

Supplementary Table 1 | Catalytic efficiency of dC and dT incorporation opposite dG by WT and mutant Pol ν

DNA / Pol ν	Correct (G – dCTP)			Incorrect (G – dTTP)			Fidelity
	K_M (μ M)	k_{cat} (min^{-1})	k_{cat} / K_M ($\text{M}^{-1} \text{min}^{-1}$)	K_M (μ M)	k_{cat} (min^{-1})	k_{cat} / K_M ($\text{M}^{-1} \text{min}^{-1}$)	dC/dT selectivity
13/19mer-G / WT	149 \pm 6	62.4 \pm 1.2	0.42 x 10 ⁶	627 \pm 82	14.4 \pm 1.0	2.3 x 10 ⁴	18.3
Coldspot / WT	16.7 \pm 0.8	54.8 \pm 0.8	3.3 x 10 ⁶	294 \pm 17	5.8 \pm 0.1	2.0 x 10 ⁴	165
Hotspot / WT	7.9 \pm 0.9	10.4 \pm 0.3	1.3 x 10 ⁶	86 \pm 6	14.5 \pm 0.3	16.9 x 10 ⁴	7.7
Coldspot / K679A	9.7 \pm 0.3	103 \pm 2	10.6 x 10 ⁶	386 \pm 87	2.3 \pm 0.1	6.0 x 10 ³	1782
Hotspot / K679A	8.8 \pm 0.4	15.1 \pm 0.2	1.7 x 10 ⁶	422 \pm 47	11.2 \pm 0.6	2.7 x 10 ⁴	63.0
Hotspot / E675R	3.7 \pm 0.3	1.4 \pm 0.1	0.38 x 10 ⁶	35 \pm 5	2.7 \pm 0.1	7.7 x 10 ⁴	4.9
Hotspot / Δ Ins2	21.4 \pm 0.7	13.2 \pm 0.1	0.62 x 10 ⁶	240 \pm 7	14.4 \pm 0.2	6.0 x 10 ⁴	10.3

13/19mer-G	5' GAGTCAACAGGTC 3' CTCAGTTGTCCAG <u>G</u> TATGG
G(-66) coldspot	5' TGAGTGAGCTAACT 3' ACTCACTCGATTGAG <u>T</u> GTAAATTAA
G(164) hotspot	5' GCGATCGGTGCGGGC 3' CGCTAGCCACGCCCG <u>G</u> GAGAAGCGA

Supplementary Table 2 | Catalytic efficiency of WT and Δ Ins2 Pol ν with looped out primers

	K_m (μ M)	k_{cat} (min^{-1})	k_{cat}/K_m ($\text{M}^{-1} \text{min}^{-1}$)	Relative efficiency (%)	
<u>Normal DNA</u>				to WT & normal DNA	
WT	245 \pm 20	16.2 \pm 0.50	66.1 x 10 ³	100	
Δ Ins2	789 \pm 47	17.4 \pm 0.57	22.1 x 10 ³	33.3	
<u>P[3,1] DNA</u>				to WT & P[3,1]	
WT	251 \pm 12	7.0 \pm 0.1	27.9 x 10 ³	42.2	100
Δ Ins2	459 \pm 41	2.3 \pm 0.1	5.0 x 10 ³	7.6	17.9
<u>P[3,2] DNA</u>				to WT / P[3,2]	
WT	369 \pm 22	2.9 \pm 0.1	7.9 x 10 ³	12.0	100
Δ Ins2	310 \pm 53	0.4 \pm 0.02	1.3 x 10 ³	2.0	16.4%