

Schistosome vaccines: a critical appraisal

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*An effective schistosome vaccine is a desirable control tool but progress towards that goal has been slow. Protective immunity has been difficult to demonstrate in humans, particularly children, so no routes to a vaccine have emerged from that source. The concept of concomitant immunity appeared to offer a paradigm for a vaccine operating against incoming larvae in the skin but did not yield the expected dividends. The mining of crude parasite extracts, the use of monoclonal antibodies and protein selection based on immunogenicity produced a panel of vaccine candidates, mostly of cytoplasmic origin. However, none of these performed well in independent rodent trials, but glutathione-S-transferase from *Schistosoma haematobium* is currently undergoing clinical trials as an anti-fecundity vaccine. The sequencing of the *S. mansoni* transcriptome and genome and the development of proteomic and microarray technologies has dramatically improved the possibilities for identifying novel vaccine candidates, particularly proteins secreted from or exposed at the surface of schistosomula and adult worms. These discoveries are leading to a new round of protein expression and protection experiments that will enable us to evaluate systematically all the major targets available for immune intervention. Only then will we know if schistosomes have an Achilles' heel.*

Key words: antigen - immune response - formulation - protection

For a variety of reasons, schistosomiasis remains an important parasitic disease of humans, its treatment and hence control being reliant on a single compound, praziquantel. An effective vaccine would be a useful adjunct and if sufficiently potent, a replacement for chemotherapy but the development of such a product has proved elusive. This review, primarily on *Schistosoma mansoni*, was compiled at the invitation of the organising committee and delivered at the Xth International Symposium on Schistosomiasis.

Why develop a schistosome vaccine?

It is pertinent at the outset to ask why we should want a schistosome vaccine – would it offer any advantages over chemotherapy? A strong argument has to be that schistosome infections are cryptic; they frequently go undetected and significant pathology has developed before chemotherapy is administered. Indeed, recent studies suggest that both morbidity and mortality have been seriously underestimated (van der Werf et al. 2003, King et al. 2005), especially what is sometimes referred to as subtle morbidity. It should also be borne in mind that single-dose chemotherapy is not completely effective in all patients. Praziquantel is inactive against the developing parasites and its efficacy at sub-curative levels may rely on host immune responses (Doenhoff et al. 1987). Furthermore, chemotherapy cannot prevent re-infection.

Low cure rates have been reported that may presage the appearance of drug resistance (Doenhoff et al. 2002). Lastly, schistosome infections are potent immunomodulators, a process that appears related to worm burden (Silveira et al. 2004), and may alter host responses to other infections in a deleterious manner.

It is also pertinent to ask what we want from a schistosomiasis vaccine. Unlike viral, microbial, and protozoan pathogens, schistosomes do not multiply in the mammalian host. The eggs they deposit in host tissues are the principal cause of pathology, which develops roughly in proportion to parasite burden. This means that sterile immunity induced by a schistosome vaccine is not essential. A significant reduction in worm burden would be a useful attribute as would a vaccine-induced reduction in the fecundity of female worms or the viability eggs.

The first obstacle

It is axiomatic that if a pathogen, however virulent, induces protective immunity as a result of an active infection, then the development of a vaccine against it is a feasible proposition. Schistosomes do not measure up well against this yardstick. The adult worms are adapted to life in the human bloodstream, a very hostile environment, where they can survive for > 30 years (Harris et al. 1984) so must possess very effective immune evasion strategies. The standard experimental design to investigate human immunity, which relies on curative chemotherapy to permit reinfection rates to be determined, reveals little evidence for protection before puberty. Lower reinfection rates have been observed in adults, but it is by no means clear that they are solely the product of acquired immunity. Moreover, whilst there have been many studies of human immune responses, no obvious immune mechanisms have emerged on which to base a vaccine. This is especially true of mechanisms invoking IgE in effector responses. Given current knowledge of immunoglobulin heavy chain class switching, it is diffi-

Financial support: The Wellcome Trust, Biotechnology and Biological Sciences Research Council, National Institutes of Health, the UNICEF/UNDP/World Bank/WHO Special Programme for Training in Tropical Diseases

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Received 25 May 2006

Accepted 26 June 2006

cult to envisage how antigens could be administered to elicit schistosome-specific IgE without also triggering bystander responses to irrelevant (environmental) antigens, with consequent risk of allergy. It is apparent that developing a schistosome vaccine for use in humans was never going to be easy.

An early disappointment

The first attempts to develop a schistosome vaccine began half a century ago. By analogy with successful microbial and viral vaccines, they involved the vaccination of mice with crude worm extracts or purified components, followed by a cercarial challenge (e.g. Sadun & Lin 1959, Murrell et al. 1975, Hayunga 1985). The results were equivocal with 20, 30, and even 50% reduction in worm burden recorded, but there was a lack of consistency, even in the same laboratory. It seemed apparent that crude extracts were inadequate vaccines. Perhaps there were a few key antigens that needed to be identified, but this begs the question about the mechanisms that a vaccine was intended to elicit. Would the possession of good models of protection assist the process of antigen identification?

Concomitant immunity: the big idea 1969-1979

The concept of concomitant immunity, introduced by Smithers and Terry (1969), dominated thinking about schistosome immunology and vaccine research for two decades. These researchers reported that when the rhesus macaque (*Macaca mulatta*) was given an infection, the primary worm burden persisted but the animals were resistant to a secondary cercarial challenge. McCullough and Bradley (1973) tentatively suggested that the concept could be applied to *S. haematobium* infections in humans and the idea became firmly established in the minds of researchers. It posed a very simple and elegant paradigm: discover how the mature primary worms induce protection and you have the basis for a vaccine. Almost simultaneously, the description of the in vitro killing of newly transformed schistosomula by a combination of antibodies, complement and eosinophils (Butterworth et al. 1974), termed antibody-dependent cellular cytotoxicity (ADCC), provided a mechanism to explain how concomitant immunity might operate. Subsequent research showed that several combinations of leucocytes and antibody isotypes were equally effective at parasite killing in vitro (Butterworth 1984).

The mid 1970s was one of those periods of optimism that seem to characterise vaccine development, where the product appears to be within our grasp and just a little more research will bring it to fruition. Thirty years later, with no product near to application, it is pertinent to ask what went wrong. One reason is that work on the rhesus macaque model ceased, before it had been fully characterised, and experimentation shifted to the laboratory mouse. We can speculate that factors such as cost and the need for adequate replication of samples were important criteria driving this change. It was shown both that chronically infected mice could develop concomitant immunity (Dean et al. 1978a), and that the protection could be passively transferred to naïve mice with infection se-

rum (Sher et al. 1975). However, the mouse with a chronic infection was always a tricky model, since even a single worm pair could elicit significant morbidity and mortality.

This led researchers to question the immunological basis of the concomitant immunity displayed by chronically infected mice, initially by demonstrating that it required a bisexual infection (Dean et al. 1978b). This implicated eggs as causative agents yet eggs alone, whilst eliciting high antibody titres, could not induce protection (Harrison et al. 1982). The prosaic explanation for concomitant immunity in the mouse was that it represented an artefact of pathology (Wilson et al. 1983). The well-documented egg-induced pathology causes porta-caval anastomoses to form that allow schistosomula, newly arrived in the portal system, to escape back to the general circulation. Was this also the explanation for concomitant immunity in the rhesus macaque? Recent experiments with this model suggest not (Coulson, unpublished data) and it would be instructive to take another look using modern immunological techniques. However, it seems that the concept of a primary worm burden remaining whilst the host was immune to a challenge was misplaced, with the rhesus macaque's response to infection best being described as a "self cure" phenomenon.

Attenuated parasites induce protective immunity

Whilst scepticism was growing about the possibility of basing a vaccine on the "immunity" displayed by hosts with a chronic infection, research on the induction of immunity using attenuated cercariae was gathering momentum. Vaccination with attenuated cercariae was first attempted in the early 1960s, with detailed analysis being undertaken from 1978 onwards (Minard et al. 1978). These studies have demonstrated that high levels of protection can be achieved in mice (>90%) and primates (86%) (Wynn et al. 1995, Mountford et al. 1996, Kariuki et al. 2004), and provide the strongest possible evidence that a schistosome vaccine is feasible. The salient features of this attenuated vaccine have been reviewed (Coulson 1997) and, in mice at least, its success appears to require a truncated migration of the attenuated parasites. A small proportion of the larvae enters and persists in the skin-draining lymph nodes where they provoke an immune response greater in intensity and differing in quality from that induced by normal parasites. Other larvae travel only as far as the lungs where they recruit lymphocytes to arm that organ (Coulson & Wilson 1997).

Both cell-mediated and humoral mechanisms appear to operate, but consecutively rather than simultaneously. Thus, in 1× vaccinated mice lung schistosomula are the target of focal cellular effector responses mediated by CD4⁺ T cells of the Th1 subset, with interferon γ (INT- γ) and tumor-necrosis factor (TNF) as key cytokines. The bulk of evidence suggests that the effector mechanism relies on blocking intravascular migration rather than cytotoxic killing. With multiple exposures of mice to attenuated cercariae, the immune response progressively develops a Th2 profile, with protection passively transferable to naïve recipients with immune serum. However, the level of protection attained with antibodies is never as great as for the cell-mediated mechanism. The immune responses

of primates to the attenuated vaccine well illustrate this switch, with the cytokine profile changing from Th1 to Th2 with successive exposures to attenuated cercariae (Eberl et al. 2001). In addition, multiple vaccinations are necessary to elicit adequate protection, and in 5× vaccinated baboons the mechanism is probably antibody-mediated (Kariuki et al. 2004). The big challenge has been, and remains, to identify the antigens that mediate protection in the irradiated vaccine model.

Routes to antigen identification 1980-1995

The attempts to identify protective antigens, whether based on a particular model, or opportunistically, have tracked new developments in technology, but usually with a lag of several years, as illustrated below:

1975: hybridoma technology developed (Kohler & Milstein 1975)

1982: first anti-schistosome mAbs (Zodda & Phillips 1982)

1974: recombinant DNA technology (Berg et al. 1974)

1982: first application of cloning techniques to development of a synthetic vaccine against schistosomiasis (Knopf et al. 1982)

1986: polymerase chain reaction (Mullis et al. 1986)

1991: first application to schistosome antigen cloning (Ali et al. 1991)

A full appraisal of the avenues that have been used to select potentially protective schistosome antigens can be found in Wilson and Coulson (2006) and only the barest outline is given here.

i) Mining crude extracts

Soluble extracts of schistosomula and adult worms, administered to mice, with BCG as adjuvant (James & DeBlois 1986), led to the identification of a 97 kDa antigen, paramyosin (Pearce et al. 1988), used in its native form for vaccination experiments.

ii) Monoclonal antibody (mAb) targets

When first developed, mAb technology seemed to provide a very direct route to the identification of protective antigens. Develop a panel on mAbs, test them for protective potential either directly via ADCC assays or by passive administration to mice, and then identify and clone the antigenic targets. A lot of laboratories followed this route, and as we can testify from personal experience, results were meagre. Nevertheless, three groups reported success, and this led to the development of triose phosphate isomerase (Harn et al. 1992), Sm23 (membrane tetraspanin; Reynolds et al. 1992) and 9B-Ag (Hazdai et al. 1985) as vaccine antigens.

iii) Anti-idiotypes

The development of anti-idiotypic vaccines was a briefly fashionable approach, especially since it offered a way to progress glycan epitopes, not amenable to recombinant DNA technology. Antigens of 38kDa (Grzych et al. 1985) and 68kDa (Olds & Kresina 1987) on the surface of schistosomula were selected in this way, but the approach petered out.

iv) Expression library screening

Molecular biology seemed to offer the best prospect for a vaccine, with unlimited quantities of antigens produced as recombinants in microbial or eucaryotic vectors. The obvious way to identify protective antigens was to screen expression libraries with immune serum. Two antigens emerged from this approach, IrV5, a cloned 62kDa fragment of myosin heavy chain (Soisson et al. 1992) and a 37kDa antigen target of immune human serum (Dessein et al. 1988) later identified as glyceraldehyde 3 phosphate dehydrogenase (GAPDH; Goudot-Crozel et al. 1989).

v) Immunogenicity

The straightforward use of immunogenicity as the criterion for selection as a vaccine candidate was popular, not least because it made cloning easy, although it rather ignored Waksman's postulate that antigens mediating protection will have been selected by immunological pressure for low reactivity. A 28kDa antigen (Balloul et al. 1985), later identified as glutathione-S-transferase (Sm28GST; Taylor et al. 1988) and fatty acid binding protein (Sm14; Moser et al. 1991) emerged by this route.

The one feature that all the above antigens had in common was an ability to elicit 30-40% protection, with occasional reports of much higher values, but set against this were papers reporting immunogenicity but not protection data. Since levels of protection elicited by a single antigen were low, several attempts were made to boost vaccine efficacy by combining antigens as protein cocktails, or their epitopes in DNA and synthetic peptide constructs. Unfortunately, no additive effect was recorded, a result that could be interpreted to mean that the various antigens involved (Sm28GST, TPI, Sm23, paramyosin) all conferred protection via the same mechanism.

Is antigen formulation the problem?

A plausible explanation for the limited success of the recombinant antigen vaccines was incorrect or inadequate formulation to achieve the desired level of protection. It was demonstrated independently that co-administration to mice of the radiation-attenuated cercarial vaccine with the recombinant cytokine IL (interleukin)-12, would increase its efficacy from 60 to > 90%, with some animals showing sterile immunity (Wynn et al. 1995, Mountford et al. 1996), so clearly formulation was an important issue. A great deal of work was published on the utility of various formulations and this is well illustrated by Sm28GST as the test antigen. Professor Capron's laboratory made a formidable effort with:

- synthetic peptides coupled to tetanus toxoid (Auriault et al. 1988) or assembled in an octomer (Wolowczuk et al. 1991);
- cloning into live *Salmonella* (Khan et al. 1994), BCG (Kremer et al. 1996) and *Bordetella pertussis* (Mielcarek et al. 1997);
- recombinant GST linked to cholera toxin delivered intranasally (Sun et al. 1999) or in biodegradable particles (Baras et al. 1999);

- a DNA construct administered with an IL-18 plasmid (Dupre et al. 2001).

Where protection data were published, it appears that no formulation gave a spectacular increase in efficacy.

The independent WHO vaccine trials

By the early 1990s, progress in expressing recombinant antigens and reports of successful protection in rodents, produced a spirit of optimism in the research community that a vaccine was feasible. This led WHO to provide funds for independent laboratory trials of candidate antigens, and calls were made for researchers to submit their antigens for testing. Two laboratories were selected for this task, The University of York, UK and BMRI, Rockville, MD, US. The antigens proffered were: IrV5, Sm28GST and Sm14 as recombinants, paramyosin as the purified native protein, and TPI and Sm23 as synthetic peptide constructs. The formulations were defined by the antigen donors as the ones that worked best in their laboratories, which precluded blind testing. The results of the trials were never made public but the WHO reported that “The consolidated results demonstrated that the stated goal of consistent induction of 40% protection or better was not reached with any of the antigen formulations tested in these trials” (TDR Newsletter, June 1996).

Clinical trials of schistosomiasis vaccines

Surprisingly, an antigen NOT involved in the WHO trials, Sh28GST, has emerged as the pacemaker for schistosome vaccines and has progressed through Phase I and Phase II clinical trials in humans (Capron et al. 2002). Its development is based on vaccination experiments with Patas monkeys (*Erythrocebus patas*) where a clear antifecundity effect was reported (Boulanger et al. 1995). If the effect could be replicated in humans, it would hopefully diminish urinary tract pathology and transmission. No adverse side effects were observed in human recipients and high titres of antibodies that neutralised Sh28GST activity were elicited in Phase I and Phase II trials (Capron et al. 2005). Phase III trials are scheduled to begin in Senegal in 2006 (Riveau, pers. commun.).

Why are high levels of protection difficult to achieve?

Although the attenuated cercarial vaccine can elicit > 80% protection in rodents and primates, no recombinant vaccine comes anywhere near this efficacy. The levels achieved by the many antigens tested seldom exceed 50% and average ~30% (it is also important to remember that journal editors are not interested in publishing negative results). We have already considered antigen formulation – are there other possibilities. For example, do we have the right antigens?

With the exceptions of Sm23 that could have exposed domains at the tegument surface, most of the antigens tested in the WHO trials, and by other workers, are cytosolic or cytoskeletal components. Thus, paramyosin and IrV5 (myosin) are muscle proteins, although there is controversy as to whether the former is present on the tegument surface (reviewed in Skelly & Wilson 2006). TPI and GAPDH are glycolytic enzymes and hence cytosolic in

location, as are GST (a detoxifying enzyme) and Sm14 (a fatty acid binding protein). In the first proteomic study undertaken on schistosomes, soluble extracts of adult worms (SWAP) were shown to contain several vaccine candidates among the top 50 spots on a 2D gel separation (Curwen et al. 2004). These authors concluded that “it seems counterintuitive that such proteins would form the basis for an effective vaccine”. This brings us back to the early trials with crude extracts (Murrell et al. 1975) – were they equivalent to a cocktail of recombinants with very similar results? Further, assuming that the internal antigens really do elicit a degree of protection, how might they trigger the immune effector mechanism?

There are several plausible explanations. It is surprising that the occurrence of ADCC *in vivo* against newly penetrated skin schistosomula has never been convincingly demonstrated. However, after *in vitro* transformation several candidate antigens (Sm28GST, IrV5, Sm23, 9B-Ag, TPI) have been reported as transiently detected on the parasite surface. Their presence coincides with shedding of the cercarial tegument membranes *in vivo* (McLaren & Hockley 1976), and could reflect a transient leakiness of the tegument surface that would explain their vulnerability to ADCC. It should be noted that by 24 h *in vitro* (and presumably *in vivo*) schistosomula are refractory to killing. However, it seems that schistosomes have adapted to this potential ambush, since the vast majority do not enter the dermis and blood vessels until > 48 h after skin penetration. Does this delay place them beyond the reach of antibodies and leucocytes until their immune evasion mechanisms are in place?

A second explanation, not involving ADCC, has been highlighted by proteomic analysis of the acetabular gland secretions of cercariae (Curwen et al. 2006). The contents of pre- and post-acetabular glands are released by holocrine secretion which means that the entire contents of the gland cell are squeezed out, cytoplasm as well as vesicles. Thus, Cyclophilin, Thioredoxin, GST28, Aldolase, TPI, and Sm14 were detected on 2D gels in descending order of concentration. Three of the WHO vaccine candidates are prominent in this list, but an effector response triggered by secreted antigens would most likely involve blockage of migration by an inflammatory response rather than ADCC (cf. pulmonary effector responses in mice after exposure to the attenuated-cercaria vaccine).

A possible limitation of ADCC was actually put forward by proponents of the mechanism, namely induction of the wrong kind of antibody. *In vitro*, evidence was obtained that some antibody isotypes (e.g. IgM) can block ADCC (Yi et al. 1986), whilst *in vivo* in humans, it has been suggested that IgG4 can block IgE-mediated mechanisms (Hagan et al. 1991). There is another limitation of ADCC as a vaccine-induced effector mechanism that has seldom been considered, namely the need for permanent high specific antibody titres because there is no time to mount a secondary response before the penetrating larva loses its susceptibility to attack. The problem is how to maintain such titres over periods of months to years using current vaccine technologies. In our personal opinion antigens eliciting antibodies that mediate ADCC are never

likely to form more than a component of a multivalent schistosome vaccine.

What are the right antigens?

In mice given a single exposure to the attenuated cercarial vaccine, the dependence of the effector response on CD4⁺ T cells means that the relevant antigens must be accessible for processing and presentation by accessory cells. In other words they must be secreted by migrating larvae, or exposed on their epithelial surfaces. Furthermore, the ability of rhesus macaques to eliminate established adult worms appears to rely on slow starvation (PS Coulson, unpublished data) and we hypothesise that gut and tegument function is impaired. Again this implies secreted or membrane antigens as targets. By what possible immune mechanisms could such antigens mediate protective responses?

In the lungs of mice exposed to attenuated cercariae where cell-mediated immunity operates, migration blocked by a tight focus of leucocytes is our favoured hypothesis, and if IFN γ is neutralised or TNF signalling is absent, the mechanism fails (Smythies et al. 1992, Street et al. 1999). Furthermore, when mice lacking an IFN γ receptor, which cannot mount a cell mediated response, are given 3-5 exposures to attenuated cercariae, their serum will confer passive protection on mice depleted of Complement or lacking a FcR γ chain (P. Coulson, unpublished data). This implies that neither complement fixation nor leucocyte attachment to the parasite is involved in the antibody-mediated protection in these defective hosts. Do such protective antibodies operate by enzyme neutralisation, or blocking of receptor/transporter function on tegument and gut epithelia?

How do we identify secreted and surface-exposed proteins?

The obvious route might appear to be to collect the secretions released by lung schistosomula cultured in vitro, raise antisera against them and screen expression libraries. When we tried this approach, the antigens detected were the same highly immunogenic constituents such as paramyosin, myosin and aldolase found by others (Harrop et al. 1999, 2000). The reason seems to be that even a few damaged or dead larvae in a culture contribute sufficient material to dominate the reactivity of the antibodies elicited, and hence of the clones picked from the expression library. Clearly this is not the way forward.

The sequencing of the *S. mansoni* transcriptome (Verjovski-Almeida et al. 2003) and genome (www.SchistoDB.org) has opened up exciting new possibilities for antigen discovery. These resources allow us to examine gene and protein expression in the target larval and adult worm stages. Such studies are sometimes referred to disparagingly as “fishing expeditions”, but at this stage of schistosome vaccine research there is simply no alternative if we are to break new ground. Provided such screens are carried out intelligently, with a clear strategy to select a specific subset of the parasite transcriptome or proteome, then they will deliver much new information. Our analysis of the immune responses involved in protective immunity in authenticated animal models has led us to concentrate on the secreted and surface-expressed

subset of larval and adult worms.

The proteome has been defined as the total protein complement of an organism, tissue, cell or sub-cellular fraction. Proteomics provides a way to link an individual protein to its encoding DNA sequence, and the first step in the process is to separate the complex mixture of proteins under investigation (Ashton et al. 2001). Reproducible 2D electrophoresis was made possible by immobilised pH gradients and is ideal for soluble proteins such as larval secretions. Protein spots are excised from the gel, trypsinised and subjected to tandem mass spectrometry. The spectra obtained can then be linked to encoding DNA by searching for matches in the transcript and genome databases. Using this approach we have identified ~80% by volume of the proteins in cercarial acetabular gland secretions (Curwen et al. 2006). Proteases and their inhibitors found included three isoforms of elastase (Salter et al. 2002). SmPepM8 (a novel metalloproteinase), dipeptyl peptidase IV (SmDPP IV) and SmSerp_c (a new serine protease inhibitor). A group of potential immunomodulators included Sm16 (Ramaswamy et al. 1995), three wasp venom homologues (SmSCP_a, b and c) and SmKK7 (a protein with homology to a potassium-channel blocker in scorpion venom). All the new proteins found here represent potential vaccine candidates on the basis of their secretion into host skin, and their biological functions in the infection process are also of considerable interest. Collectively they represent the major targets in acetabular gland secretions and so merit a full exploration.

We are applying the same approach to analyse the secretions of skin and lung stage larvae, a tougher proposition as the amount of material available is much less, straining the detection limits of mass spectrometry. We have also characterised the surface of the adult worm tegument (Braschi & Wilson 2006, Braschi et al. 2006a, b), as a prelude to the same approach with lung schistosomula, the targets of the immunity elicited in mice by attenuated cercariae. The proteins regurgitated by the adult worm in the course of blood feeding have also been investigated (Hall 2005). Potentially, given time, resources and ingenuity it should be possible to define the full range of secreted and surface-exposed proteins of larvae and adults that might serve as vaccine candidates. The greater task is in fact to clone such proteins in a correctly folded and functional state for vaccine trials in mice and, then for the most promising, in baboons that represent a realistic primate model.

The current sensitivity of mass spectrometers makes low abundance proteins difficult to identify. However, possession of the transcriptome and genome data opens up another route to the identification of vaccine candidates. It permits the construction of microarrays for subsets of the transcriptome, with which to explore e.g. stage-specific gene expression, and these data can be combined with *in-silico* analysis to pinpoint genes encoding secreted or membrane proteins. We have used the approach to compare gene expression in lung schistosomula, relative to six other life cycle stages, enabling us to pinpoint around 30 possible candidates, many with no homology to any known protein (Dillon et al. 2006). In our hands the

technique appears to be one to two orders of magnitude more sensitive than proteomics but it does suffer one disadvantage. Schistosomes are metazoan parasites with well developed tissues and organs so that the identification of a secreted or membrane antigen in a lung worm tells us nothing about whether it is likely to be accessible to the host immune system in a live worm, or sequestered deep in some internal location.

Conclusions

As in the period 1980-1990, once again new technologies and information open up the prospect of progress towards the elusive goal of an effective schistosome vaccine. Access to the genome, transcriptome and proteome provide a fantastic opportunity to search for new vaccine candidates. However, we should always bear in mind that schistosomes are not stupid. They have had tens of millions of years to evolve mechanisms that help them survive immune attack from the mammalian host, even an attack orchestrated by our vaccine strategies. Schistosomes are truly a formidable adversary that won't easily be beaten.

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