



Pentamidine uptake and resistance in pathogenic protozoa: past, present and future

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Diamidines, and pentamidine in particular, have a long history as valuable chemotherapeutic agents against infectious disease. Their selectivity is due mostly to selective accumulation by the pathogen, rather than the host cell; and acquired resistance is frequently the result of changes in transmembrane transport of the drug. Here, recent progress in elucidating the mechanisms of diamidine transport in three important protozoan pathogens, *Trypanosoma brucei*, *Leishmania* and *Plasmodium falciparum*, is reviewed, and the implications for drug resistance are discussed.

Diamidines have a long history of involvement in the treatment of human protozoan infections, and pentamidine (PMD) is still the drug of choice against early stage sleeping sickness. The first antiprotozoan diamidine, synthalin (actually a diguanidine, see Fig. 1) was developed as a synthetic analogue of insulin, and tested in experimental trypanosome infections on the basis that induced hypoglycaemia was expected to control or cure trypanosome infections in humans [1]. However, Lourie and Yorke discovered in 1937 that synthalin was as active against trypanosomes *in vitro* as trivalent arsenicals. This led to the synthesis and testing of a large number of related compounds, the most active of which was diamidino-1,11-*n*-undecane [2], and later to the aromatic diamidines, including stilbamidine, propamidine and PMD by A.J. Ewins of the pharmaceutical company May and Baker [3]. These compounds were found to be highly effective against first stage (and early second-stage, see Box 1) human *Trypanosoma brucei gambiense* infections [4] and Indian kala azar [5], although stilbamidine was found to cause serious neurological effects in some patients [6] and was later abandoned. Importantly, PMD and propamidine were proven to be effective against strains resistant to the aromatic arsenicals then in use, such as tryparsamide [7], which had developed as a result of widespread use of these drugs during the preceding 15 years. Resistance to PMD itself has never been a significant problem in the field, despite its widespread use as a prophylactic (Box 2).

In addition to their use as drugs against African trypanosomes, diamidines have been used clinically against

various other infectious diseases including *Pneumocystis carinii* pneumonia [8] and leishmaniasis. Various studies, in the decade following the first diamidine synthesis, also found antibacterial, antifungal, antineoplastic, antimalarial and antileishmanial activity (reviewed in Ref. [9]).

PMD uptake in African trypanosomes

PMD uptake in African trypanosomes was first studied by Damper and Patton [10,11], who concluded that high-affinity transporters were involved, and that the parasite continued actively to accumulate drug until a lethal intracellular concentration had been reached. This concentration appeared to be in excess of 1 mM, but it is not clear to what extent this reflects free PMD concentrations in the cell because PMD binds tightly to DNA and might be segregated in an intracellular compartment such as the mitochondrion, as fluorescent diamidines such as stilbamidine have been shown to be associated with kinetoplast DNA (kDNA) and promote linearization of kDNA minicircles [12].

Carter *et al.* [13] provided strong evidence that PMD uptake by *T. b. brucei* is mediated by the P2 aminopurine transporter, which is also implicated in the uptake of melaminophenyl arsenicals [14,15], by showing that PMD inhibited both P2-mediated adenosine transport and prevented melarsen-oxide-induced cell lysis *in vitro*. The apparent inconsistency of a selective aminopurine-only transporter efficiently mediating the uptake of diverse molecules such as melaminophenyl arsenicals and diamidines was overcome with the elucidation of the structural motif recognized by this transporter [16,17]. However, adenosine did not inhibit [³H] PMD transport by *T. b. brucei* to the expected extent [13]. Recently, it has become clear that additional transporters are capable of mediating PMD uptake in bloodstream forms of *T. brucei* (Fig. 2). No fewer than three distinct transporters are capable of mediating PMD uptake, as measured by transport of [³H] PMD [18] and [¹²⁵I] iodopentamidine [19], with K_m values ranging over three orders of magnitude (Table 1).

Only one of these transporters was sensitive to inhibition by adenosine, and was hence designated adenosine-sensitive pentamidine transporter (ASPT)1. There is strong kinetic evidence that ASPT1 is identical to the P2 adenosine transporter. First, the ASPT1-mediated

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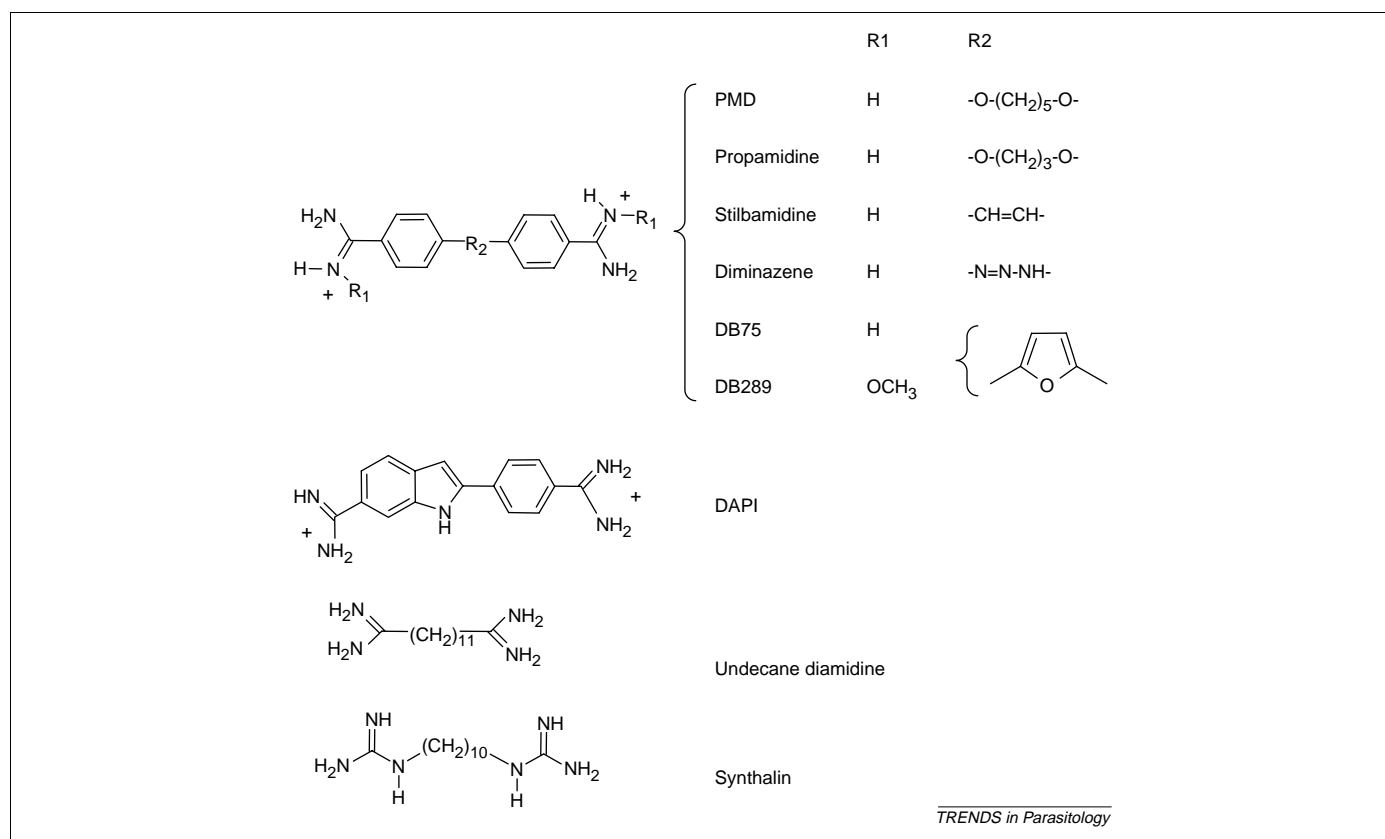


Fig. 1. Structure of various diamidines and related compounds. Structures of the classical aromatic diamidines PMD, propamidine, stilbamidine and diminazene aceturate, of the first orally available diamidine, DB289 and its active metabolite, DB75, of DAPI and of the early diamidine undecane diamidine and the diguanidinium compound synthalin. Abbreviations: DAPI, 4,6-diamidino-2-phenylindole; PMD, pentamidine; R₁, functional group substituted on amidine group; R₂, chain or ring forming the central part of the diamidine.

transport of both [³H] PMD and [¹²⁵I] iodopentamidine is inhibited by adenine and adenosine (but not by other purines, pyrimidines, pterines and amino acids), with K_i values that are virtually identical to the corresponding K_i values for P2-mediated adenosine transport [18,19].

Box 1. Diamidines in late-stage sleeping sickness

The main limitation to diamidine use against *Trypanosoma brucei rhodesiense* infections seems to be the far more rapid invasion of the cerebrospinal fluid (CSF) by this parasite. Pentamidine (PMD) is not prescribed against late-stage sleeping sickness, presumably because it does not cross the blood–brain barrier efficiently enough to attain the necessary drug levels in the central nervous system (CNS), although early trials do report considerable efficacy of PMD and propamidine against sleeping sickness with a moderate degree of change to the CSF [63,64] and PMD was found to be equally effective as tryparsamide in these cases [64]. These findings, now largely forgotten, show that diamidines are capable of crossing the blood–brain barrier. The high potential for diamidine use against late-stage sleeping sickness was demonstrated in a much more recent study by Doua *et al.* [65]. These authors showed that early late-stage *Trypanosoma brucei gambiense* human African trypanosomiasis (HAT) could be treated with ten intramuscular injections of 4 mg kg⁻¹, with a cure rate of 94%. It is of great importance to develop novel diamidines with improved CNS permeation, possibly administered as prodrugs with the amidine groups masked. A good example of this approach is the orally available experimental drug DB289, now being clinically tested against early stage HAT, and its derivatives are currently in development [66].

Likewise, the P2 transporter is inhibited by PMD and iodopentamidine, with K_i values almost identical to the K_m values for ASPT1-mediated PMD transport (Table 1). K_i values for other diamidines (diminazene aceturate) and melaminophenyl arsenicals are also identical for P2 and ASPT1 [17,18]. However, the kinetic similarities alone can offer no formal proof that the adenosine-sensitive PMD transport activity is mediated by the same transporter

Box 2. Pentamidine as a prophylactic

Field trials for the prophylactic use of pentamidine (PMD) were initiated in 1942 by Van Hoof and co-workers. It appeared that a single injection of 4 mg kg⁻¹ protected against *Trypanosoma gambiense* trypanosomiasis for at least six months [67], and a massive programme of PMD prophylaxis was started in the then Belgian and French African colonies and Nigeria, running until the late 1960s [68]. While this program was spectacularly successful in reducing the incidence of sleeping sickness, virtually eradicating the disease from Afrique Occidentale Française and Afrique Equatoriale Française [69], the initial high hopes of eradicating the disease were never realized (see Ref. [70] for review). Despite massive PMD use, regularly treating millions of people during the 1950s and 1960s, there were virtually no reports of induced PMD resistance. The 1976 WHO report on African trypanosomiasis mentions resistance to arsenicals and to diminazene aceturate (berenil) in cattle, but not to PMD [71]. PMD resistance was not detected in the comprehensive survey by the Organization de Coordination pour la Lutte contre les Endemies en Afrique Centrale in 1967, in countries where intense ‘pentamidinization’ had been practiced for over a decade [68].

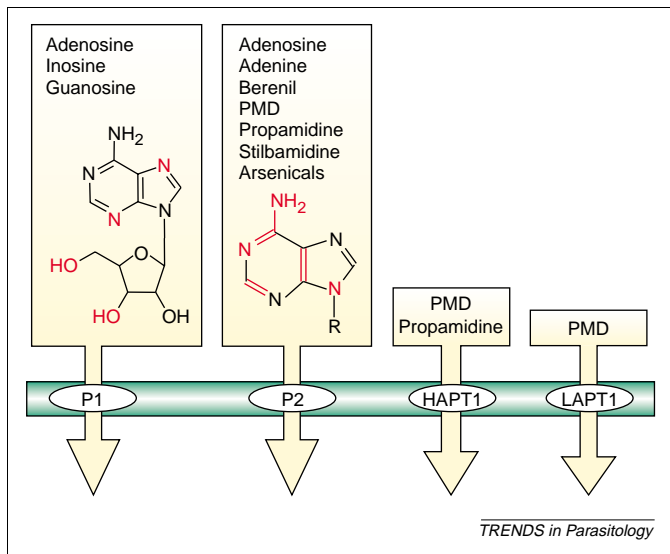


Fig. 2. Nucleoside and diamidine transporters in *Trypanosoma brucei brucei*. The substrate recognition motifs of the P1 and P2 transporters are indicated in red. P1 interacts with adenosine through hydrogen bonds to nitrogen residues 3 and 7 of the purine ring, and the 5' and 3' hydroxyl groups of ribose. P2 binds the amidine group comprising N1 and the position 6 amine. Additional interactions are formed by pi-stacking of the purine ring with an aromatic residue in the transporter-binding pocket, and N9 also contributes to binding, although not through a hydrogen bond [17]. Abbreviations: HAPT1, high-affinity PMD transporter 1; LAPT1, low-affinity PMD transporter 1; PMD, pentamidine; R, hydrogen (adenine) or ribose (adenosine).

as the PMD-sensitive adenosine transport activity. The construction of null mutants of *TbAT1* (the gene believed to encode P2 [15]) and our transport studies with [³H] PMD and [³H] adenosine in these cells are consistent with this gene encoding both P2 and ASPT1 activities (E. Matovu *et al.*, unpublished).

The P2 transporter consistently mediated 50–70% of the PMD uptake [18,19]. Two additional PMD transport activities could be distinguished: a high-affinity pentamidine transporter (HAPT1; $K_m = 36 \pm 6$ nM) and a low-affinity pentamidine transporter (LAPT1; $K_m = 56 \pm 8$ μ M). HAPT1 was competitively inhibited by propamidine, but neither transporter was sensitive to inhibition by other trypanocides, including stilbamidine and melarsen oxide, or by various amino acids, vitamins, purines and pyrimidines.

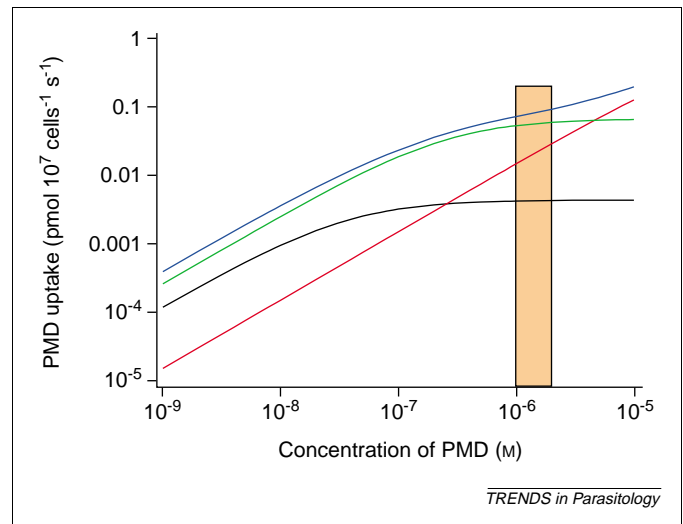


Fig. 3. PMD uptake rates by various transporters in *Trypanosoma brucei brucei* bloodstream forms. Uptake rates were plotted as a function of the extracellular PMD concentration from $V = (V_{max} \times L)/(L + K_m)$ in which L is the ligand concentrations, and the K_m and V_{max} values for individual transporters were used from Table 1. Plasma concentrations in patients treated with the standard 4 mg kg⁻¹ dosing have been determined at 0.5–2.5 μ M [20]. Over this range, the contribution of LAPT1 increased from 13% to 35% of total PMD uptake, due to its very high capacity for PMD transport. Key: black line, HAPT1; blue line, combined uptake by HAPT1, LAPT1 and P2; green line, P2; orange bar, maximum plasma concentration of PMD in patients; red line, LAPT1. Abbreviations: HAPT1, high-affinity PMD transporter 1; LAPT1, low-affinity PMD transporter 1; PMD, pentamidine.

Because of the different kinetic parameters of the three PMD transporters, the relative contribution of each to the total flux is a function of the extracellular PMD concentration (Fig. 3), which reaches up to 2.5 μ M in plasma of sleeping sickness patients treated with the standard dose of 4 mg kg⁻¹ [20]. The combination of high affinity and relatively high V_{max} of ASPT1 ensures that, at any concentration, at least 50% of the flux is mediated by this carrier. HAPT1, as a result of its very high affinity, contributes relatively little because it is saturated at low concentrations.

It is not known whether the P2 or ASPT1 transporter is energy-dependent, but the related P1 nucleoside transporter of *T. b. brucei* procyclics is a proton symporter [21], as are its purine and pyrimidine nucleobase transporters [22].

Table 1. Kinetic parameters and inhibition constants of the trypanosomal pentamidine transporters^a.

	<i>Trypanosoma brucei brucei</i> ^b			<i>Leishmania mexicana</i> ^c
	P2	HAPT1	LAPT1	PT1
Kinetic parameters				
K_m (μ M)	0.26 \pm 0.03	0.036 \pm 0.006	56 \pm 8	7.4 \pm 1.8
V_{max}	0.068 \pm 0.007 ^d	0.0044 \pm 0.0004 ^c	0.85 \pm 0.15 ^d	452 \pm 42 ^e
K_i values of inhibitors (μM)				
Stilbamidine	2.4 \pm 0.3	56 \pm 3	> 250	21 \pm 5
Propamidine	1.9 \pm 0.8	4.6 \pm 0.7	> 250	12 \pm 1
Diminazene	2.5 \pm 0.8	63 \pm 3	> 250	27 \pm 3
Melarsen oxide	0.63 \pm 0.20	> 100	NE, 100	ND

^aAbbreviations: HAPT1: high-affinity pentamidine transporter 1; LAPT1: low-affinity pentamidine transporter 1; ND, not determined; NE, no effect at indicated concentration; P2, purine transporter 2; PT1, pentamidine transporter 1.

^bFrom Ref. [18]

^cFrom Ref. [39]

^dIn pmol (10^7 cells)⁻¹

^eIn fmol s⁻¹ μ g protein⁻¹

to high levels of PMD *in vitro* [27]. There was no crossresistance to other diamidines and the resistance mechanism is still unknown. However, it is unlikely that this resistance will arise or persist 'in the field' because the resistant clone was much less virulent than the parent strain [27].

It is recommended that diamidine analogues considered for future use against human African trypanosomiasis (HAT) are tested for uptake by all three PMD transporters to delay the onset of resistance. The recent suggestion that diminazene aceturate could be used against HAT [28] does not seem sensible in the light of this information because it would quickly induce high levels of resistance, as well as a low-level resistance to melarsoprol. Because *T. b. gambiense* isolates from melarsoprol-treated relapse patients display only a low level of melarsoprol resistance [29], this could rapidly escalate the current crises of melarsoprol refractoriness [30].

PMD uptake and resistance in *Leishmania mexicana*

Pentavalent antimonial drugs are the standard first-line treatment for leishmaniasis [31], although resistance is a growing problem [32]. PMD is used clinically as an alternative to pentavalent antimony in leishmaniasis treatment [33–35]. A definitive mode of action has yet to be elucidated for the drug, although studies on PMD uptake and distribution in drug-sensitive and -resistant parasites have recently given some insight into events.

PMD was found to be a competitive inhibitor of arginine transport [36] in *Leishmania donovani* and a noncompetitive inhibitor of putrescine and spermidine transport in *Leishmania infantum* [37], *L. donovani* and *L. mexicana* [38]. The availability of radiolabelled PMD helped to determine whether the drug shared a transporter with arginine, putrescine or spermidine [39]. These metabolites did not inhibit [³H] PMD uptake; therefore, the observed inhibition by PMD must relate to the drug's ability to inhibit permeases through some mechanism other than it being a competitive substrate. The drug enters both promastigote and amastigote forms of *L. mexicana* via a carrier-mediated process that recognizes the drug with measured K_m values of 7.4 μM in promastigotes and 13.9 μM in amastigotes [39]. The maximum velocity of uptake is substantially higher in amastigotes than in promastigotes (8540 $\text{fmol s}^{-1} \mu\text{g protein}^{-1}$ in amastigotes and 452 $\text{fmol s}^{-1} \mu\text{g protein}^{-1}$ for promastigotes) [39]. A wide variety of common metabolites also failed to inhibit PMD uptake. Three inhibitors of P-glycoproteins (trifluoperazine, prochlorperazine and verapamil) were competitive inhibitors of uptake, but the physiological role of the *Leishmania* transporter that carries PMD into the cell is unknown. Some divalent inorganic cations also showed inhibitory activity and it is tempting to speculate that a divalent cation channel or transporter could operate as the transporter for PMD, although with a mechanism markedly different from that normally used for the natural substrates.

Resistance to PMD has been observed in *Leishmania* [40–46]. The mechanism(s) of resistance are poorly understood, but events at the mitochondrion appear to be associated with the drug's action. Electron microscopy

revealed that treatment of *Leishmania* with PMD leads to disintegration of the network of intercatenated circular DNA molecules that comprise the mitochondrial genome, termed the kinetoplast, and also the mitochondrion [47,48], and a collapse in the mitochondrial membrane potential (MMP) [49] is one of the first manifestations of this drug treatment. In yeast, a mitochondrial localization was identified for the product of the *PNT1* gene that confers resistance to PMD when overexpressed [50,51]. In mammalian cells, PMD affects MMP as well [52].

The affinity for PMD shown by its transporter in *Leishmania* is unaltered in resistant parasites (7.4 \pm 1.8 μM and 5.9 \pm 2.1 μM , in wild-type and resistant strains, respectively). However, PMD accumulates to a lower level, and the apparent V_{max} for uptake is diminished in resistant parasites. A similar situation has been noted in resistance to isometamidium in *Trypanosoma congolense* [53]. Resistance in this case was concluded not to associate with alterations at the level of a plasma membrane transporter, but with changes in the MMP. Isometamidium rapidly enters the *T. congolense* mitochondrion where it binds tightly to kDNA. The apparent decrease in V_{max} in resistant cells correlated with diminished capacity to accumulate drug intracellularly. Similar to isometamidium, diamidines are organic cations, which also bind to the kinetoplast. 4,6-diamidino-2-phenylindole (DAPI), a fluorescent analogue of PMD (Fig. 1), enters cells and accumulates rapidly in the mitochondrion, binding to the kinetoplast in wild-type, but not in high-level PMD-resistant, *Leishmania* [39]. The fluorescent diamidine binds equally well to wild-type and resistant kinetoplasts, so access to the kinetoplast, rather than its ability to bind to the DNA, differentiates wild-type and resistant cells. A diminution of MMP in PMD-resistant *Leishmania* [39], as measured by accumulation of rhodamine-123, indicates that MMP is diminished in the drug-resistant line. Because rhodamine-123 is a substrate for P-glycoproteins, a diminished accumulation of this substance could also indicate enhanced P-glycoprotein efflux activity at the leishmanial plasma membrane. However, while an efflux mechanism for PMD was identified in *Leishmania* [39], the activity was the same in wild-type and resistant parasites [39].

The data on a resistance mechanism are consistent with a model, outlined in Fig. 4b, whereby PMD enters both resistant and wild-type cells via the same plasma membrane carrier that is unaltered in the resistant lines [39]. However, in wild-type but not in resistant parasites, PMD is rapidly accumulated into the mitochondrion with uptake being driven by the greater MMP of wild-type cells. Inhibitors of the MMP (e.g. potassium cyanide, sodium azide, 2,4-dinitrophenol and oligomycin) decrease PMD uptake into wild-type cells. By contrast, in resistant parasites, the drug does not accumulate rapidly into the mitochondrion, but remains free in the cytosol. Efflux pumps appear to be operative in removing cytosolic or membrane-associated PMD from the cell (as judged by the ability of inhibitors of pumps shown to be operative in *Leishmania*, e.g. verapamil, trifluoperazine and prochlorperazine, to allow accumulation of drug in resistant cells). It is likely that this PMD efflux system can remove the

majority of PMD from the non-mitochondrial compartment of both cell types but, in the wild-type cells, most of the PMD becomes bound to the kinetoplast within the mitochondrion, and so is not available for efflux. This is not the case in resistant parasites. A balance involving rate of uptake into the cell, efflux from the cell at the plasma membrane, and accumulation into the mitochondrion would then underlie the events reported here. Agents that inhibit efflux might lead to reversal of resistance by allowing PMD to accumulate in the cytosol to levels that drive accumulation into the mitochondrion, through an unknown transporter or channel.

PMD against *Plasmodium falciparum*

Diamidines were tested against experimental malaria almost immediately after their synthesis [54]. Monkeys infected with *Plasmodium knowlesi* at parasitaemias that would, without intervention, be fatal could be saved with a daily dose of 2.5 mg kg⁻¹ of undecane diamidine [54] and, in 1940, PMD and stilbamidine were shown to have similar curative effects in monkeys [55]. Sen Gupta also observed that, even in highly endemic areas, patients treated for kala azar with stilbamidine were rarely suffering from malaria [56]. However, other drugs, such as quinine and atebirin were effective, and diamidines were not pursued as much as antimalarials until recently. In 1990, Tidwell and co-workers synthesized new classes of diamidines, and demonstrated that these have considerable antimalarial activity [57]. Our recent studies confirm the potential of diamidines against *P. falciparum*, which showed a lack of crossresistance with standard antimalarials [58].

PMD is selectively accumulated into the *Plasmodium*-infected erythrocyte, reaching concentrations 500-fold greater than the surrounding medium. By contrast, no significant accumulation into uninfected erythrocytes is observed. Hence, it is probable that this specific uptake process underlies the antimalarial activity of PMD and other diamidines [58].

Kinetic studies reveal an initial rapid phase of PMD uptake (half life = 90 s) followed by a slower phase that is linear over several hours. The initial phase is nonsaturable in the micromolar range, but the slower phase shows two saturable processes with dissociation constants (K_d) of $\sim 2 \mu\text{M}$ and $200 \mu\text{M}$.

At first, we focused on the rapid initial phase of uptake because this most likely represents the passage of PMD across the host erythrocyte membrane. As well as being nonsaturable, the initial uptake phase is insensitive to known inhibitors of PMD transport into other protozoan parasites. Instead, initial uptake is effectively blocked by inhibitors of the new permeability pathway (NPP) [58]. The NPP is induced by the malaria parasite on the host cell membrane and has properties of an inwardly rectified anion channel rather than a transporter [59]. It is nonsaturable and anion-selective, but it exhibits significant permeability to cations [60,61]. Changing the counter ion (major permeant ion) in the bathing medium markedly influences passage of cations through the NPP [60]. We find similar effects of changing the counter ion on the initial uptake of PMD [58].

Taken together, all of these data suggest that PMD is

not transported as in other protozoan parasite species, but penetrates the *Plasmodium*-infected erythrocyte membrane via the induced NPP channel (Fig. 4c). Once through the host cell membrane, PMD binds to haemoglobin but with low affinity. Here, the drug will have direct access to the external membranes of the intracellular parasite. From here, the drug must be able to access the interior of the intracellular parasite to exert its antimalarial effect, but the means by which this is achieved is not known. It is possible that the drug is taken up by the intracellular parasite in the cytostomal feeding process (f, [61]). Alternatively, a putative transporter could carry the drug across the parasite plasma membrane (Fig. 4c). Regardless of the way PMD is taken up by the intracellular parasite, it is likely that a significant amount of the drug eventually ends up in the food vacuole: PMD binds avidly to ferriprotoporphyrin IX (FPIX) *in vitro* with an affinity of $\sim 2 \mu\text{M}$. It also binds to aggregated FPIX with lower affinity. FPIX is generated inside the food vacuole from the digestion of haemoglobin and it is likely that PMD binds to FPIX in this location. During the slower phase of PMD accumulation, a saturable process with an affinity of $2 \mu\text{M}$ is revealed by Scatchard analysis [58]. We believe that this saturable process reflects the binding of PMD to FPIX in the food vacuole. This argument is based on the K_d being identical to that obtained for PMD–FPIX binding *in vitro* and is supported by the data obtained using plasmepsin inhibitors [58]. Plasmepsins are the food vacuole enzymes thought to be responsible for the first steps in the degradation of haemoglobin [62]. Blocking these enzymes stops haemoglobin digestion in the parasite and effectively stops the generation of FPIX [61]. We were able to demonstrate that this action effectively blocks the high-affinity binding of PMD to infected cells and antagonizes the antimalarial activity of all of the diamidines [58].

All of the diamidines tested so far are efficient blockers of haemozoin crystallization *in vitro*. As such, their ultimate mode of action is probably similar to that of chloroquine (CQ). Interestingly, however, there is no crossresistance with CQ or any other standard antimalarials [58], indicating that the diamidine motif could be an interesting lead structure for the development of new antimalarials.

Concluding remarks

Diamidines have a long and distinguished history in the fight against infectious disease and in particular protozoan infections. While much remains unclear about their precise mechanism of action in kinetoplastids, longstanding issues such as selective uptake and mechanisms of (cross)resistance have now largely been resolved. New classes of orally available diamidines with fewer side-effects or improved central nervous system penetration are being developed, and we are now in a better position to predict or prevent the emergence of resistance to the new drugs. The fact that this class of drugs has action against a variety of organisms including opportunistic fungal infections, such as *Pneumocystis carinii* and *Aspergillus*, will make development more commercially attractive. Although prophylaxis on the massive scale of the colonial era is now impracticable, it could be considered to bring epidemics in

defined foci under control, especially in areas with high incidence of melarsoprol treatment failure.

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