Substrate Inhibition of Uracil Phosphoribosyltransferase by Uracil Can Account for the Uracil Growth Sensitivity of Leishmania donovani Pyrimidine Auxotrophs

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Background: Leishmania donovani salvage all pyrimidines through uracil phosphoribosyltransferase (LdUPRT).

Results: LdUPRT phosphoribosylates uracil, 5-fluorouracil, and 4-thiouracil and is susceptible to substrate inhibition.

Conclusion: LdUPRT recognizes pyrimidine analogs, and substrate inhibition by LdUPRT explains the supersensitivity of pyrimidine auxotrophs to uracil.

Significance: Substrate inhibition of LdUPRT provides a mechanism for uracil susceptibility and offers a protective function for the parasite.

The pathogenic protozoan parasite Leishmania donovani is capable of both de novo pyrimidine biosynthesis and salvage of pyrimidines from the host milieu. Genetic analysis has authenticated Leishmania donovani uracil phosphoribosyltransferase (LdUPRT), an enzyme not found in mammalian cells, as the focal enzyme of pyrimidine salvage because all exogenous pyrimidines that can satisfy the requirement of the parasite for pyrimidine nucleotides are funneled to uracil and then phosphoribosylated to UMP in the parasite by LdUPRT. To characterize this unique parasite enzyme, LdUPRT was expressed in Escherichia coli, and the recombinant enzyme was purified to homogeneity. Kinetic analysis revealed apparent $K_m$ values of 20 and 99 $\mu$M for the natural substrates uracil and phosphoribosylpyrophosphate, respectively, as well as apparent $K_m$ values 6 and 7 $\mu$M for the pyrimidine analogs 5-fluorouracil and 4-thiouracil, respectively. Size exclusion chromatography revealed the native LdUPRT to be tetrameric and retained partial structure and activity in high concentrations of urea. Leishmania donovani mutants deficient in de novo pyrimidine biosynthesis, which require functional LdUPRT for growth, are hypersensitive to high concentrations of uracil, 5-fluorouracil, and 4-thiouracil in the growth medium. This hypersensitivity can be explained by the observation that LdUPRT is substrate-inhibited by uracil and 4-thiouracil, but 5-fluorouracil toxicity transpires via an alternative mechanism. This substrate inhibition of LdUPRT provides a protective mechanism for the parasite by facilitating purine and pyrimidine nucleotide pool balance and by sparing phosphoribosylpyrophosphate for consumption by the nutritionally indispensable purine salvage process.

Leishmania donovani is a protozoan parasite and etiologic agent of visceral leishmaniasis, a disease that is ultimately fatal if untreated. Leishmania are digenetic parasites subsisting as the motile, extracellular promastigote in the female Phlebotomine sandfly vector and as the nonmotile, intracellular amastigote within the phagolysosome of macrophages inside the infected mammalian host. There is no vaccine against leishmaniasis, and the current assortment of drugs used to treat leishmaniasis is far from ideal. These drugs are toxic to the host, require invasive means of administration, and trigger resistance in the field. Thus, the need to discover new drugs and validate new drug targets for the treatment of leishmaniasis—or for that matter any disease of parasitic origin—is imperative.

Among the pathways that have been touted as potential antiparasitic targets are those for purines and pyrimidines, the building blocks for nucleic acid synthesis. Leishmania, like all protozoan parasites studied to date, are incapable of synthesizing purine nucleotides de novo, and therefore, each genus must obligatorily scavenge purines from its hosts (1). In contrast, most, but not all, protozoan parasites, including Leishmania, are prototrophic for pyrimidines (1). The de novo pathway for pyrimidine biosynthesis is conserved among eukaryotes and prokaryotes and consists of six enzymes that generate UMP from CO$_2$, amino acids, and 5-phosphoribosyl-1-pyrophosphate (PRPP)$^2$ (Fig. 1). UMP is then distributed into ribonucleotides via nucleotide kinases and into deoxyribonucleotides by thymidylate synthase. Gene sequencing, supported by biochemical studies, has revealed a number of significant differences between the pyrimidine biosynthetic pathways of Leishmania and mammals: 1) the genes encoding the pyrimidine pathway of Leishmania are syntenic (2–5), whereas the mammalian pyrimidine genes are not; 2) the genes encoding the first three enzymes in Leishmania are dis-

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$^2$ The abbreviations used are: PRPP, 5-phosphoribosyl-1-pyrophosphate; UMPs, UMP synthase; CPS, carbamoyl phosphate synthetase; LdUPRT, L. donovani uracil phosphoribosyltransferase; IPTG, isopropyl $\beta$-D-1-thiogalactopyranoside; TgUPRT, T. gondii uracil phosphoribosyltransferase; PR, uracil phosphoribosyltransferase; Ni-NTA, nickel-nitrilotriacetic acid; CHES, 2-cyclohexylaminoethanesulfonic acid; Tricine, N-[2-hydroxy-1,1-bis(2-hydroxyethyl)glycine. 

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create, unlike the mammalian pathway in which there is a single gene encoding a trifunctional protein (6, 7); 3) the last two enzymes of the pyrimidine biosynthetic pathway are expressed as a single bifunctional protein, designated UMP synthase (UMPS), although the domain order in *Leishmania* and mammalian cells is reversed (3, 8); and 4) the UMP synthase of *Leishmania* is localized within the glycosome (3), a unique peroxisomal-like organelle that is found uniquely among *Leishmania* and related parasites (9, 10). Genetic ablation of either carbamoyl phosphate synthetase (CPS), the first enzyme of pyrimidine biosynthesis, or UMPS in *L. donovani* conveys pyrimidine auxotrophy that can be circumvented by supplementation of the defined growth medium with uracil, uridine, deoxyuridine, cytidine, or deoxycytidine (3, 5). Furthermore, both the Δcps and the Δumps null mutants exhibit a striking collateral supersensitivity to uracil, which is innocuous to wild type parasites, that is not observed with any of the ribonucleosides (5). A comparable growth susceptibility toward uracil is also observed in other species of protozoan parasites in which the de novo pyrimidine pathway has been genetically disrupted. These uracil-sensitive mutants include Δcps strains of *Toxoplasma gondii* (11) and *Trypanosoma cruzi* (12), the causative agents of toxoplasmplasmis and Chagas disease in humans, respectively, as well as a Δumps null mutant in *T. brucei* (13), which causes African sleeping sickness. Furthermore, repressing expression of dihydroorotate dehydrogenase, the fourth enzyme in the pyrimidine biosynthesis pathway, by RNA interference elicits susceptibility to 5-fluorouracil in *Trypanosoma brucei* (14).

Despite the pyrimidine auxotrophy observed for Δcps and Δumps *L. donovani* promastigotes in culture, both knock-out lines sustain relatively robust infections in mice (Ref. 5 and data not shown). These findings imply that the null mutants within the macrophage phagolysosome can access a source of host pyrimidines that can satisfy the pyrimidine nucleotide requirements of the parasite. Thus, *L. donovani*, in contrast to the purine pathway, has two routes for pyrimidine nucleotide synthesis, biosynthesis, and salvage. Genetic analysis has also authenticated that this salvage of preformed pyrimidines in both promastigotes and amastigotes of *L. donovani* is mediated through uracil phosphoribosyltransferase (LdUPRT) and that pyrimidine nucleosides (other than thymidine) are converted to uracil within the parasite (5). Uridine and deoxyuridine are cleaved to form uracil via nucleoside hydrolase enzymes, whereas a cytidine deaminase converts cytidine and deoxycytidine to their uracil-containing counterparts (15, 16) (Fig. 1). Thus, LdUPRT plays an exclusive role in pyrimidine salvage in the parasite, a function that profoundly impacts the capacity of the parasite to survive as the amastigote in a rodent model.

To characterize the biochemical and kinetic properties of LdUPRT and to evaluate the involvement of LdUPRT in the noteworthy vulnerability of three different genera of protozoan parasite to uracil- or 5-fluorouracil-mediated growth inhibition when the pyrimidine biosynthetic pathway is genetically compromised, recombinant LdUPRT was purified and characterized. Kinetic parameters to the naturally occurring substrates, as well as to several important uracil analogs, were determined, and the *L. donovani* enzyme, unlike its *T. gondii* counterpart (17), was shown to form a stable tetramer in the absence of GTP. Furthermore, profound substrate inhibition of LdUPRT to nucleobase substrates was demonstrated, providing a mechanism by which *L. donovani, T. gondii, T. cruzi*, or *T. brucei*, genetically deficient in pyrimidine biosynthesis, would exhibit a dramatic growth sensitivity to exogenous uracil. It is conjectured that this substrate inhibition of LdUPRT by uracil affords the parasite a protective mechanism to protect its nutritionally indispensable purine salvage mechanism and to maintain an equilibrium between purine and pyrimidine nucleotide pools in the parasite.

**EXPERIMENTAL PROCEDURES**

**Chemicals and Reagents**—Uracil, 5-fluorouracil, 4-thiouracil, PRPP, GTP, isopropyl β-D-1-thiogalactopyranoside (IPTG), and metal salts were purchased from Sigma-Aldrich. Ni-NTA agarose beads were from Qiagen. Complete Mini EDTA-free protease inhibitor was bought from Roche Applied Science. BioSafe Coomassie and Bio-Rad protein dye were acquired from Bio-Rad Laboratories Life Science Research. Oligonucleotide primers were obtained from Integrated DNA Technologies, Inc. (Corvalle, IA), and Phusion® High-Fidelity PCR Master Mix was from Fisher Scientific. Champion™ pET Directional TOPO expression kit was purchased from Invitrogen. The Agilent 8453 UV-visible diode array spectrophotometer was from Agilent Technologies (Santa Clara, CA). All other chemicals and reagents were of the highest quality commercially available.

Expression and Purification of LdUPRT and *T. gondii* UPRT (TgUPRT) in *Escherichia coli*—The cloning of LdUPRT into the pET 200/D-TOPO® *E. coli* expression vector has been previously reported (5). The full-length TgUPRT cDNA was amplified by PCR using the forward primer 5'-GAGGCGCATGGGCAGGATCCAGGAC-3' and reverse primer 5'-GAGGCACAGCCGCTACTAGTTCTAAAA-GTACCGTACCCGAA-3' (SfiII restriction sites are in bold, and unique triads are underlined) from a previously reported TgUPRT cDNA construct (18). The insert was ligated into the pET 200/D-TOPO® vector containing an NH₂-terminal His₆ tag.
tag and transformed into One Shot® Top10 chemically competent E. coli. The fidelity and orientation of the recombinant plasmid was verified by DNA sequencing. The LdUPRT and TgUPRT expression constructs in the pET 200/D-TOPO® vector were transformed into BL21Star™ (DE3) One Shot® E. coli according to the Champion™ pET Directional TOPO user manual and plated on LB plates containing 50 µg/ml kanamycin. Transformants were picked and expanded in 200 ml or 1 liter of LB medium to an A₆₀₀ ~0.6, and protein expression was induced with 1 mM IPTG for 16 h at 37 °C with constant shaking.

The E. coli from the 16-h culture was harvested by centrifugation and resuspended in a buffer containing 50 mM NaH₂PO₄, 300 mM NaCl, 10 mM imidazole, pH 8.0, and EDTA-free protease inhibitors. The cells were ruptured by sonication on ice with six 10-s bursts at 200–300 W with a 10-s cooling period between each burst. The lysate was centrifuged at 10,000 × g for 30 min at 4 °C to pellet the cellular debris, and the supernatant was collected. The clarified lysate was incubated with a 50% Ni-NTA slurry at 4 °C with continuous shaking. The lysate-Ni-NTA mixture was washed twice in 50 mM NaH₂PO₄, 1 mM NaCl, 20 mM imidazole, pH 8.0 buffer and eluted in buffer consisting of 50 mM NaH₂PO₄, 300 mM NaCl, and 250 mM imidazole, pH 8.0. The purified recombinant proteins were buffer-exchanged into a final storage buffer containing 50 mM KCl, 50 mM KH₂PO₄, 5% glycerol, pH 8.0, using 7,000 molecular weight cutoff Zeba™ spin desalting columns (ThermoFisher Scientific). Concentrated LdUPRT and TgUPRT preparations were obtained by ultrafiltration employing Amicon Ultra-10K centrifugal filter units (EMD Millipore Corp., Billerica, MA), and protein concentrations were determined using the Bio-Rad Bradford total protein assay system.

**LdUPRT Kinetics**—All kinetic parameters were determined using a published spectrophotometric method based on monitoring the absorbance change at a specified wavelength under steady state conditions (18, 19). Each assay mixture was prepared by adding the substrates to a buffer containing 50 mM Tris-HCl, 5 mM MgCl₂, and 2 mM DTT, pH 7.5 (TMD 50), unless otherwise noted. For each substrate, the assay mixture was blanked prior to assay to remove background caused by substrates or reagents. Kinetic traces were initiated by addition of LdUPRT enzyme to the pre-equilibrated assay mixture, and data were collected for total of 120 s at a fixed wavelength. The assays were based on the small but significant differential molar absorption coefficients (Δε) between the substrate and product, e.g., uracil and UMP at a given wavelength. The fixed wavelengths and extinction coefficients employed varied depending upon the nucleobase substrate and are described below. All kinetic parameters were calculated by the suite of algorithms available in GraphPad Prism 4.0.

The kinetic parameters for uracil were determined using an assay mixture consisting of TMD 50 buffer, 1 mM PRPP, and various concentrations of uracil ranging from 5 µM to 1.5 mM, whereas the Kₘ value for PRPP was ascertained in TMD 50 buffer, 250 µM uracil, and PRPP concentrations ranging from 25 µM to 2.5 mM. To evaluate the effect of GTP on PRPP kinetics, 2 mM GTP was added to the assay mixture described above. UMP formation was determined at a wavelength of 280 nm, and the rates were calculated using a differential molar extinction coefficient of Δε = 1419 M⁻¹ cm⁻¹.

The activity of LdUPRT toward the nucleobase analogs 5-fluorouracil and 4-thiouracil were determined in TMD 50 buffer, 1 mM PRPP, pH 7.5, containing either 5–150 µM 5-fluorouracil or 1–80 µM 4-thiouracil. Steady state kinetics were performed based on the expenditure of 5-fluorouracil at 303 nm and the formation of 4-thiouridine-5'-monophosphate at 320 nm, respectively. The differential extinction coefficient for 5-fluorouracil was calculated to be Δε = 923 M⁻¹ cm⁻¹ (at pH 7.5; λₘₐₓ, 303 nm), and the differential extinction coefficient for 4-thiouridine-5'-monophosphate was previously determined to be ε = 16300 M⁻¹ cm⁻¹ (at pH 7.5; λₘₐₓ, 320 nm according to the brochure from Jena Bioscience (Jena, Germany).

**Effect of pH on LdUPRT Activity**—The pH optimum of LdUPRT was evaluated by measuring the activity between pH 6.0 and 10.0 at 0.5 pH unit increments and also at pH 5.8 using either a 50 mM CHES, 50 mM Bis-Tris or 50 mM Tricine buffer. Concentrated NaOH and HCl were used to adjust the pH in the three buffer solutions. To measure LdUPRT activity, 250 µM of uracil, 1 mM PRPP, 2 mM MgCl₂, and 2 mM DTT were added to each buffer, and the rate of UMP formation was measured spectrophotometrically as described above.

**Ion Dependence of LdUPRT**—The effects of an assortment of divalent cations on LdUPRT activity was verified in 50 mM Tris-HCl, 2 mM DTT, 250 µM uracil, 1 mM PRPP, pH 7.5, to which 2 mM of one of the following cations was added: MgCl₂, MnCl₂, BaCl₂, CoCl₂, CaCl₂, NiCl₂, and ZnCl₂. Control experiments were performed both in the absence of divalent cation and in the presence of 10 mM EDTA. UMP formation was determined as described above.

**Size Exclusion Chromatography**—Either LdUPRT or TgUPRT at a concentration of 1.0 mg/ml was injected in a volume of 100 µl of 50 mM KCl, 50 mM KH₂PO₄, pH 8.0 buffer onto a Superose 12 10/300 GL column (GE Healthcare) and eluted with 1 column volume of 50 mM KCl, 50 mM KH₂PO₄, pH 8.0 buffer at a flow rate of 0.4 ml/min. Parallel runs were also conducted in the presence of 2 mM GTP in both the loading and elution buffers. Protein in the eluates was monitored by absorption at 280 nm. Estimated molecular weights were calibrated using a gel filtration marker kit from Sigma-Aldrich. Size exclusion chromatography of LdUPRT and TgUPRT was performed after 3-h incubations in 3 mM urea, and 3 mM urea was added to both the loading and elution buffers.

**Parasite Cell Culture**—The creation and characterization of Δuprt, Δcps, and Δumps L. donovani lines have been reported previously (3, 5). Wild type and null mutant promastigotes were continuously cultured in 26 °C in pH 7.4 DME-L medium that was supplemented with 10% Serum Plus® (SAFC Biosciences, Lenexa, KS), 1 mg/ml hemin, and 100 µM hypoxanthine as a purine source. The Δuprt, Δcps, and Δumps transgenic strains were routinely maintained in the drugs in which the homologous gene replacement events were selected.

**Growth Assays**—The abilities of wild type and mutant cells to grow in a range of uracil concentrations (4 µM to 4 mM) were determined by placing 5.0 × 10⁵ promastigotes into individual wells of a 96-well cell culture plate containing 0.2 ml of growth
medium. Additional uracil sensitivity experiments were performed using the same protocol but with either 250 μM cytidine, 2 mM dihydroorotate, or 2 mM orotate added to the growth medium. 5-Fluorouracil and 4-thiouracil sensitivity experiments were conducted using the same protocol, again as a function of multiple 5-fluorouracil (40 nm to 40 μM) or 4-thiouracil (1 μM to 1 mM) concentrations. 250 μM cytidine was added to the growth medium in these growth experiments with the two uracil analogs. At the end of each growth experiment, parasites were counted using the AlamarBlue® (BIOSOURCE) cell viability assay (20). Reduction of AlamarBlue was monitored at 570 and 600 nm on a Multiskan Ascent plate reader (Thermo Labsystems, Vantaa, Finland). The percentage of dye reduction was calculated according to the formula delineated in the manufacturer's pamphlet, and the largest reduction was expressed as maximum growth.

Substrate Inhibition Profiles—The ability of high concentrations of uracil, 5-fluorouracil, and 4-thiouracil to inhibit their own phosphoriboyslation by LdUPRT was gauged in TMD 50 μM concentrations of the uracil or 5-fluorouracil or 75–1500 μM concentrations of 4-thiouracil employing the spectrophotometric methods described above. To evaluate whether uracil-mediated substrate inhibition was reversible or irreversible, 2.0 μg of purified LdUPRT was incubated in the absence or presence of either 1.0 mM PRPP, 1.5 mM uracil, or both 1.0 mM PRPP and 1.5 mM uracil for 5 min and diluted in TMD buffer just prior to assay, and LdUPRT activity assessed in the presence of 1.0 mM PRPP and either 75 μM or 1.5 mM uracil. The thermostability of LdUPRT was evaluated by incubating 2.0-μg aliquots of purified LdUPRT in the absence or presence of either 1.0 mM PRPP, 75 μM uracil, 1.5 mM uracil, both 1.0 mM PRPP and 75 μM uracil, or both 1.0 mM PRPP and 1.5 mM uracil at 62 °C for various time points up to 20 min; diluting into TMD buffer; and assaying residual LdUPRT activity in TMD buffer to which 1.0 mM PRPP and 200 μM uracil were added.

RESULTS

LdUPRT Expression and Purification—LdUPRT was robustly expressed from the pET 200/D-TOPO construct using the BL21Star™ (DE3) One Shot® E. coli expression system and IPTG induction (Fig. 2). A visible band was observed at 27 kDa, consistent with the predicted molecular mass (Fig. 2). This band was not observed in uninduced E. coli. His6-LdUPRT (henceforth just referred to as LdUPRT) was subsequently purified to homogeneity over a Ni-NTA column.

Substrate inhibition profiles of LdUPRT for the two nucleobase analogs were similar, 1.1-fold greater than the experimentally determined K_m, value. Because the affinity of TgUPRT, the only uracil phosphoribosyltransferase (UPRT) of parasitic origin that has been previously characterized, for PRPP was reduced by the addition of GTP (17), the K_m, value for PRPP was also determined in the presence of GTP. Incubation of LdUPRT with 2 mM GTP treatment did not, however, affect the K_m, value of LdUPRT for PRPP (Fig. 4B). A V_max, value of 13.6 ± 1.4 μmol/min/mg protein for LdUPRT was computationally derived (Fig. 4, A and B). A k_cat, value of 6.19 s⁻¹ was then calculated from the kinetic data, and the catalytic efficiency (k_cat/K_m) was computed to be 0.303 s⁻¹ μmol⁻¹, LdUPRT also displayed high affinities for the pyrimidine analogs 4-thiouracil and 5-fluorouracil with calculated K_m values of 7.1 and 6.4 μM, respectively (Fig. 4, C and D). The calculated V_max values of LdUPRT for the two nucleobase analogs were similar, 1.1 ± 0.18 and 1.3 ± 0.12 μmol/min/mg protein for 4-thiouracil and 5-fluorouracil, respectively.

Oligomerization State—Size exclusion chromatography indicated that LdUPRT migrated with a molecular mass just under 100 kDa, consistent with a tetrameric oligomerization state (Fig. 5). In contrast, purified recombinant TgUPRT migrated on the size exclusion column with a molecular mass consistent with a dimeric quaternary state. A dimeric structure has been previously reported for TgUPRT based on its sedimentation properties in sucrose gradients (17). The addition of 2 mM GDP to TgUPRT, known to stabilize higher order structures of the protein (17), induced an oligomeric state that was either a trimeric or quaterneric structure (Fig. 5). The quaternary structure of LdUPRT, as expected, was unaffected by 2 mM GTP (data not shown).

Effect of Added Pyrimidines on the Sensitivity of L. donovani to Uracil—It has been previously reported that the growth of L. donovani rendered auxotrophic for pyrimidines through
genetic lesions in the de novo pyrimidine biosynthesis pathway, specifically strains in which either the CPS or UMPS ORFs have been deleted, is inhibited by high concentrations of uracil in the growth medium (5). Uracil is not, however, growth inhibitory to wild type L. donovani (5). Similarly, T. gondii (11), T. cruzi (12), and T. brucei (13) with genetic defects in pyrimidine biosynthesis pathway are susceptible to uracil-mediated growth inhibition, whereas their wild type counterparts are not. The previously published uracil sensitivity experiments with L. donovani were conducted in the absence of additional pyrimidines in the culture medium (5). To determine whether this curious growth inhibitory effect of uracil on pyrimidine auxotrophic L. donovani was impacted by the presence of other pyrimidines, the uracil susceptibility of Δcps and Δumps promastigotes to uracil was evaluated in the absence or presence of pyrimidine biosynthetic or salvage intermediates (Fig. 6). Whereas both Δcps and Δumps promastigotes were sensitive to uracil in the absence or presence of 250 μM cytidine in the culture medium, the uracil supersensitivity of the Δcps line was abrogated by the addition of either 2 mM orotate or 2 mM dihydroorotate (Fig. 6). Neither orotate nor dihydroorotate, however, affected the sensitivity of the Δumps null mutant to uracil (Fig. 6, C and D).

**UPRT from L. donovani**

![Figure 3. pH and divalent cation profiles of LdUPRT.](image1)

**Figure 3.** pH and divalent cation profiles of LdUPRT. Initial rates of LdUPRT activity were determined as a function of pH at 0.5 pH units from pH 6.0 to 10.0 and at pH 5.8 as described under “Experimental Procedures.” The data are presented as percentages of maximum activity (pH 8.0) as a function of pH (A). LdUPRT activity was also assessed as a function of the divalent cation in the assay mixture (B). All cations were present at a concentration of 2 mM as the chloride salt. Controls included no divalent cation and 10 mM EDTA. The data are calculated as percentages of maximum LdUPRT activity. The data points in both panels are the averages ± standard deviations obtained for three separate experiments.

![Figure 4. Michaelis-Menten kinetics for LdUPRT.](image2)

**Figure 4.** Michaelis-Menten kinetics for LdUPRT. LdUPRT activity was measured spectrophotometrically as a function of uracil concentration at a fixed 1.0 mM PRPP concentration (A) and as a function of PRPP concentration at 250 μM uracil in the absence and presence of 2 mM GTP (B). Michael-Menten kinetics were also collected as a function of 4-thiouracil (C) and 5-fluorouracil (D) concentrations in the presence of 1.0 mM PRPP. All data are the means ± standard deviations of three replicates. Kinetic parameters were calculated in GraphPad Prism 4.0.
implied that uracil was triggering a pyrimidine deficiency explicitly in cells with genetic lesions in pyrimidine biosynthesis. One possible mechanism by which pyrimidine deficiency could be selectively induced in pyrimidine auxotrophic L. donovani is by uracil-mediated substrate inhibition of LdUPRT. To test this conjecture, the ability of high concentrations of uracil to inhibit nucleoside phosphoribosylation by LdUPRT was determined. As shown in Fig. 7A, concentrations of uracil 10-fold higher than the $K_m$ value dramatically diminished the capacity of LdUPRT to convert uracil to UMP. 1.5 mM uracil diminished LdUPRT activity by $\sim 90\%$. To test whether TgUPRT was also prone to substrate inhibition by uracil, TgUPRT activity was also measured as a function of uracil concentration in the assay. As shown in Fig. 7B, TgUPRT catalytic activity is also markedly inhibited by high uracil in a dose-dependent manner similar to the pattern of inhibition obtained with LdUPRT.

To examine in more detail the mechanism by which uracil elicited inhibition of LdUPRT, the reversibility of this substrate inhibition was examined. Purified LdUPRT samples were preincubated with 1.0 mM PRPP, 1.5 mM uracil, or both 1.0 mM PRPP and 1.5 mM uracil and then examined for LdUPRT activity. None of the preincubation conditions affected LdUPRT activity, and the activity detected remained sensitive to inhibition by high uracil concentrations (Fig. 8A). The thermobility of LdUPRT was also tested using the same conditions as those employed for the reversibility assays. 1.0 mM PRPP stabilized LdUPRT to heat inactivation at 62°C, whereas 1.5 mM uracil did not impact thermobility. Furthermore, addition of either 75 $\mu$M or 1.5 mM uracil to the enzyme in the absence or presence of 1.0 mM PRPP had no effect on the heat inactivation profile of the enzyme (Fig. 8B and data not shown). Attempts to examine changes in LdUPRT secondary structure induced by uracil by circular dichroism spectroscopy were precluded by the high level of absorbance of 1.5 mM uracil in the far ultraviolet spectral range.

Inhibition of LdUPRT Activity and L. donovani Growth by Uracil Analogs—LdUPRT was also susceptible to substrate inhibition by high concentrations of the uracil analogs 4-thiouracil and 5-fluorouracil (Fig. 7, C and D). To determine whether the substrate sensitivity of LdUPRT to 4-thiouracil and 5-fluorouracil could also affect the growth susceptibility of $\Delta cps$ and $\Delta uprt$ L. donovani to the two uracil analogs, the growth of wild type and the two pyrimidine auxotrophs was assessed over a range of analog concentrations. These experiments were performed in medium supplemented with 250 $\mu$M cytidine, which does not affect uracil sensitivity of $\Delta cps$ and $\Delta uprt$ L. donovani (Fig. 6B) but is required for their survival and growth. Both the $\Delta cps$ and $\Delta uprt$ null mutants exhibited sensitivity to 4-thiouracil with EC$_{50}$ values of 82.3 and 67.3 $\mu$M, respectively, whereas wild type parasites, as well as a $\Delta uprt$ L. donovani strain, were refractory to 4-thiouracil concentrations >500 $\mu$M killed the mutant parasites. The pyrimidine auxotrophs were also more sensitive to 5-fluorouracil than the wild type or $\Delta uprt$ mutant, although the differences were not as dramatic as for uracil (Fig. 6A) and 4-thiouracil (Fig. 9A). EC$_{50}$ values obtained for wild type, $\Delta uprt$, $\Delta cps$, and $\Delta uprt$ promastigotes were 1.65, 1.33, 0.49, and 0.24 $\mu$M, respectively (Fig. 9B). These 5-fluorouracil growth sensitivity experiments were also carried out in growth medium supplemented with 250 $\mu$M cytidine as the requisite source of preformed pyrimidine for the pyrimidine biosynthesis null mutants.

DISCUSSION

A genetic dissection of the pyrimidine pathway has established that LdUPRT is the sole enzyme capable of salvaging preformed pyrimidines to the nucleotide level in L. donovani and that all pyrimidine nucleosides that can satisfy the nutritional requirements of strains genetically auxotrophic for pyrimidines are ultimately converted to uracil prior to phosphoribosylation by LdUPRT (5). Introduction of a $\Delta uprt$ lesion into wild type L. donovani obliterates pyrimidine salvage in both life cycle stages of the parasite and, when introduced into a $\Delta cps$ line that lacks an intact biosynthetic pathway but is still capable of manifesting a robust infection, reduces parasite burdens in both liver and spleen to zero (5). To investigate this key pyrimidine salvage enzyme in more detail, recombinant LdUPRT was purified to homogeneity, and its kinetic parameters were determined (Figs. 2 and 4). LdUPRT displays a neutral pH optimum, requires a divalent cation for activity, and exhibits affinities toward its naturally occurring substrates that are equivalent to that previously reported for the T. gondii enzyme (18). LdUPRT also recognizes the uracil analogs 4-thiouracil and 5-fluorouracil with affinities similar to that of uracil (Fig. 4, C and D). Molecular sizing of LdUPRT implied that the L. donovani enzyme was a physiologically active tetramer (Fig. 5), similar to the T. gondii counterpart for which high resolution crystal structures were determined in a number of catalytic states (17, 21). In contrast to the T. gondii enzyme (17), the activity of LdUPRT was not augmented by GTP (Fig. 4B). It has been previously observed that $\Delta cps$ and $\Delta uprt$ L. donovani promastigotes exhibit a collateral supersensitivity to ura-
cil, a nucleobase that is neither growth inhibitory nor cytotoxic toward wild type parasites (5). This effect of uracil on pyrimidine auxotrophic L. donovani is specific for the nucleobase and is growth inhibitory in nature rather than lethal (5). This substrate inhibition of LdUPRT by uracil was reversible and did not affect the thermostability of the enzyme (Fig. 8). The fact that

![Figure 6. Inhibition of L. donovani growth and LdUPRT activity by high substrate concentrations.](image)

![Figure 7. Substrate inhibition of LdUPRT and TgUPRT.](image)
two distinct genetic lesions in the pyrimidine pathway instigate this susceptibility to uracil substantiates that it is triggered by a deficit in pyrimidine biosynthesis capacity and not by some ancillary event in either of the null mutant lines. Analogous pyrimidine biosynthetic mutants of other protozoan parasites, including Δcps T. gondii (11), Δcps T. cruzi (12), and Δumps T. brucei (13), all display this predisposition to be growth inhibited by high concentrations of uracil in the culture medium. Heretofore, no mechanism has been established for this intriguing growth inhibition of the normally nondetrimental nucleobase toward pyrimidine biosynthesis mutants of protozoan parasites, although Ali et al. (13) conjectured that excessive uracil influx might bring about nucleotide pool imbalances in T. brucei.

The ability of orotate and dihydroorotate, which are both metabolic intermediates in the pyrimidine biosynthetic pathway (Fig. 1), to eliminate the uracil hypersensitivity of the Δcps line (Fig. 6, C and D) implied that the provision of an exogenous source of pyrimidine nucleotides was alleviating a pyrimidine starvation state that was being instigated by uracil. Cytidine, however, which is deaminated to uridine, cleaved to uracil, phosphoribosylated to UMP, and ultimately distributed into all pyrimidines in the parasite, does not impact the uracil supersensitivity of the Δcps or Δumps strain (Fig. 6B). It should be noted that neither orotate nor dihydroorotate, which cannot be converted into UMP by L. donovani harboring a Δumps lesion (Fig. 1), alleviated the susceptibility of the Δumps strain to uracil (Fig. 6, C and D). Taken together, we conjectured that uracil at high concentrations was instigating an intracellular depletion of pyrimidine nucleotides by impairing salvage because the growth inhibition in the Δcps knock-out could be alleviated by de novo production of UMP but not by a precursor of UMP synthesis that is incorporated through LdUPRT. This hypothesis was tested directly by verifying that high concentrations of uracil (>10-fold K_m concentrations) triggered substrate inhibition of LdUPRT that inhibited activity >90% at the highest uracil concentration tested, i.e., 1.5 mM (Fig. 7A). Purified TgUPRT was also inhibited by high uracil concentrations (Fig. 7B), intimating that substrate inhibition by uracil may be a common feature of protozoan UPRTs. Substrate inhibition of UPRT by uracil remains to be evaluated for the T. cruzi and T. brucei UPRT enzymes but is a plausible mechanism to account for the susceptibility of pyrimidine auxotrophs to uracil-instigated growth inhibition in those species. Both T. cruzi and T. brucei accomplish a UPRT gene within their respective genomes (2, 22, 23), and the UPRT enzyme has been detected in both species (24), although neither has been investigated in detail. The biological significance for LdUPRT substrate inhibition may pertain to PRPP sparing. PRPP is a critical substrate for purine salvage, an indispensable nutritional function for all protozoan parasites (1), and a process that is known to be mediated through two PRPP-dependent phosphoribosyl-
transfers: HGPRT and XPRT, in *L. donovani* (25). Thus, high concentrations of the nucleobase would prioritize PRPP for usage in purine salvage when pyrimidine pools are replete. Because *L. donovani* amastigotes within the phagolysosome are exposed to RNA degradation products that they can salvage, this substrate inhibition of LdUPRT by uracil can also ensure the maintenance of a balanced supply of pyrimidine and purine nucleotides for the parasite under conditions when pyrimidine pools are replete.

Similarly, 4-thiouracil, which has been used as a tag for evaluating gene expression and transcriptional profiling on a genome-wide level (26, 27), was not toxic to wild type *L. donovani* promastigotes, although the Δcps and Δumps null mutants were killed by high concentrations of the analog (Fig. 9A). The marked inhibition of LdUPRT activity by concentrations of 4-thiouracil that also induced parasite growth inhibition (Fig. 9A) implicates the disruption of pyrimidine salvage as the sole mechanism of growth disruption induced by 4-thiouracil treatment of the pyrimidine auxotrophs. This conclusion is supported by the observed greater sensitivity of LdUPRT to uracil (Fig. 7, A and C) that was reflected in the greater sensitivity of the Δcps and Δumps parasites to growth inhibition by 4-thiouracil compared with uracil (Figs. 6A and 8A).

Although LdUPRT was also predisposed to substrate inhibition by high levels of 5-fluorouracil (Fig. 7D), it is difficult to reconcile this incomplete substrate inhibition of the enzyme with the observed toxicity toward *L. donovani* promastigotes that was observed at concentrations of 5-fluorouracil 2–3 orders of magnitude lower than that which inhibit LdUPRT. Although the precise mechanism of 5-fluorouracil toxicity toward *L. donovani* promastigotes is unknown, the Δuprt cell line, which is capable of de novo pyrimidine synthesis, was just as susceptible to 5-fluorouracil as the wild type strain (Fig. 9B). Despite the demonstration herein that 5-fluorouracil is a substrate for LdUPRT, *L. donovani* promastigotes in which LdUPRT has been genetically deleted (Δuprt) are essentially as sensitive to 5-fluorouracil as wild type parasites (EC<sub>50</sub> = 1.65 μM versus 1.33 μM, respectively; Fig. 9B), revealing another comparably efficient means for phosphoribosylating 5-fluorouracil. Collectively, these data intimate that 5-fluorouracil is phosphoribosylated via both LdUPRT and orotate phosphoribosyltransferase, a component of the UMPs bifunctional enzyme (Fig. 1), and biochemical evidence that purified recombinant UMPs is capable of phosphoribosylating 5-fluorouracil directly supports this contention (data not shown). Because it is not feasible to generate a conditionally lethal Δumps/Δuprt double knock-out caused by the lack of a pyrimidine salvage bypass mechanism (Fig. 1), it cannot be definitively determined using genetic approaches that UPRT and UMPs are the exclusive routes by which the fluorinated pyrimidine is salvaged. Collectively, these data suggest that the production of the 5-fluoro-UMP from 5-fluorouracil by LdUPRT and/or UMPs is sufficient to account for the observed toxicity of the fluorinated pyrimidine in pyrimidine prototrophs. The enhanced 5-fluorouracil toxicity observed in pyrimidine auxotrophs is likely more complicated but may result from a combination of factors including competition between 5-fluorouracil and uracil for phosphoribosylation by LdUPRT, competition between UMP and 5-fluoro-UMP for further metabolism by downstream enzymes, and/or more efficient incorporation of 5-fluorouracil into the nucleotide pool and RNA in the absence of de novo UMP production. This is consistent with the model proposed by Ali *et al.* (13) to explain the hypersensitivity of *T. brucei* pyrimidine auxotrophs to 5-fluorouracil.

In summary, we have performed a biochemical and kinetic characterization of LdUPRT, the sole enzyme capable of incorporating preformed host pyrimidines into the parasite nucleotide pool. The kinetic characterization of LdUPRT revealed a remarkable inhibition of the enzyme by its uracil substrate that can account for the unique susceptibility of *L. donovani* harboring genetic lesions in the pyrimidine biosynthetic pathway to the nucleobase. This substrate inhibition of UPRT enzymes appears to be a general mechanism by which purine auxotrophs can equilibrate purine and pyrimidine nucleotide pools and offers a means by which purine incorporation, an essential nutritional function, is ceded preference over pyrimidine salvage, a nonessential process, when the parasites have access to nucleotide precursors. Finally, because mammalian cells lack an analogous UPRT enzyme (28), the capacity of LdUPRT to recognize cytotoxic nucleobase analogs offers a potential therapeutic strategy by which cytotoxic uracil derivatives could be selectively incorporated into the parasite nucleotide pool without impacting the human host. 5-Fluorouracil and 4-thiouracil, however, are not prospective candidates as pro-drugs for which LdUPRT activation is required because 5-fluorouracil, an anti-neoplastic agent employed pervasively in the treatment of gastrointestinal cancers (29), exhibits unacceptable toxicity toward mammalian cells (30), whereas 4-thiouracil is nontoxic toward wild type *L. donovani* (Fig. 9A). A structure-activity study of uracil analogs has been previously carried out with a partially purified TgUPRT preparation (31), but no similar study has been performed with LdUPRT. Extrapolating the results of such a structure-activity analysis performed on LdUPRT to intact *Leishmania* parasites, however, is complicated somewhat by the existence of another pyrimidine phosphoribosyltransferase activity, i.e., the orotate phosphoribosyltransferase, the penultimate enzyme in de novo pyrimidine biosynthesis that can also presumably recognize pyrimidine analogs (3). Indeed, 5-fluorouracil is a known substrate for the mammalian orotate phosphoribosyltransferase enzyme (32). Regardless, the absence of a mammalian equivalent raises the possibility of exploiting LdUPRT as a selective mechanism for activating potential antileishmanial pyrimidine nucleobase analogs.

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UPRT from L. donovani

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