

Theoretical study on ternary complex stability and Michael addition reactivity of Thymidylate synthase/mTHF/XdUMP

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Thymidylate synthase (TS) catalyzes the reductive methylation of dUMP to dTMP during the DNA synthesis process. Herein, binding affinity, and activation energy of Michael addition and a covalent complex formation of TS/mTHF/XdUMP (where X is -H, -F, -Cl or -Br) were investigated using molecular dynamics (MD) and high level quantum mechanics-molecular mechanics (QM-MM) methods. Note that FdUMP is metabolite form of available anticancer agent, 5-FU, targeted at this enzyme. In MD results, the unique H-bonding between Y94 and the substituted fluorine of FdUMP was detected whereas the other systems share almost similar pattern in the overall H-bonding interactions. The B3LYP/6-31+G*-CHARMM potential energy surface according to the two reacting distances, d1: S-(C14)-C6(dUMP) and d2: C5(dUMP)-CH₂(mTHF) suggested that the Michael addition and covalent complex formation occurred in the concerted mechanism. In addition, the SCS-MP2/cc-pVTZ-CHARMM//B3LYP/6-31+G*-CHARMM barriers of this mechanism were 19.7, 23.6 and 25.1 kcal/mol for dUMP, FdUMP and CldUMP complexes.

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