

# 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 *L. 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  $\mu\text{M}$  for the natural substrates uracil and phosphoribosylpyrophosphate, respectively, as well as apparent  $K_m$  values 6 and 7  $\mu\text{M}$  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. *L. 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  $\text{CO}_2$ , amino acids, and 5-phosphoribosyl-1-pyrophosphate (PRPP)<sup>2</sup> (Fig. 1). UMP is then distributed into ribonucleotides via nucleotide kinases and into deoxyribonucleotides by ribonucleotide reductase and 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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<sup>2</sup> 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; UPRT, uracil phosphoribosyltransferase; Ni-NTA, nickel-nitrilotriacetic acid; CHES, 2-(cyclohexylamino)ethanesulfonic acid; Tricine, N-[2-hydroxy-1,1-bis(hydroxymethyl)ethyl]glycine.

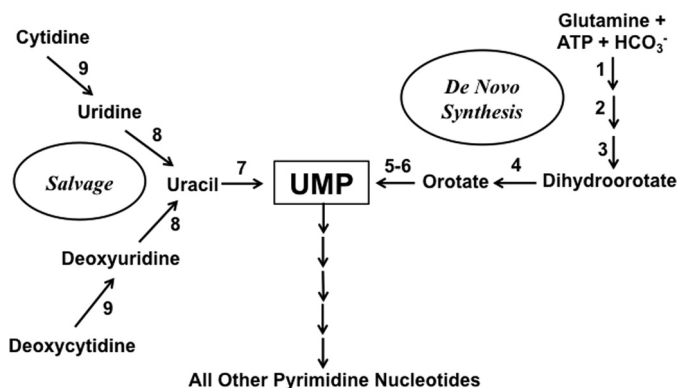


FIGURE 1. **Schematic diagram of the pyrimidine biosynthetic and salvage pathways of *L. donovani*.** The portions of the pyrimidine biosynthetic and pyrimidine salvage pathways that are pertinent to this manuscript are depicted. The following enzymes are depicted: CPS (1), aspartate transcarbamoylase (2), dihydroorotase (3), dihydroorotate dehydrogenase (4), orotate phosphoribosyltransferase (5), orotidylate decarboxylase (6), LdUPRT (7), uridine nucleosidase (8), and cytidine deaminase (9). Orotate phosphoribosyltransferase and orotidylate decarboxylase are the enzymatic activities encoded by UMPS.

crete, 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* confer 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  $\Delta cps$  and the  $\Delta umps$  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  $\Delta cps$  strains of *Toxoplasma gondii* (11) and *Trypanosoma cruzi* (12), the causative agents of toxoplasmosis and Chagas disease in humans, respectively, as well as a  $\Delta umps$  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  $\Delta cps$  and  $\Delta umps$  *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  $\beta$ -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. (Coralville, IA), and Phusion<sup>®</sup> High-Fidelity PCR Master Mix was from Fisher Scientific. Champion<sup>™</sup> 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<sup>®</sup> *E. coli* expression vector has been previously reported (5). The full-length *TgUPRT* cDNA was amplified by PCR using the forward primer 5'-GAGGCCAC-CTGGGCCATGGCGCAGGTCCAGCGAG-3' and reverse primer 5'-GAGGCCAGCCCGGCCCTACATGGTTCCAAAGTACCGGTCACCGAA-3' (SfiI restriction sites are in bold, and unique triads are underlined) from a previously reported *TgUPRT* cDNA construct (18). The insert was ligated in to the pET 200/D-TOPO<sup>®</sup> vector containing an NH<sub>2</sub>-terminal His<sub>6</sub>

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  $A_{600} \sim 0.6$ , 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  $\text{NaH}_2\text{PO}_4$ , 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  $10,000 \times g$  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 for 1 h with continuous shaking. The lysate-Ni-NTA mixture was loaded onto a column and washed twice in 50 mM  $\text{NaH}_2\text{PO}_4$ , 1 M NaCl, 20 mM imidazole, pH 8.0 buffer and eluted in buffer consisting of 50 mM  $\text{NaH}_2\text{PO}_4$ , 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  $\text{KH}_2\text{PO}_4$ , 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  $\text{MgCl}_2$ , 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 ( $\Delta\epsilon$ ) 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  $K_m$  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  $\Delta\epsilon = 1419 \text{ M}^{-1} \text{ cm}^{-1}$ .

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  $\Delta\epsilon = 923 \text{ M}^{-1} \text{ cm}^{-1}$  (at pH 7.5;  $\lambda_{\text{max}}$  303 nm), and the differential extinction coefficient for 4-thiouridine-5'-monophosphate was previously determined to be  $\epsilon = 16300 \text{ M}^{-1} \text{ cm}^{-1}$  (at pH 7.5;  $\lambda_{\text{max}}$  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  $\text{MgCl}_2$ , 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:  $\text{MgCl}_2$ ,  $\text{MnCl}_2$ ,  $\text{BaCl}_2$ ,  $\text{CoCl}_2$ ,  $\text{CaCl}_2$ ,  $\text{NiCl}_2$ , and  $\text{ZnCl}_2$ . 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  $\text{KH}_2\text{PO}_4$ , 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  $\text{KH}_2\text{PO}_4$ , 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 M urea, and 3 M urea was added to both the loading and elution buffers.

**Parasite Cell Culture**—The creation and characterization of  $\Delta\text{uprt}$ ,  $\Delta\text{cps}$ , and  $\Delta\text{umps}$  *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  $\Delta\text{uprt}$ ,  $\Delta\text{cps}$ , and  $\Delta\text{umps}$  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  $5.0 \times 10^3$  promastigotes into individual wells of a 96-well cell culture plate containing 0.2 ml of growth

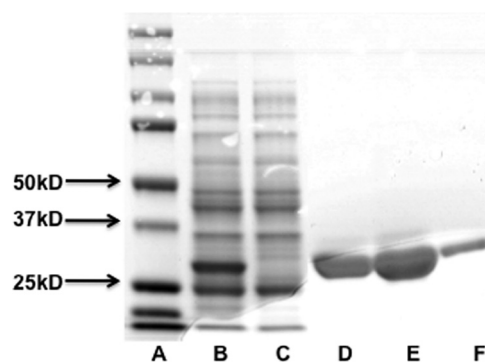
medium. Additional uracil sensitivity experiments were performed using the same protocol but with either 250  $\mu\text{M}$  cytidine, 2 mM dihydroorotate, or 2 mM orotate added to the growth medium. 5-Fluorouracil and 4-thiouracil sensitivity experiments were conducted using the same protocol, again as a function of multiple 5-fluorouracil (40 nM to 40  $\mu\text{M}$ ) or 4-thiouracil (1  $\mu\text{M}$  to 1 mM) concentrations. 250  $\mu\text{M}$  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<sup>®</sup> (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 phosphoribosylation by LdUPRT was gauged in TMD 50 buffer containing 1 mM PRPP and 150–1500  $\mu\text{M}$  concentrations of the uracil or 5-fluorouracil or 75–1500  $\mu\text{M}$  concentrations of 4-thiouracil employing the spectrophotometric methods described above. To evaluate whether uracil-mediated substrate inhibition was reversible or irreversible, 2.0  $\mu\text{g}$  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  $\mu\text{M}$  or 1.5 mM uracil. The thermostability of LdUPRT was evaluated by incubating 2.0- $\mu\text{g}$  aliquots of purified LdUPRT in the absence or presence of either 1.0 mM PRPP, 75  $\mu\text{M}$  uracil, 1.5 mM uracil, both 1.0 mM PRPP and 75  $\mu\text{M}$  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  $\mu\text{M}$  uracil were added.

## RESULTS

**LdUPRT Expression and Purification**—LdUPRT was robustly expressed from the pET 200/D-TOPO construct using the BL21Star<sup>™</sup> (DE3) One Shot<sup>®</sup> *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*. His<sub>6</sub>-LdUPRT (henceforth just referred to as LdUPRT) was subsequently purified to virtual homogeneity by affinity chromatography (Fig. 2, lanes D–F). Overall yield was roughly 4–6 mg of purified recombinant protein per liter of bacterial culture. The purified LdUPRT was stable with no measurable loss of enzymatic activity for >1 month when maintained at 4 °C in the elution buffer to which 5% glycerol was added.

**Enzyme Kinetics**—Freshly purified recombinant LdUPRT was enzymatically active and catalyzed the phosphoribosylation of uracil efficiently. LdUPRT displayed a pH optimum between 7.5 and 8.5 with sharp drops in activity 0.5 pH units outside this range, as well as an absolute requirement for the presence of a divalent cation (Fig. 3). Maximum activity of LdUPRT was achieved with Mg<sup>2+</sup>, although LdUPRT was almost



**FIGURE 2. Purification of LdUPRT.** LdUPRT was overexpressed in *E. coli* and purified to homogeneity over a Ni-NTA column. Lane A, Precision Plus Protein Standards; lane B, 10,000  $\times$  g crude cell lysates of IPTG-treated BL21Star<sup>™</sup> (DE3) One Shot<sup>®</sup> *E. coli* transformed with LdUPRT; lane C, 10,000  $\times$  g cell lysates of uninduced BL21Star<sup>™</sup> (DE3) One Shot<sup>®</sup> *E. coli*; lanes D–F, final elution fractions from the Ni-NTA column. Molecular weight markers are shown in lane A.

as efficient with Co<sup>2+</sup> and Mn<sup>2+</sup>. Ba<sup>2+</sup>, Ca<sup>2+</sup>, Ni<sup>2+</sup>, and Zn<sup>2+</sup> did not support LdUPRT catalysis. Michaelis-Menten plots of steady state kinetic data obtained at pH 7.5 and 5.0 mM MgCl<sub>2</sub> revealed apparent  $K_m$  values of 20.4 and 99.3  $\mu\text{M}$  for uracil and PRPP, respectively (Fig. 4, A and B). The kinetic data for the bisubstrate reaction were collected at concentrations of invariant substrate that were  $\sim$ 10-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  $V_{\text{max}}$  value of  $13.6 \pm 1.4$   $\mu\text{mol}/\text{min}/\text{mg}$  protein for LdUPRT was computationally derived (Fig. 4, A and B). A  $k_{\text{cat}}$  value of  $6.19 \text{ s}^{-1}$  was then calculated from the kinetic data, and the catalytic efficiency ( $k_{\text{cat}}/K_m$ ) was computed to be  $0.303 \text{ s}^{-1} \mu\text{M}^{-1}$ . 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  $\mu\text{M}$ , respectively (Fig. 4, C and D). The calculated  $V_{\text{max}}$  values of LdUPRT for the two nucleobase analogs were similar,  $1.1 \pm 0.18$  and  $1.3 \pm 0.12$   $\mu\text{mol}/\text{min}/\text{mg}$  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 GTP to TgUPRT, known to stabilize higher order structures of the protein (17), induced an oligomeric state that was either a trimeric or quaternary 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

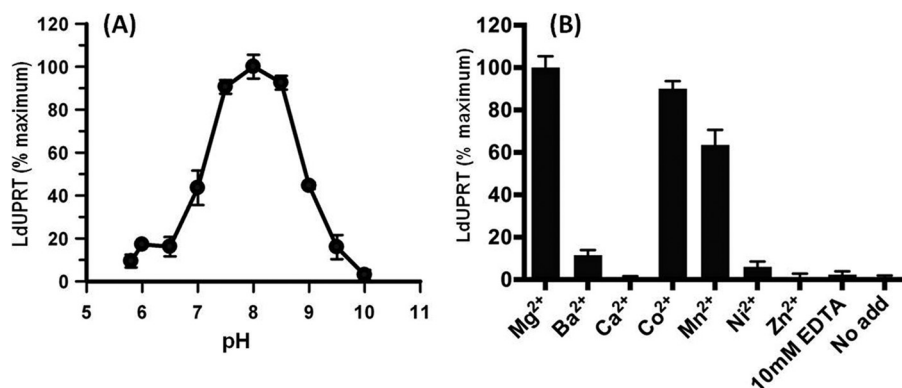


FIGURE 3. **pH and divalent cation profiles of LdUPRT.** Initial rates of LdUPRT activity were determined as a function of pH at 0.5 pH units from pH 6.0 to 10.0 and at pH 5.8 as described under "Experimental Procedures." The data are presented as percentages of maximum activity (pH 8.0) as a function 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  $\pm$  standard deviations obtained for three separate experiments.

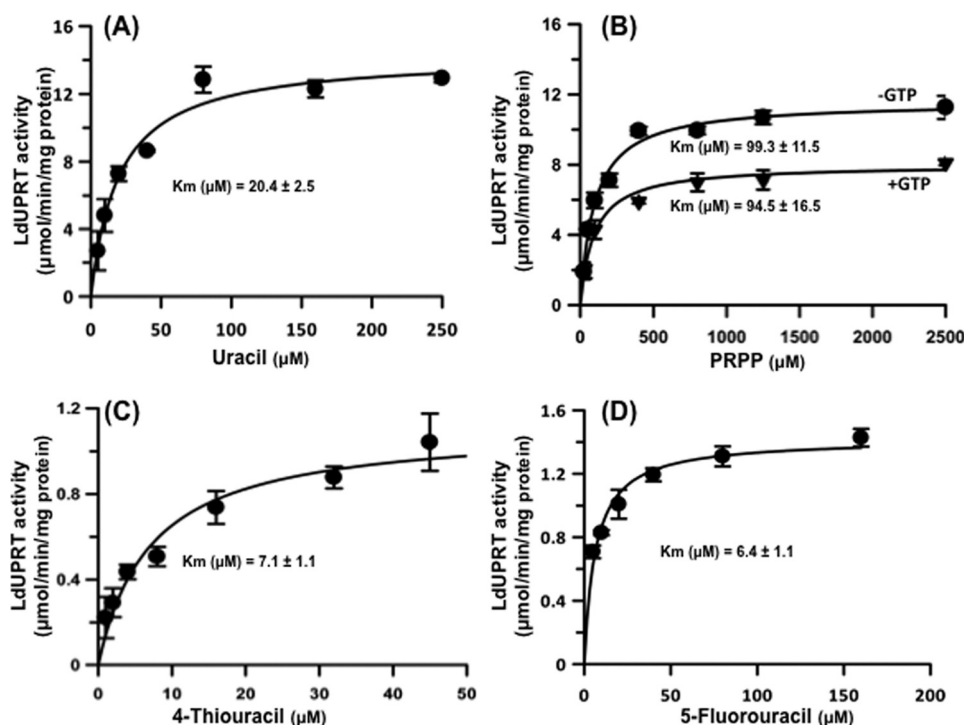
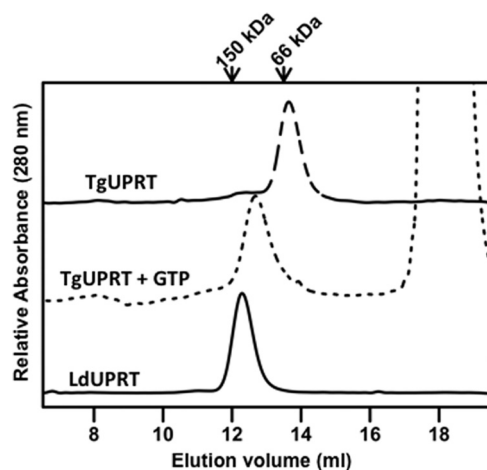


FIGURE 4. **Michaelis-Menten kinetics for LdUPRT.** LdUPRT activity was measured spectrophotometrically as a function of uracil concentration at a fixed 1.0 mM PRPP concentration (A) and as a function of PRPP concentration at 250  $\mu\text{M}$  uracil in the absence and presence of 2 mM GTP (B). Michael-Menten 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  $\pm$  standard deviations of three replicates. Kinetic parameters were calculated in GraphPad Prism 4.0.

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 pyrimi-

dines, the uracil susceptibility of  $\Delta cps$  and  $\Delta umps$  promastigotes to uracil was evaluated in the absence or presence of pyrimidine biosynthetic or salvage intermediates (Fig. 6). Whereas both  $\Delta cps$  and  $\Delta umps$  promastigotes were sensitive to uracil in the absence or presence of 250  $\mu\text{M}$  cytidine in the culture medium, the uracil supersensitivity of the  $\Delta cps$  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  $\Delta umps$  null mutant to uracil (Fig. 6, C and D).

**LdUPRT Substrate Inhibition**—The ability of dihydroorotate and orotate to alleviate uracil-mediated growth inhibition of *L. donovani* promastigotes harboring a genetic lesion in *CPS*



**FIGURE 5. Size exclusion chromatography of purified LdUPRT and TgUPRT.** 1.0 mg/ml purified TgUPRT in the absence or presence of 2 mM GTP or purified LdUPRT were chromatographed over a Superose 12 10/300 GL column, and eluted protein was monitored by absorbance at 280 nm and plotted as a function of elution volume. Molecular weight standards included carbonic anhydrase ( $M_r = \sim 29$  kDa), bovine serum albumin ( $M_r = \sim 66$  kDa), and alcohol dehydrogenase ( $M_r = \sim 150$  kDa).

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 nucleobase 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 thermolability 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 thermolability. 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  $\Delta umps$  *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  $\Delta umps$  *L. donovani* (Fig. 6B) but is required for their survival and growth. Both the  $\Delta cps$  and  $\Delta umps$  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 in the medium as high as 1 mM (Fig. 9A). Unlike uracil, which was growth inhibitory toward the pyrimidine auxotrophs, 4-thiouracil at 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  $\Delta umps$  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  $\Delta umps$  *L. donovani* promastigotes exhibit a collateral supersensitivity to ura-

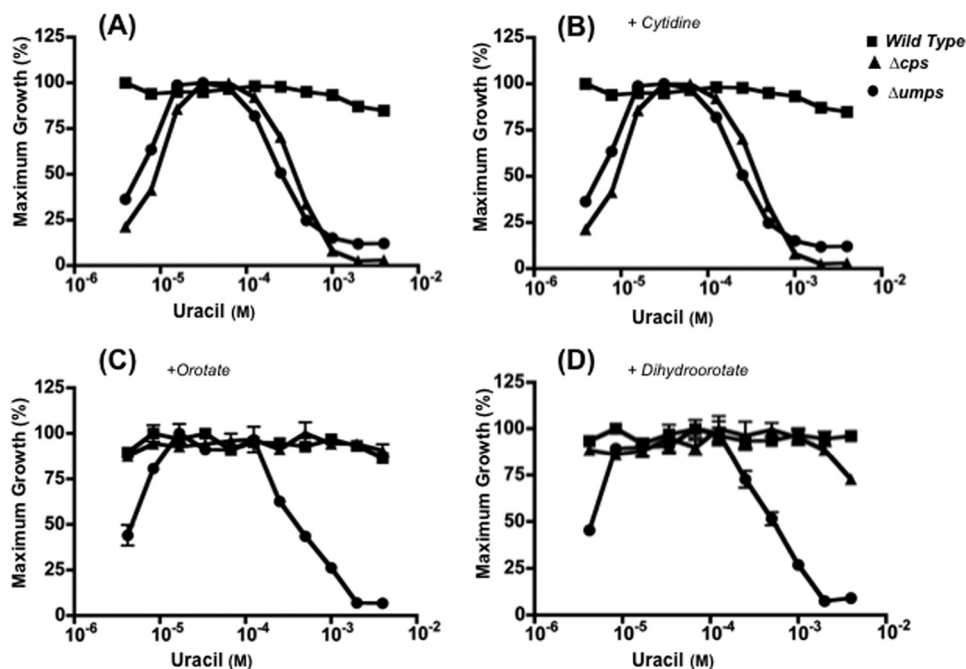


FIGURE 6. Inhibition of *L. donovani* growth and LdUPRT activity by high substrate concentrations. The abilities of wild type (■),  $\Delta cps$  (▲), and  $\Delta ump s$  (●) *L. donovani* promastigotes to grow in various concentrations of uracil (A) or in various concentrations of uracil plus 250  $\mu M$  cytidine (B), 2 mM orotate (C), or 2 mM dihydroorotate (D) were assessed as described under "Experimental Procedures." The data are those from one of at least three independent experiments, all of which gave virtually identical results.

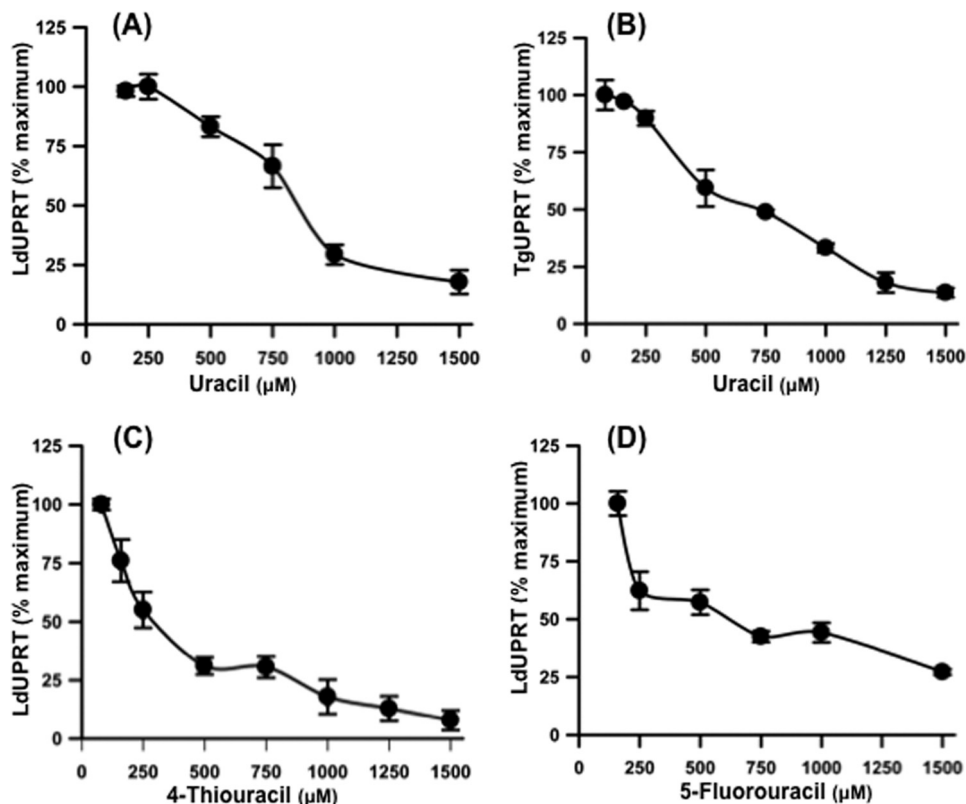


FIGURE 7. Substrate inhibition of LdUPRT and TgUPRT. LdUPRT (A) and TgUPRT (B) activity was determined spectrophotometrically at a variety of high uracil concentrations in the assay mixture. LdUPRT activity was also ascertained spectrophotometrically as a function of high concentrations of 4-thiouracil (C) or 5-fluorouracil (D). The data presented are the means  $\pm$  standard deviations from three distinct measurements.

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

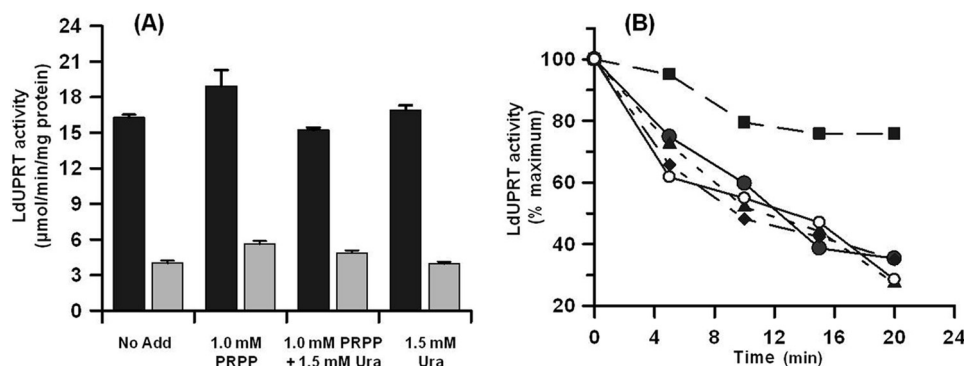


FIGURE 8. **Reversibility and thermostability of uracil-induced substrate inhibition of LdUPRT.** Purified LdUPRT was preincubated in the absence or presence of either 1.0 mM PRPP, both 1.0 mM PRPP and 1.5 mM uracil, or 1.5 mM uracil alone, as described under "Experimental Procedures," then diluted into TMD buffer, and assayed for LdUPRT activity in the presence of 1.0 mM PRPP and either 75  $\mu$ M (black bars) or 1.5 mM uracil (gray bars), respectively (A). The thermostability of LdUPRT was assessed at 62 °C in TMD buffer in the absence (●) or presence of either 1.0 mM PRPP (■), both 1.0 mM PRPP and 75  $\mu$ M uracil (▲), both 1.0 mM PRPP and 1.5 mM uracil (◆), 75  $\mu$ M uracil alone (data not shown), or 1.5 mM uracil alone (○), and residual activity was measured as described under "Experimental Procedures" (B).

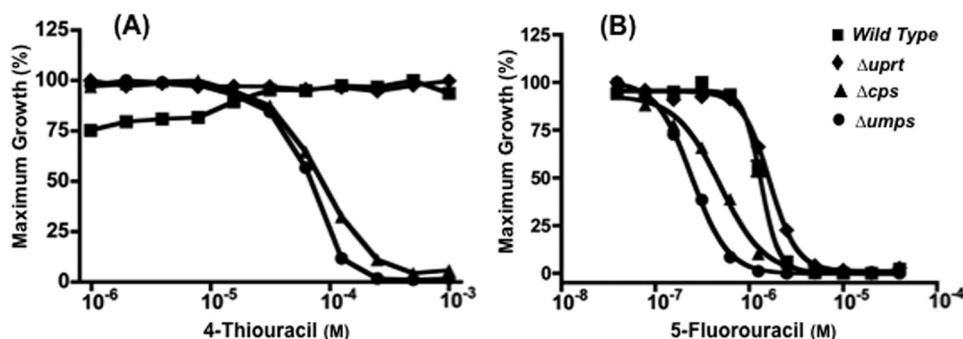


FIGURE 9. **Sensitivity of wild type and null mutant *L. donovani* to 4-thiouracil and 5-fluorouracil.** The abilities of wild type (■),  $\Delta$ uprt (◆),  $\Delta$ cps (▲), and  $\Delta$ umps (●) *L. donovani* promastigotes to grow in various concentrations of 4-thiouracil (A) or 5-fluorouracil (B) were assessed as described under "Experimental Procedures." Growth data are presented from a single representative experiment, which has been repeated at least three times with similar results.

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  $\Delta$ cps *T. gondii* (11),  $\Delta$ cps *T. cruzi* (12), and  $\Delta$ umps *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  $\Delta$ cps 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 other pyrimidines in the parasite, does not impact the uracil supersensitivity of the  $\Delta$ cps or  $\Delta$ umps strain (Fig. 6B). It should be noted that neither orotate nor dihydroorotate, which cannot be converted into UMP by *L. donovani* harboring a  $\Delta$ umps

lesion (Fig. 1), alleviated the susceptibility of the  $\Delta$ umps 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  $\Delta$ cps 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* accommodate 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-



## UPRT from *L. donovani*

transferases: 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  $\Delta cps$  and  $\Delta umps$  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 substrate inhibition by 4-thiouracil versus uracil (Fig. 7, A and C) that was reflected in the greater sensitivity of the  $\Delta cps$  and  $\Delta umps$  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 those that inhibit LdUPRT. Although the precise mechanism of 5-fluorouracil toxicity toward *L. donovani* promastigotes is unknown, the  $\Delta uprt$  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 ( $\Delta uprt$ ) are essentially as sensitive to 5-fluorouracil as wild type parasites ( $EC_{50} = 1.65 \mu M$  versus  $1.33 \mu M$ , 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  $\Delta umps/\Delta uprt$  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 intimate 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-fluorou-

racil 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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