

# Correction to Probing the Reaction Mechanism of Spore Photoproduct Lyase (SPL) via Diastereoselectively Labeled Dinucleotide SP TpT Substrates

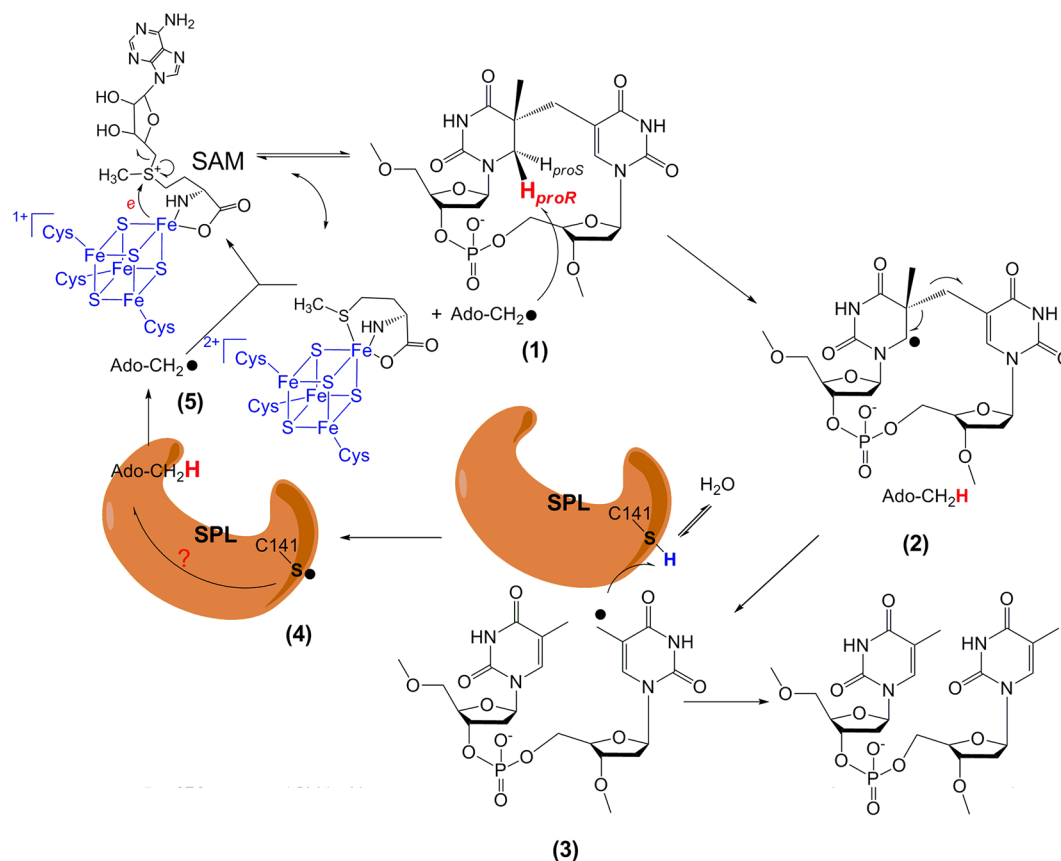
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Page 10443: After carefully calibrating the 5'-dA and SP repair products generated from the wild-type spore photoproduct lyase (SPL) reaction, it was found that 3 equiv of SAM support 5.4 turnovers upon a 3 h enzyme reaction, not the 12 turnovers reported in Figure 9A. The reported 5'-dA consumption in Figure 9B was due to an unknown contaminant enzyme but not the SPL activity to regenerate the S-adenosylmethionine. The corrected description of the reactivity of the wild-type SPL enzyme can be

found in the following paper: Yang, L.; Lin, G.; Nelson, R. S.; Jian, Y.; Telser, J.; Li, L. *Biochemistry*, **2012**, 51 (36), 7173–7188.

Page 10445: We are indebted to the cordial suggestion from Professor John-Stephen Taylor at Washington University, St. Louis, MO, that the stereoconfiguration of SP shown in Figure 10 is not correct. The corrected Figure 10 is shown.



**Figure 10.** Newly proposed reaction mechanism for SPL-catalyzed SP dimer repair. The 5'-dA radical generated from SAM reductive cleavage reaction takes the 6-H<sub>proR</sub> atom to yield a C6 radical on SP, and the SP methylene bridge subsequently undergoes a homolytic cleavage to give a thymine methyl radical. (Note: This allyl radical is likely to delocalize to the thymine aromatic ring; the current drawing as a methyl radical is a simplified model to facilitate discussion.) This radical abstracts an H atom back from an unknown protein residue, presumably C141, to generate a thiyl radical, releasing the repaired TpT. This thiyl radical either takes a H-atom back from the 5'-dA directly or reacts with other protein residues before the 5'-dA is reoxidized to the radical form again. The resulting 5'-dA radical recombines with methionine to regenerate SAM and finish one catalytic cycle.

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