).
https://doi.org/10.1101/2024.08.29.610291
4 Loker, E. S. & Brant, S. V. Diversification, dioecy and dimorphism in schistosomes. *Trends
Parasitol* **22**, 521-528 (2006). https://doi.org/10.1016/j.pt.2006.09.001
5 Chen, R. *et al.* A male-derived nonribosomal peptide pheromone controls female
schistosome development. *Cell* **185**, 1506-1520.e1517 (2022).
https://doi.org/10.1016/j.cell.2022.03.017
6 Shakir, E. M. N., Rinaldi, G., Kirk, R. S. & Walker, A. J. Schistosoma mansoni excretory-
secretory products induce protein kinase signalling, hyperkinesia, and stem cell proliferation
in the opposite sex. *Commun Biol* **6**, 985 (2023). https://doi.org/10.1038/s42003-023-05333-
9
7 Elkrewi, M., Moldovan, M. A., Picard, M. A. L. & Vicoso, B. Schistosome W-Linked
Genes Inform Temporal Dynamics of Sex Chromosome Evolution and Suggest Candidate
for Sex Determination. *Mol Biol Evol* **38**, 5345-5358 (2021).
https://doi.org/10.1093/molbev/msab178
8 Fitzpatrick, J. M. & Hoffmann, K. F. Dioecious Schistosoma mansoni express divergent
gene repertoires regulated by pairing. *Int J Parasitol* **36**, 1081-1089 (2006).
https://doi.org/10.1016/j.ijpara.2006.06.007
9 Lu, Z. *et al.* Schistosome sex matters: a deep view into gonad-specific and pairing-
dependent transcriptomes reveals a complex gender interplay. *Sci Rep* **6**, 31150 (2016).
https://doi.org/10.1038/srep31150
10 Picard, M. A. L. *et al.* 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 (2016).
https://doi.org/10.1371/journal.pntd.0004930
11 Wangwiwatsin, A. *et al.* Transcriptome of the parasitic flatworm Schistosoma mansoni
during intra-mammalian development. *PLoS Negl Trop Dis* **14**, e0007743 (2020).
https://doi.org/10.1371/journal.pntd.0007743
12 Diaz Soria, C. L. *et al.* Single-cell atlas of the first intra-mammalian developmental stage of
the human parasite Schistosoma mansoni. *Nat Commun* **11**, 6411 (2020).
https://doi.org/10.1038/s41467-020-20092-5
13 Li, P. *et al.* Single-cell analysis of Schistosoma mansoni identifies a conserved genetic
program controlling germline stem cell fate. *Nat Commun* **12**, 485 (2021).
https://doi.org/10.1038/s41467-020-20794-w
14 Wendt, G. *et al.* A single-cell RNA-seq atlas of identifies a key regulator of blood feeding.
Science **369**, 1644-1649 (2020). https://doi.org/10.1126/science.abb7709
15 Pance, A. & Rinaldi, G. Editorial: Stem cell technologies meet stem cell biology to shine
new light into tropical infectious diseases. *Front Cell Infect Microbiol* **14**, 1411728 (2024).
https://doi.org/10.3389/fcimb.2024.1411728
16 Sutrave, S. & Richter, M. H. The Truman Show for Human Helminthic Parasites: A Review
of Recent Advances in In Vitro Cultivation Platforms. *Microorganisms* **11** (2023).
https://doi.org/10.3390/microorganisms11071708
17 Ahmed, N. H. Cultivation of parasites. *Trop Parasitol* **4**, 80-89 (2014).
https://doi.org/10.4103/2229-5070.138534
18 Mann, V. H., Morales, M. E., Rinaldi, G. & Brindley, P. J. Culture for genetic manipulation
of developmental stages of Schistosoma mansoni. *Parasitology* **137**, 451-462 (2010).
https://doi.org/10.1017/S0031182009991211

579	19	Mellin, R. & Boddey, J. A. Organoids for Liver Stage Malaria Research. *Trends Parasitol*
580		**36**, 158-169 (2020). https://doi.org/10.1016/j.pt.2019.12.003
581	20	Zorrinho-Almeida, M., de-Carvalho, J., Bernabeu, M. & Silva Pereira, S. Leveraging
582		microphysiological systems to expedite understanding of host-parasite interactions. *PLoS*
583		*Pathog* **21**, e1013088 (2025). https://doi.org/10.1371/journal.ppat.1013088
584	21	Faral-Tello, P., Pagotto, R., Bollati-Fogolín, M. & Francia, M. E. Modeling the human
585		placental barrier to understand ´s vertical transmission. *Front Cell Infect Microbiol* **13**,
586		1130901 (2023). https://doi.org/10.3389/fcimb.2023.1130901
587	22	Korwin-Mihavics, B. R., Dews, E. A., di Genova, B. M. & Huston, C. D. Organoid-based
588		systems to model infection in 2D and 3D. *bioRxiv* (2023).
589		https://doi.org/10.1101/2023.09.29.560165
590	23	Zhou, Y. *et al.* Full maturation of in vitro Plasmodium falciparum oocysts using the
591		AlgiMatrix 3D culture system. *Malar J* **23**, 251 (2024). https://doi.org/10.1186/s12936-024-
592		05079-7
593	24	Britton, C. *et al.* New technologies to study helminth development and host-parasite
594		interactions. *Int J Parasitol* **53**, 393-403 (2023). https://doi.org/10.1016/j.ijpara.2022.11.012
595	25	Duque-Correa, M. A. *et al.* Defining the early stages of intestinal colonisation by
596		whipworms. *Nat Commun* **13**, 1725 (2022). https://doi.org/10.1038/s41467-022-29334-0
597	26	Faber, M. N. *et al.* Development of Bovine Gastric Organoids as a Novel Model to Study
598		Host-Parasite Interactions in Gastrointestinal Nematode Infections. *Front Cell Infect*
599		*Microbiol* **12**, 904606 (2022). https://doi.org/10.3389/fcimb.2022.904606
600	27	Vitkauskaite, A. *et al.* In vitro co-culture of Fasciola hepatica newly excysted juveniles
601		(NEJs) with 3D HepG2 spheroids permits novel investigation of host–parasite interactions.
602		*Virulence* (2025). https://doi.org/10.1080/21505594.2025.2482159
603	28	White, R., Blow, F., Buck, A. H. & Duque-Correa, M. A. Organoids as tools to investigate
604		gastrointestinal nematode development and host interactions. *Front Cell Infect Microbiol* **12**,
605		976017 (2022). https://doi.org/10.3389/fcimb.2022.976017
606	29	Herz, M. *et al.* Genome-wide transcriptome analysis of larvae and germinative cell cultures
607		reveals genes involved in parasite stem cell function. *Front Cell Infect Microbiol* **14**,
608		1335946 (2024). https://doi.org/10.3389/fcimb.2024.1335946
609	30	Koziol, U. & Brehm, K. Recent advances in Echinococcus genomics and stem cell research.
610		*Vet Parasitol* **213**, 92-102 (2015). https://doi.org/10.1016/j.vetpar.2015.07.031
611	31	Basch, P. F. Cultivation of Schistosoma mansoni in vitro. II. production of infertile eggs by
612		worm pairs cultured from cercariae. *J Parasitol* **67**, 186-190 (1981).
613	32	Basch, P. F. Cultivation of Schistosoma mansoni in vitro. I. Establishment of cultures from
614		cercariae and development until pairing. *J. Parasitol.* **67**, 179-185 (1981).
615	33	Basch, P. F. & Humbert, R. Cultivation of Schistosoma mansoni in vitro. III. implantation of
616		cultured worms into mouse mesenteric veins. *J Parasitol* **67**, 191-195 (1981).
617	34	Wang, J., Chen, R. & Collins, J. J., 3rd. Systematically improved in vitro culture conditions
618		reveal new insights into the reproductive biology of the human parasite Schistosoma
619		mansoni. *PLoS Biol* **17**, e3000254 (2019). https://doi.org/10.1371/journal.pbio.3000254
620	35	You, Y. *et al.* An improved medium for in vitro studies of female reproduction and
621		oviposition in Schistosoma japonicum. *Parasit Vectors* **17**, 116 (2024).
622		https://doi.org/10.1186/s13071-024-06191-y
623	36	Anisuzzaman *et al.* Host-Specific Serum Factors Control the Development and Survival of.
624		*Front Immunol* **12**, 635622 (2021). https://doi.org/10.3389/fimmu.2021.635622
625	37	Buchter, V., Schneeberger, P. H. H. & Keiser, J. Validation of a human-serum-based in vitro
626		growth method for drug screening on juvenile development stages of Schistosoma mansoni.
627		*PLoS Negl Trop Dis* **15**, e0009313 (2021). https://doi.org/10.1371/journal.pntd.0009313
628	38	Maharjan, S., Kirk, R. S., Lawton, S. P. & Walker, A. J. Further evaluation and validation of
629		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 (2021). https://doi.org/10.1016/j.ijpara.2020.12.009

Reimers, N. *et al.* Drug-induced exposure of Schistosoma mansoni antigens SmCD59a and SmKK7. *PLoS Negl Trop Dis* **9**, e0003593 (2015). https://doi.org/10.1371/journal.pntd.0003593

Abla, N. *et al.* Evaluation of the pharmacokinetic-pharmacodynamic relationship of praziquantel in the Schistosoma mansoni mouse model. *PLoS Negl Trop Dis* **11**, e0005942 (2017). https://doi.org/10.1371/journal.pntd.0005942

Milligan, J. N. & Jolly, E. R. Cercarial transformation and in vitro cultivation of Schistosoma mansoni schistosomules. *J Vis Exp* (2011). https://doi.org/10.3791/3191

Nation, C. S., Da'dara, A. A., Marchant, J. K. & Skelly, P. J. Schistosome migration in the definitive host. *PLoS Negl Trop Dis* **14**, e0007951 (2020). https://doi.org/10.1371/journal.pntd.0007951

Canepa, M., Ding, A. & Fechner, C. Cellular Adaptation: Hypertrophy, Hyperplasia, Atrophy. *General Pathology Student Guide*, 50-64 (2024). https://doi.org/10.1007/978-3-662-67962-3_3

Nanes Sarfati, D., Li, P., Tarashansky, A. J. & Wang, B. Single-cell deconstruction of stem-cell-driven schistosome development. *Trends Parasitol* **37**, 790-802 (2021). https://doi.org/10.1016/j.pt.2021.03.005

Wang, B. *et al.* Stem cell heterogeneity drives the parasitic life cycle of. *Elife* **7** (2018). https://doi.org/10.7554/eLife.35449

Arunsan, P. *et al.* Programmed knockout mutation of liver fluke granulin attenuates virulence of infection-induced hepatobiliary morbidity. *Elife* **8** (2019). https://doi.org/10.7554/eLife.41463

Kalinna, B. H., Ross, A. G. & Walduck, A. K. Schistosome Transgenesis: The Long Road to Success. *Biology (Basel)* **13** (2024). https://doi.org/10.3390/biology13010048

Quinzo, M. J., Perteguer, M. J., Brindley, P. J., Loukas, A. & Sotillo, J. Transgenesis in parasitic helminths: a brief history and prospects for the future. *Parasit Vectors* **15**, 110 (2022). https://doi.org/10.1186/s13071-022-05211-z

Stephens, D. R. *et al.* 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 (2025). https://doi.org/10.1073/pnas.2505710122

Xie, Y. *et al.* RNAi screening of uncharacterized genes identifies promising druggable targets in Schistosoma japonicum. *PLoS Pathog* **21**, e1013014 (2025). https://doi.org/10.1371/journal.ppat.1013014

Loker, E. S. REVIEW: THE ORIGIN OF DIGENETIC TREMATODES: DID THE STORY BEGIN WITH THE BLOOD FLUKES? *J Parasitol* **111**, 787-814 (2025). https://doi.org/10.1645/25-27

Moné, H. & Boissier, J. Sexual biology of schistosomes. *Adv Parasitol* **57**, 89-189 (2004). https://doi.org/10.1016/S0065-308X(04)57002-1

Puck, T. T., Cieciura, S. J. & Robinson, A. Genetics of somatic mammalian cells. III. Long-term cultivation of euploid cells from human and animal subjects. *J Exp Med* **108**, 945-956 (1958). https://doi.org/10.1084/jem.108.6.945

van der Valk, J. *et al.* Fetal Bovine Serum (FBS): Past - Present - Future. *ALTEX* **35**, 99-118 (2018). https://doi.org/10.14573/altex.1705101

Gstraunthaler, G., Lindl, T. & van der Valk, J. A plea to reduce or replace fetal bovine serum in cell culture media. *Cytotechnology* **65**, 791-793 (2013). https://doi.org/10.1007/s10616-013-9633-8

Weber, T. *et al.* Fetal bovine serum: how to leave it behind in the pursuit of more reliable science. *Front Toxicol* **7**, 1612903 (2025). https://doi.org/10.3389/ftox.2025.1612903

Rosolowski, J., Weber, T., Malakpour-Permlid, A. & Oredsson, S. Revisiting 3Rs: rethinking replacement and new approach methodologies. *Front Toxicol* **7**, 1664209 (2025). https://doi.org/10.3389/ftox.2025.1664209

Subbiahanadar Chelladurai, K. *et al.* Alternative to FBS in animal cell culture - An overview and future perspective. *Heliyon* **7**, e07686 (2021). https://doi.org/10.1016/j.heliyon.2021.e07686

Xiao, S.-H. & Sun, J. Schistosoma hemozoin and its possible roles. *Int J Parasitol* **47**, 171-183 (2017). https://doi.org/10.1016/j.ijpara.2016.10.005

Sun, J. *et al.* Hemozoin: a waste product after heme detoxification? *Parasit Vectors* **18**, 83 (2025). https://doi.org/10.1186/s13071-025-06699-x

Truscott, M., Evans, D. A., Gunn, M. & Hoffmann, K. F. Schistosoma mansoni hemozoin modulates alternative activation of macrophages via specific suppression of Retnla expression and secretion. *Infect Immun* **81**, 133-142 (2013). https://doi.org/10.1128/IAI.00701-12

Miller, P. & Wilson, R. A. Migration of the schistosomula of Schistosoma mansoni from the lungs to the hepatic portal system. *Parasitology* **80**, 267-288 (1980). https://doi.org/10.1017/s0031182000000743

Corrêa Soares, J. B. R. *et al.* Interference with hemozoin formation represents an important mechanism of schistosomicidal action of antimalarial quinoline methanols. *PLoS Negl Trop Dis* **3**, e477 (2009). https://doi.org/10.1371/journal.pntd.0000477

Frahm, S. *et al.* 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 (2019). https://doi.org/10.1371/journal.pntd.0006590

Collins, J. J., 3rd *et al.* Adult somatic stem cells in the human parasite Schistosoma mansoni. *Nature* **494**, 476-479 (2013). https://doi.org/10.1038/nature11924

Moescheid, M. F. *et al.* 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** (2025). https://doi.org/10.1093/nar/gkae1228

Doenhoff, M. J., Modha, J. & Walker, A. J. 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 (2019). https://doi.org/10.1016/j.ijpara.2019.05.004

Geyer, K. K. *et al.* 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 (2017). https://doi.org/10.1371/journal.pntd.0005246

Buddenborg, S. K., Lu, Z., Sankaranarayan, G., Doyle, S. R. & Berriman, M. The stage- and sex-specific transcriptome of the human parasite Schistosoma mansoni. *Sci Data* **10**, 775 (2023). https://doi.org/10.1038/s41597-023-02674-2

Grevelding, C. G. Genomic instability in Schistosoma mansoni. *Mol. Biochem. Parasitol.* **101**, 207-216 (1999). https://doi.org/10.1016/s0166-6851(99)00078-x

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

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.

Author ORCIDs

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

Mary K. Evans, https://orcid.org/0009-0001-9120-6017

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

Geetha Sankaranarayanan, https://orcid.org/

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, http://orcid.org/0000-0002-9581-0377

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

758 **Tables**

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

Reagent name	Final Concentration	Reagent source
DMEM, high glucose, sodium pyruvate		Fisher Scientific (Cat. 13476146)
Lactalbumin hydrolysate	1mg/ml	Merk Life Sciences (Cat. 61300-500G)
Hypoxanthine	500nM	Merk Life Sciences (Cat. H9636-1G)
Serotonin	1μM	Merk Life Sciences (Cat. H9523)
Hydrocortisone	1μM	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 Foetal Bovine Serum	20%	HS: NHSBT serum (Cat. NC02) FBS: Fisher Scientific (Cat. 11550356)
Antibiotic antimycotic solution	2X	Fisher Scientific (Cat. 15140-122)
Insulin solution	8μg/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)

Figure legends

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 (µm²) ± SEM. HS: Human Serum (light brown); FBS: Foetal Bovine Serum (blue). *: P-value < 0.01 in indicated pairwise comparisons (Supplementary Table S6).

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. (Statistical analyses in Supplementary Table S6.) **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≤0.05 (*) considered significant (Supplementary Table S6). **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).

Supplementary information

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 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 µm.

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 µm.

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 µm **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 µm.

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 µm.

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 µm.

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 µm.

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 µm.

Supplementary Video S4. Representative video of *in vitro* developed females and males *in copula* at ~80 days in culture. Scale bar: 100 µm

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 µm

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 µm (top panel) and 50 µm (bottom panel)

Figure 1

Figure 2

Figure 3

A

B

Figure 5

Figure 6