

Supporting Information for

Crystallographic Structures of Two Bisphosphonate:1-Deoxyxylulose-5-Phosphate Reductoisomerase Complexes

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Figure S1. Stereo representation of the overall structure of the complex between **2** and DXR (dimer structure), PDB file 1T1R. The NADP domain is shown in red, the catalytic domain in yellow, the active site loop in purple and the C-terminal domain in green. The inhibitor **2** is shown in cyan.

