Correlations of Inhibitor Kinetics for *Pneumocystis jirovecii* and Human Dihydrofolate Reductase with Structural Data for Human Active Site Mutant Enzyme Complexes

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Supplemental Information: Figures S1 and S2.

Figure S1. Illustration of the dose-response curve for PY957 with the N64F mutant protein of hDHFR.

Figure S2. Graphical illustration of the effect of hDHFR active site mutations on K_i values for the inhibitor PY957.