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*Correspondence:

Roger Woodgate woodgate@nih.gov

§These authors share first authorship

[†]ORCID:

Alexandra Vaisman orcid.org/0000-0002-2521-1467 John P. McDonald orcid.org/0000-0003-2482-148X Mallory R. Smith orcid.org/0000-0003-1450-7825 Sender L. Aspelund orcid.org/0000-0003-0726-4028 Thomas C. Evans Jr. orcid.org/0000-0001-5406-0146 Roger Woodgate orcid.org/0000-0002-2521-1467

[‡]Present address:

Sender L. Aspelund, Novavax, Inc., Gaithersburg, MD, United States

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Corrigendum: Identification and Characterization of Thermostable Y-Family DNA Polymerases η, ι, κ and Rev1 From a Lower Eukaryote, *Thermomyces lanuginosus*

Alexandra Vaisman^{1†§}, John P. McDonald^{1†§}, Mallory R. Smith^{1†}, Sender L. Aspelund^{1†‡}, Thomas C. Evans Jr^{2†} and Roger Woodgate^{1*†}

¹Laboratory of Genomic Integrity, National Institute of Child Health and Human Development, National Institutes of Health, 9800 Medical Center Drive, Bethesda, MD, United States, ²New England Biolabs Incorporated, Ipswich, MA, United States

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In the original article, there was a formatting issue in **Figure 6** as published. This occurred when the image was converted from a PC generated pdf to an Apple Macintosh generated tif for publication. The corrected **Figure 6** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 6 | TLS past *trans-S*-BPDE-dA by *T. lanuginosus* pols. The ability to bypass BPDE-dA was assayed for (A) pol₁ in the presence of 4 mM Mg²⁺, (B) pol_k in the presence of 4 mM Mg²⁺, (C) pol₁ in the presence of 4 mM Mn²⁺, (D) pol_k in the presence of 4 mM Mn²⁺, (E) pol_k in the presence of 4 mM Mn²⁺, (C) pol₁ in the presence of 4 mM Mn²⁺, (D) pol_k in the presence of 4 mM Mn²⁺, (E) pol_k in the presence of 4 mM Mn²⁺, (C) pol₁ in the presence of 4 mM Mn²⁺, (C) pol₁ in the presence of 4 mM Mn²⁺, (C) pol_k in the presence of 4 mM Mn²⁺, (C) pol₁ in the presence of 4 mM Mn²⁺. The substrate used in these assays was made by annealing of the ³²P labeled primer 5'-CACTGCAGACTCTAAA-3' and either an undamaged or BPDE-containing template 5'- GCTCGTCAGCAGACTCTGCAGTG-3', where the underlined bold A stands for the undamaged, or BPDE modified dA. Reactions contained 100 µM each of individual nucleotide (dC, dG, dA, and dT) or a mixture of all four dNTPs as indicated in the figure and were carried out at 37°C for 10 min. Concentrations of enzymes were 0.17 pM for pol₁, 0.32 pM for pol_k, and 0.15 pM for pol₁. The sequence of the template immediately downstream of the primer (pr) is shown on the left-hand side of each gel pair. The star (*) indicates the position of the adduct.