

Supplementary Data

Mechanistic Studies of the Bypass of a Bulky Single-Base Lesion Catalyzed by a Y-Family DNA Polymerase

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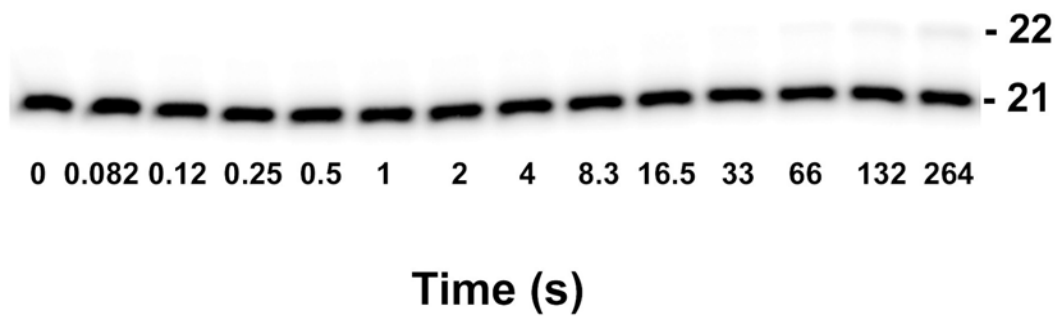
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Supplementary Table 1. Kinetic parameters of dNTP incorporation into undamaged DNA

	K_d, dNTP (μM)	k_p (s^{-1})	$k_p/K_d, \text{dNTP}$ ($\mu\text{M}^{-1}\text{s}^{-1}$)	Fidelity ^a
19/26-mer				
dGTP	183 ± 54	4.6 ± 0.4	2.5 x 10 ⁻²	-
dATP	731 ± 175	(1.0 ± 0.1) x 10 ⁻²	1.4 x 10 ⁻⁵	5.6 x 10 ⁻⁴
dCTP	350 ± 107	(5.0 ± 0.5) x 10 ⁻²	1.4 x 10 ⁻⁴	5.6 x 10 ⁻³
dTTP	1440 ± 305	(1.3 ± 0.2) x 10 ⁻²	8.9 x 10 ⁻⁶	3.6 x 10 ⁻⁴
20/26-mer				
dCTP	205 ± 64	11.6 ± 0.9	5.7 x 10 ⁻²	-
dATP	631 ± 136	(1.2 ± 0.1) x 10 ⁻²	1.9 x 10 ⁻⁵	3.3 x 10 ⁻⁴
dGTP	77 ± 16	(3.5 ± 0.2) x 10 ⁻³	4.5 x 10 ⁻⁵	7.9 x 10 ⁻⁴
dTTP	489 ± 52	(4.0 ± 0.2) x 10 ⁻²	8.3 x 10 ⁻⁵	1.5 x 10 ⁻³
21/26-mer				
dGTP	437 ± 19	1.62 ± 0.03	3.7 x 10 ⁻³	-
dATP	859 ± 168	(2.0 ± 0.2) x 10 ⁻³	2.4 x 10 ⁻⁶	6.5 x 10 ⁻⁴
dCTP	701 ± 30	(2.14 ± 0.04) x 10 ⁻²	3.1 x 10 ⁻⁵	8.3 x 10 ⁻³
dTTP	1180 ± 144	(2.1 ± 0.2) x 10 ⁻³	1.8 x 10 ⁻⁶	4.9 x 10 ⁻⁴
22/26-mer				
dCTP	129 ± 19	4.4 ± 0.2	3.4 x 10 ⁻²	-
dATP	1313 ± 154	(5.5 ± 0.4) x 10 ⁻³	4.2 x 10 ⁻⁶	1.2 x 10 ⁻⁴
dGTP	567 ± 95	(4.8 ± 0.4) x 10 ⁻³	8.5 x 10 ⁻⁶	2.5 x 10 ⁻⁴
dTTP	1340 ± 454	(1.5 ± 0.3) x 10 ⁻²	1.1 x 10 ⁻⁵	3.2 x 10 ⁻⁴
23/26-mer				
dGTP	116 ± 24	2.8 ± 0.1	2.4 x 10 ⁻²	-
dATP	431 ± 25	(6.6 ± 0.1) x 10 ⁻³	1.5 x 10 ⁻⁵	6.2 x 10 ⁻⁴
dCTP	918 ± 102	(1.5 ± 0.1) x 10 ⁻³	1.6 x 10 ⁻⁶	6.7 x 10 ⁻⁵
dTTP	1220 ± 59	(3.8 ± 0.1) x 10 ⁻³	3.1 x 10 ⁻⁶	1.3 x 10 ⁻⁴

^aCalculated as $(k_p/K_d, \text{dNTP})_{\text{incorrect}} / [(k_p/K_d, \text{dNTP})_{\text{correct}} + (k_p/K_d, \text{dNTP})_{\text{incorrect}}]$.



Supplementary Fig. 1. Effectiveness of the DNA trap for biphasic kinetic assays. A preincubated solution of Dpo4 (120 nM), 5'-[³²P]-labeled 21/26-mer-dG^{AP} (30 nM) and DNA trap (5 μM, 21/41-mer D-1) was rapidly mixed with dGTP (1.2 mM) and quenched at various time intervals with 0.37 M EDTA. The products were resolved on a 17% polyacrylamide gel with 8 M urea. The autoradiographed gel image revealed minimal product formation (22-mer) after 264 s. Thus, a molar ratio of 167:1 for the D-1 DNA trap to the radiolabeled damaged DNA substrate was effective at sequestering free Dpo4 that dissociated from the 5'-[³²P]-labeled 21/26-mer-dG^{AP}.