Identifying structural determinants of product specificity in Leishmania major farnesyl diphosphate synthase

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Table S1. List of primers used for generating *Lm*FPPS mutants

Primer	Sequence (5′-3′)
T164F_F	GACGTCGATCTCACTACCTTTATTGGTCAGCTGTACGAC
T164F_R	GTCGTACAGCTGACCAATAAAGGTAGTGAGATCGACGTC
T164Y_F	GACGTCGATCTCACTACCTATATTGGTCAGCTGTACGAC
T164Y_R	GTCGTACAGCTGACCAATATAGGTAGTGAGATCGACGTC
T164W_F	CGTCGATCTCACTACCTGGATTGGTCAGCTGTAC
T164W_R	GTACAGCTGACCAATCCAGGTAGTGAGATCGACG
E97F_F	GGCCCACTTCCTTGTGTTTGACGACATCATGGACC
E97F_R	GGTCCATGATGTCGTCAAACACAAGGAAGTGGGCC
E97Y_F	GGCCCACTTCCTTGTGTATGACGACATCATGGACC
E97Y_R	GGTCCATGATGTCGTCATACACAAGGAAGTGGGCC
E97W_F	CCCACTTCCTTGTGTGGGACGACATCATGG
E97W_R	CCATGATGTCGTCCCACACAAGGAAGTGGG

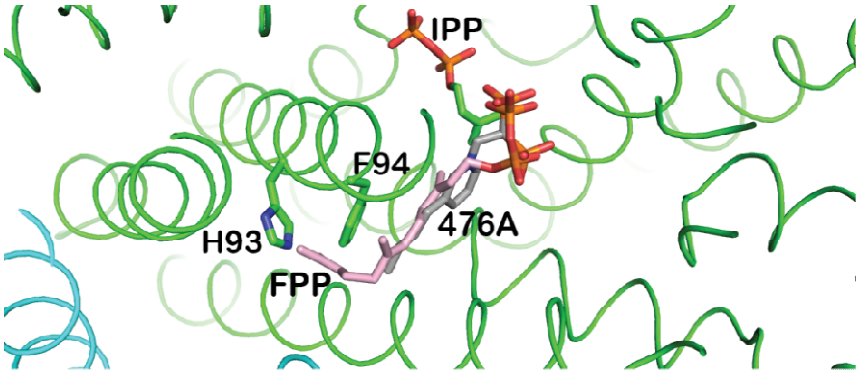

Figure S1. Active site of WT-*Lm*FPPS (PDB ID: 4JZX) in complex with bisphosphonate inhibitor 476A (grey chain; colored by atoms), substrate IPP (colored by atoms). FPP is modeled from avian FPPS structure (PDB ID: 1UBX). The amino acid residues, H93 and F94, are shown as sticks.

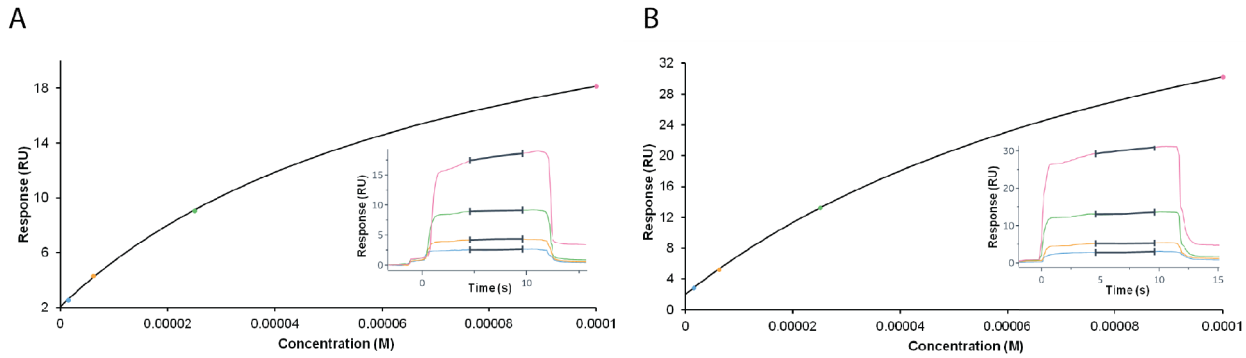

Figure S2. Binding of FPP to T164Y-*Lm*FPPS (A) and T164W-*Lm*FPPS (B) determined by SPR. Steady-state dose response curves are shown against concentration of FPP-1.5, 6.2, 25 and 100 μM. Sensorgrams for multi-cycle kinetics are depicted as inserts.