

Supplementary Data

Binding and Cleavage Specificities Define the Ordered Processing of Human Okazaki Fragments by Dna2 and FEN1

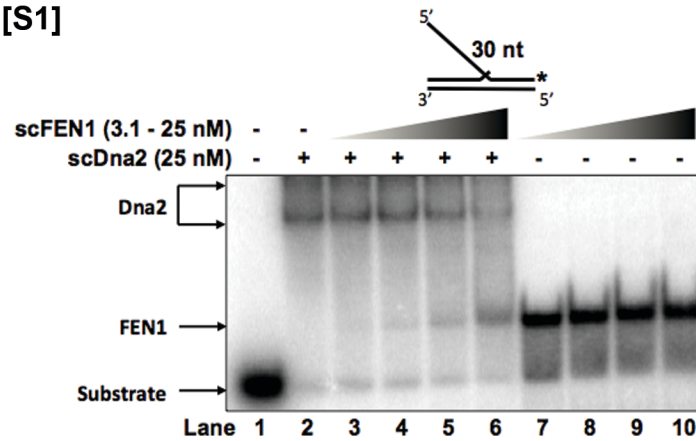
Jason W. Gloor¹, Lata Balakrishnan¹, Judith L. Campbell² and Robert A. Bambara^{1,*}

¹Department of Microbiology and Immunology, University of Rochester School of Medicine and Dentistry, Rochester, New York, 14642.

²Braun Laboratories, California Institute of Technology, Pasadena, California, 91125.

* To whom correspondence should be addressed. Tel: 585-275-3269; Fax: 585-275-6007; Email: robert_bambara@urmc.rochester.edu.

[S1]



Supplementary Figure S1. Yeast FEN1 (scFEN1) displaces Dna2 (scDna2) from long double flap structures. scDna2 (25 nM) was pre-incubated with the experimental 30 nt double flap substrate (U2:T3:D2.30) prior to the addition of increasing concentrations of scFEN1 (3.1, 6.25, 12.5, or 25 nM in lanes 3-6). Lane 1 shows the substrate alone, lane 2 shows scDna2 bound without scFEN1 and lanes 7-10 show scFEN1 alone bound to the substrate at the same concentrations as in 3-6. The position of the substrate alone, scDna2-substrate complex, and scFEN1-substrate complex are indicated to the left of the figure.