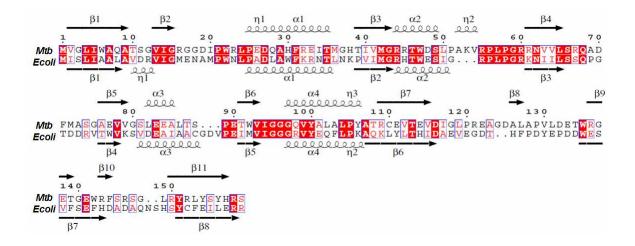
## Kinetic and Chemical Mechanism of the Dihydrofolate Reductase from *Mycobacterium*tuberculosis

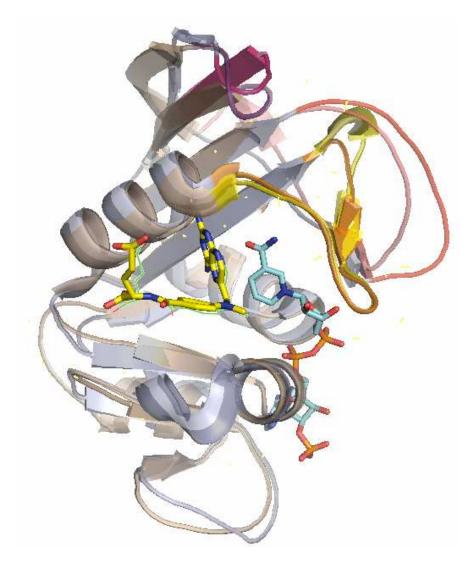
Clarissa M. Czekster<sup>‡</sup>, An Vandemeulebroucke<sup>‡</sup>, John S. Blanchard<sup>‡</sup>\*

<sup>‡</sup> Department of Biochemistry, Albert Einstein College of Medicine, 1300 Morris Park Avenue, Bronx, New York 10461.

Supplementary Figure 1: Primary sequence alignment of the MtDHFR, EcDHFR, and HsDHFR.



Supplementary Figure 2: Structure-based sequence alignment of the MtDHFR and EcDHFR made using ESPript.



Supplementary Figure 3: Structure alignment of the *Mt*DHFR (blue), and the *Ec*DHFR (light orange). The main differences between the two proteins are in the M20 loop (orange in the *Mt*DHFR, yellow in the *Ec*DHFR), the F/G loop (pink in the MtDHFR, light pink in the *Ec*DHFR), and the G/H loop (light purple in the MtDHFR, dark purple in the *Ec*DHFR). NADPH is shown in blue/orange, methotrexate is shown in yellow.