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# The Acidocalcisome as a Target for Chemotherapeutic Agents in Protozoan Parasites

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### Abstract

Acidocalcisomes are acidic organelles rich in calcium and phosphorus that have been conserved from bacteria to man. In parasitic protozoa acidocalcisomes possess enzymes that are absent or different from their mammalian counterparts and could be potential targets for chemotherapy, such as the vacuolar proton translocating pyrophosphatase, and the soluble inorganic pyrophosphatase, both of which are inhibited by pyrophosphate analogs (bisphosphonates). In addition, a number of drugs, including bisphosphonates, and diamidines appear to accumulate in these organelles and/or induce an increase in their numbers, potentially enhancing their toxicity. Bisphosphonates mechanism of action, however, is by inhibition of the isoprenoid pathway and more specifically the prenyl diphosphate synthases.

## INTRODUCTION

The acidocalcisome is a dense, acidic organelle (Fig. 1A) with a high concentration of phosphorus present as pyrophosphate and polyphosphate (poly P) complexed with calcium, and other cations. The acidocal cisome membrane contains a number of pumps ( $Ca^{2+}$ -ATPase. V-H<sup>+</sup>-ATPase, H<sup>+</sup>-PPase), exchangers (Na<sup>+</sup>/H<sup>+</sup>, Ca<sup>2+</sup>/H<sup>+</sup>), and channels (aquaporins), while its matrix contains enzymes related to pyrophosphate and poly P metabolism [1] (Fig. 1B). Acidocalcisomes have been found in several pathogenic microorganisms [2] as well as in the green alga Chlamydomonas reinhardtii [3], and the slime mold Dictyostelium discoideum [4]. The identification of acidocal cisomes in bacteria [5,6] and the finding that human platelet dense granules are similar to acidocalcisomes [7,8], indicated that these are organelles have been conserved from bacteria to humans. Some of the potential functions of the acidocalcisome are the storage of cations and phosphorus, and its participation in pyrophosphate and polyphosphate metabolism, calcium homeostasis, maintenance of intracellular pH homeostasis, and osmoregulation [1]. The discovery of novel enzymes in this organelle that are absent from mammalian cells led to the finding of compounds (bisphosphonates) that produced radical cures in animal models of diseases caused by several parasites [9]. Further exploration of the structure and function of acidocalcisomes in protozoan parasite could lead to the identification of new targets for drug action.

# POTENTIAL TARGET ENZYMES LOCATED IN THE ACIDOCALCISOME

Of the enzymes present in the acidocalcisomes, two have been found to be targets for drugs with *in vitro* and/or *in vivo* activity against different protozoan parasites: a vacuolar proton

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translocating pyrophosphatase (V-H<sup>+</sup>-PPase) and a soluble inorganic pyrophosphatase (PPase).

The V-H<sup>+</sup>-PPase activity has been detected in the following parasitic protozoa: *Trypanosoma cruzi* [10], *T. brucei* [11,12], *Leishmania donovani* [13], *L. amazonensis* [14], *Phytomonas françai* [15], *Leptomonas wallacei* [16], *Herpetomonas spp.* [16,17], *Plasmodium spp.* [18, 19], *and Toxoplasma gondii* [20,21]. This enzyme also localizes to acidocalcisomes in all these species (Fig. 1B). The V-H<sup>+</sup>-PPase from *T. cruzi* functions in yeast [22]. The acidocalcisomal V-H<sup>+</sup>-PPase is K<sup>+</sup>-stimulated (type I), and can be used as a marker for acidocalcisome purification [3,4,10–13,23]. Although it is not restricted to the acidocalcisome it is concentrated in this organelle. The *T. cruzi* V-H<sup>+</sup>-PPase is also found in the Golgi complex and in the plasma membrane [24]. The *Plasmodium* spp. V-H<sup>+</sup>-PPase is also localized in the digestive vacuole [19,25,26].

In earlier work, it was found that some pyrophosphate analogs, bisphosphonates (containing a non-hydrolyzable P-C-P, rather that a P-O-P, backbone) as well as imidodiphosphate (containing a non-hydrolyzable P-N-P group), were inhibitors of a plant (mung bean, *Vigna radiata* L.) V-H<sup>+</sup>-PPase [27]. A more extensive investigation of the structural aspects of the effectiveness of bisphosphonates as competitive inhibitors of this enzyme was reported later [28]. More recently the results of a three-dimensional quantitative structure-activity relationship (3D-QSAR) comparative molecular field analysis (ConMFA) of the activity of 18 bisphosphonates and imidodiphosphate in the inhibition of a mung beam (*Vigna radiata* L.) V-H<sup>+</sup>-PPase was reported [29], and it was shown that the activities of the V-H<sup>+</sup>-PPase inhibitors could be predicted to within about a factor of two. Several of the compounds investigated were active against the parasite enzymes [10,11,13,20,30]. One of the best known inhibitors of the V-H<sup>+</sup>-PPase, aminomethylenediphosphonate (AMDP) [31], was able to impair intracellular replication of *T. gondii* in tissue culture cells exerting little or no effect on host cell invasion [20,30]. Some of the treated parasites had ultrastructural alterations compatible with acidocalcisome disruption [30].

The vacuolar soluble pyrophosphatase (VSP1) present in *Trypanosoma brucei*, is essential for growth of bloodstream forms in their mammalian host and is located in acidocalcisomes [32] (Fig. 1B). Depending on the nature of its divalent metal ion cofactor, this soluble enzyme can act either as a pyrophosphatase (PPase; with +Mg<sup>2+</sup>) or as a short-chain polyphosphatase (PPX; with +Zn<sup>2+</sup>). It was found that the exopolyphosphatase (tripolyphosphatase) activity (in the presence of Zn<sup>2+</sup>, which is abundant in acidocalcisomes) of TbVSP1 was inhibited by bisphosphonates [33]. The inhibition of the recombinant TbVSP1 by a panel of 81 bisphosphonates was reported [33]. The IC<sub>50</sub> values for enzyme inhibition were found to vary from 2 to 850  $\mu$ M. In general the most active compounds contained both a single aromatic ring and a hydrogen bond donor feature. Thirteen of the most potent compounds were tested *in vivo* in a mouse model of *T. brucei* infection. The most active compound *in vivo* provided a 40% protection from death with no apparent side effects, suggesting that further development of such compounds might be of interest [33].

# ROLE OF THE ACIDOCALCISOME IN THE MECHANISM OF ACTION OF BISPHOSPHONATES

Bisphosphonates are used to treat a variety of bone resorption diseases including osteoporosis, Paget's disease, hypercalcemia caused by malignancy, and tumor metastases in bone [34]. Bisphosphonates have also been shown to have activity against different protozoan parasites *in vitro* and *in vivo* [9].

Selective action on bone is based on the binding of the bisphosphonate moiety to the bone mineral [34]. It has been postulated that the acidocalcisomes are equivalent in composition to the bone mineral and that accumulation of bisphosphonates in these organelles, as they do in bone mineral, facilitates their antiparasitic action [35].

The primary target for the nitrogen-containing or amino-bisphosphonates is though to be the isoprenoid pathway at the level of the enzyme farnesyl diphosphate synthase (FPPS) [36–41] (Fig. 2). By inhibiting this enzyme, bisphosphonates inhibit the formation of farnesyl diphosphate (FPP), a compound used in protein prenylation of proteins like Ras, Rho and Rap [42–44], and in the production of dolichols, ubiquinones, heme a, and sterols. FPPS inhibition results, in addition to decreased prenylation of proteins and generation of FPP derivatives, in the accumulation of isopentenyl diphosphate (IPP), a known  $\gamma\delta$  Tcell activator [45]. These alterations lead to apoptotic cell death [46–48]. Recent work has shown that bisphosphonates can also target other enzymes of the isoprenoid pathway, like for example geranylgeranyl diphosphate synthase (GGPPS) [49] (Fig. 2) and that they can target multiple sites in prenyltransferases [50]. Interestingly, some bisphosphonates are also able to inhibit the activity of *T. cruzi* hexokinase, an enzyme that in contrast to the mammalian enzyme, is inhibited by PPi [51,52]

Nitrogen-containing bisphosphonates (Fig. 3) were first found to be effective in the inhibition of *T. cruzi in vitro* and *in vivo* without toxicity to the host cell [53]. Later, a series of bisphosphonates was tested on the growth of *T. gondii*, *T. b. rhodesiense*, *L. donovani* and *P. falciparum in vitro* showing that bisphosphonates could effectively inhibit the growth of these parasites [35]. The bisphosphonate risedronate (Fig. 3) was shown to inhibit *Cryptosporidium parvum* growth *in vivo* using a xenograft model [54].

The use of another bisphosphonate, pamidronate (Fig. 3), resulted in the radical cure of experimental cutaneous leishmaniasis in mice [55]. Pamidronate was also active *in vivo* against *L. donovani* by intravenous administration [56]. Risedronate (Fig. 3) had a 50% effective dosage of five 2.6 mg/kg of body weight intraperitoneal doses against *L. donovani*-infected mice.

In vivo testing against *T. gondii* in mice showed that risedronate can significantly increase the survival of mice infected by this parasite [57]. *In vitro* testing of risedronate in *T. cruzi* showed that it had selective antiproliferative effects against the intracellular amastigotes, and at 100  $\mu$ M, was able to prevent completely the development of *T. cruzi* infection of murine muscle heart or Vero cells, and to cure cultures which were already infected [58]. *In vivo* testing of bisphosphonates against *T. cruzi* has shown that risedronate can significantly increase the survival of mice infected by *T. cruzi* [57,59].

The effect of a series of 102 bisphosphonates on the inhibition of growth of *Entamoeba histolytica* and *Plasmodium falciparum in vitro* was also determined [60]. The most active compounds (IC<sub>50</sub> 4–9  $\mu$ M) against *E. histolytica* were nitrogen-containing bisphosphonates with relatively large aromatic side chains. Five bisphosphonates were selected and screened for their ability to delay the development of amebic liver abscess formation in an *E. histolytica* infected hamster model and 2 compounds were found to decrease liver abscess formation at 10 mg/kg ip with little or no effect on normal liver mass [60]. With *P. falciparum*, the most active compounds were n-alkyl bisphosphonates (Fig. 4). Five compounds were selected for *in vivo* investigation in a *Plasmodium berghei* ANKA Balb/c mouse suppressive test. The most active compound caused an 80% reduction in parasitemia with no overt toxicity [60].

The activity of 60 bisphosphonates against the replication of *T. gondii in vitro* and of three of the most active compounds, *in vivo* has been investigated [61]. The two most active compounds

found were n-alkyl bisphosphonates containing long (n = 9 or 10) hydrocarbon chains (Fig. 4), not the nitrogen-containing species used in bone resorption therapy. The three most active compounds found *in vitro* were tested *in vivo* in a Smith-Webster mouse model and the two most active bisphosphonates were found to provide up to an 80% protection from death, a considerable improvement over that found previously with nitrogen-containing bisphosphonates [61]. This effect may originate in the much higher therapeutic indices of these alkyl bisphosphonates, as deduced from in vitro assays using LD<sub>50</sub> values for growth inhibition of a human cell line.

Alkyl bisphosphonates (Fig. 4) were also shown to be potent inhibitors of *T. cruzi* amastigotes growth *in vitro* [62]. Overall, these results indicate that alkyl bisphosphonates are promising compounds for further development as agents against parasite growth, *in vivo* [61,62], especially against Apicomplexan parasites [60,61].

There is strong evidence that the main target of the most active bisphosphonates in protozoan parasites is the isoprenoid biosynthesis pathway enzyme farnesyl diphosphate synthase (FPPS): (1) There is excellent correlation between inhibition of the enzyme in *T. cruzi* [63], *T. brucei* [65], *T. gondii* [65], and *L. major* [66] and growth of these parasites *in vitro*; (2) *In vitro* "rescue" experiments showed reversal of risedronate-induced growth inhibition of *T. b. rhodesiense* by GGPP, FPP, or farnesol [67]; (3) Molecular modeling and structure-activity investigations of enzyme and *in vitro* growth inhibition data in *T. brucei* resulted in similar pharmacophores [64]; (4) a *T. gondii* strain engineered to overexpress FPPS required considerably higher levels of bisphosphonates to achieve 50% growth inhibition, while the IC<sub>50</sub> for atovaquone (which does not inhibit FPPS) remained the same in the overexpressing strain [65]; (5) Promastigotes of *L. major* overexpressing FPPS were highly resistant to risedronate and the degree of resistance correlated with the increase in enzyme activity [68]; (7) RNAi experiments in *T. brucei* has shown that FPPS is an essential enzyme thus validating it as a target for chemotherapeutic agents [65].

The farnesyl diphosphate synthase genes from *T. cruzi* [63], *T. brucei* [64], *L. major* [68], and *T. gondii* [65] have been cloned and their protein products purified and characterized biochemically. The tridimensional structures of *T. brucei* [69,70], *T. cruzi* [71], *P. berghei*, and *C. parvum* [72] FPPSs have been solved, providing mechanistic insights that will have important implications for future drug design.

The reasons why alkyl-bisphosphonates have higher activity than nitrogen-containing bisphosphonates in Apicomplexan parasites is now becoming clear. In the Apicomplexans, the putative "FPPS" enzymes actually produce not only FPP, but much longer ( $C_{20}$ ,  $C_{25}$  and up) prenyl diphosphates [65]. Long chain bisphosphonates are able to block the TgFPPS active site (since it is bifunctional). Moreover, the availability of the closely related *P. berghei* and *C. parvum* X-ray structures [72] strongly suggests a structural explanation in that the Apicomplexans have a F to C, S substitution in the fifth aminoacid upstream of the first aspartate rich domain (FARM) region, enabling longer chain inhibitors to bind in the active site [65]. However, these inhibitors are expected to have a steric clash with the FF groups in the host cell FPPS, resulting in no FPPS inhibition [70]. Interestingly, it has been shown that inhibition of TgFPPS, which is a bifunctional enzyme generating longer chain isoprenoids (GGPP) [65], correlates better with inhibition of solanesyl diphosphate synthase from *T. cruzi* (TcSPPS), which is an enzyme that generates the 45-carbon solanesyl diphosphate (SPP, [73a]) than with inhibition of other FPPSs, that generate only FPP [73b).

In summary, there are several reasons for bisphosphonates to be good candidate drugs for treatment of parasitic disease. First, they have already been developed to treat other diseases and consequently have low toxicity; second, their structures are simple, so they are easy to

synthesize; third, experimental results have shown that several bisphosphonates have excellent inhibitory activity against different parasites *in vitro* and *in vivo*.

### OTHER DRUGS TARGETING THE ACIDOCALCISOME

Acidocalcisomes are also known by the names 'volutin granules' or 'polyphosphate granules' [1], and early work by Ormerod [74] proposed that they become more visible under light microscopy when cells are treated with drugs. For this reason they were also named as 'chemotherapy granules' [75]. Hawkins and Smiles in 1941 [76] were able to show accumulation of the fluorescent drug stilbamidine in trypanosome granules. Other drugs, like quinapyramine, suramin, hydroxystilbaminine [77], and acriflavine [76] were also found to concentrate in these granules. Interestingly, some of these drugs are first concentrated in the kinetoplast and nucleus, then diffuse to the cytosol, and finally concentrate in granules [76, 77]. Recent work on other diamidines such as DB75 (furamidine) and DB820, which are in phase III clinical trials against human African trypanosomiasis, revealed a similar pattern of accumulation, first in DNA-containing regions such as the nucleus and kinetoplast and later in acidocalcisomes [78]. However, the impact that acidocalcisome accumulation has on the mechanism of action of these compounds in not known [79].

Ketoconazole and terbinafine, two sterol biosynthesis inhibitors, were shown to induce the formation of numerous and diverse acidocalcisomes in promastigotes and amastigotes of *L. amazonensis*, which were enclosed by in larger compartments with access to endocytic tracers [80]. Naphthoimidazole compounds were found to decrease the electron density of acidocalcisomes of *T. cruzi* [81].

Azithromicin, a drug used against toxoplasmosis has also been shown to accumulate in acidic compartments within *T. gondii* tachyzoites [82]. Other chemotherapeutic agents used against malaria (e.g. chloroquine) have also been shown to accumulate in acidic compartments [83] and Na<sup>+</sup>/H<sup>+</sup> exchangers such as monensin are used in the treatment of coccidiosis. Chloroquine accumulates in the acidocalcisomes of *T. brucei*, slows down growth in vivo and prolongs the survival time of infected mice [83].

### CONCLUSIONS

In conclusion, acidocalcisomes are potential targets for the chemotherapy of protozoan parasitic diseases not only because they possess enzymes that are absent or different from their mammalian counterparts, but also because of their acidic characteristics, which allow them to accumulate basic drugs, potentially enhancing their toxicity.

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Figure 1B

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### Figure 1. Ultrastructure and composition of acidocalcisomes

**A.** Transmission electron microscopy of a procyclic stage of *Trypanosoma brucei* showing the acidocalcisomes (dark granules). Bar = 3  $\mu$ m. Reprinted with permission from ref. [87]. **B**. Schematic representation of an acidocalcisome. A H<sup>+</sup> gradient is established by a vacuolar ATPase (V-H<sup>+</sup>-ATPase) and a vacuolar pyrophosphatase (V-H<sup>+</sup>-PPase). Ca<sup>2+</sup> transport is driven by a Ca<sup>2+</sup>-ATPase. Other transporters include Na<sup>+</sup>/H<sup>+</sup>, and Ca<sup>2+</sup>/H<sup>+</sup> exchangers, a Cl<sup>-</sup> channel, and a water channel or aquaporin. Transporters for basic amino acids, P<sub>i</sub>, PP<sub>i</sub>, and cations are potentially present. The matrix is rich in PPi and polyphosphatase (PPX), and pyrophosphatase (PPase). Not all the enzymes and transporters are present in all acidocalcisomes.



### Figure 2. Overview of the pathway for isoprenoid synthesis

The DOXP/MEP pathway, present in higher plants, green algae, some bacteria, *Plasmodium* spp. and yeast as well as the Mevalonate pathway, present in mammals, higher plants, some bacteria, trypanosomatids, and yeast, generate isopentenyl diphosphate (IPP), which isomerizes to dimethylallyl diphosphate (DMAPP). The farnesyl diphosphate synthase (FPPS) catalizes the reaction of DMAPP with IPP to generate geranyl diphosphate (GPP), which incorporates another IPP to generate farnesyl diphosphate (FPP). FPP is the precursor for ubiquinones, heme a, sterols, dolichols and geranylgeranyl diphosphate (GGPP) through the action of GGPP synthase. Bisphosphonates (BP) inhibit the short chain prenyl transferases (FPPS and GGPPS).



Figure 3. Structure of GPP, FPP, and different bisphosphonates commercially available for the treatment of bone resorption diseases



**Figure 4. Structure of n-alkyl bisphosphonates effective against Apicomplexan parasites** The figure shows the structure of compounds with 9 and 10-carbon chain.