Supplementary Information

Structural basis of methotrexate and pemetrexed action on serine hydroxymethyltransferases revealed using plant models

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Supplementary figure S1

Determination of glycine and 5-formyl-THF (CHO-THF) binding parameters. Panel A-B show *At*SHMT2 with variable 5-formyl-THF; C-D, *At*SHMT2 with variable glycine; E-F, *At*SHMT4 with variable 5-formyl-THF; G-H *At*SHMT4 with variable glycine. Assays were performed at 30 °C, in 20 mM KPi buffer at pH 8.0 for *At*SHM2 and pH 7.3 for *At*SHMT4. Absorbance was measured at 505 nm (*At*SHMT2) and 500 nm (*At*SHMT4) from the quinonoid intermediate formed upon addition of 5-formyl-THF to samples conteining protein (5 mM) and glycine. Samples with varied 5-formyl-THF (glycine at different fixed concentrations) and samples with varied glycine (5-formyl-THF at different fixed concentrations) were measured at equilibrium. The obtained saturation curves were fitted to obtained K_d and αK_d values as explained in Tramonti *et al.*, 2018.



Supplementary figure S2

Methotrexate and pemetrexed inhibition experiments with *At*SHMT2 and *At*SHMT4. Panel A shows data for *At*SHMT2 inhibition by methotrexate; B, *At*SHMT4 by methotrexate; C, *At*SHMT2 by pemetrexed; D, *At*SHMT4 by pemetrexed. Absorbance at 505 nm (*At*SHMT2) or 500 nm (*At*SHMT4) from the quinonoid intermediate formed upon addition of 5-formyl-THF to samples containing enzyme (5 mM), glycine (10 mM), and the antifolate was measured at equilibrium. Right-side charts show double reciprocal plots of data shown in left-side panels fitted to a linear equation.