Supplementary Information

Structural basis for drug resistance conferred by β -tubulin mutations: a molecular modeling study on native and mutated tubulin complexes with epothilone B

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Figure S1. Conformational reorganization of the M loop region of β -tubulin upon epothilones' binding, as revealed by the formation of the intra-protein electrostatic interaction between Arg276 and Asp224 residues in complexed β -tubulin.