CORRECTIONS

Telomeric Protein–DNA Point Contacts Identified by Photo-Cross-Linking Using 5-Bromodeoxyuridine, by Brian J. Hicke, Michael C. Willis, Tad H. Koch, and Thomas R. Cech*, Volume 33, Number 11, March 22, 1994, pages 3364–3373.

Page 3367. In Figure 3, the line art is incorrectly positioned with respect to the gray scale. Also, in part B, the bands in the upper left portion are obscured. The figure should appear as follows:

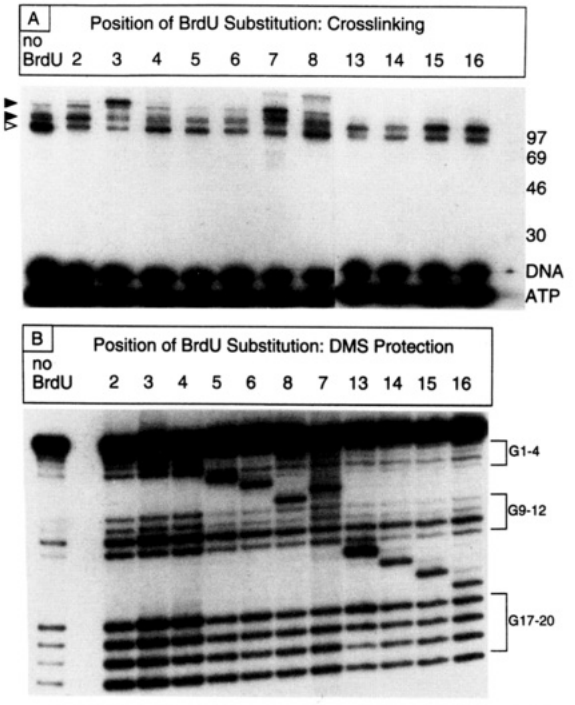

FIGURE 3: Methylation protection footprint on BrdU-substituted DNAs. In each DNA, BrdU was substituted by automated DNA synthesis for one nucleotide in O2T. Position 1 is the 3′ G. (A) Irradiation of BrdU-substituted nucleoprotein complexes, followed by SDS–PAGE/autoradiography. Open arrowhead: cross-link to the β subunit. Filled arrowheads: cross-links to α. (B) BrdU-substituted nucleoprotein complexes analyzed by methylation protection as described (Fang et al., 1993). Not shown is the methylation of protein-free DNA, which produces a uniform ladder of cleavage at Gs; cf. Gray et al. (1991).

Murine Heparin Cofactor II: Purification, cDNA Sequence, Expression, and Gene Structure, by Guang Sen Zhang, Julie H. Mehringer, Vivianna M. D. Van Deerlin, Christine A. Kozak, and Douglas M. Tollefsen*, Volume 33, Number 12, March 29, 1994, pages 3632–3642.

Page 3638. In Figure 9, the data in lanes a and b for the bands marked I1 and I2 did not reproduce. The figure should appear as follows:

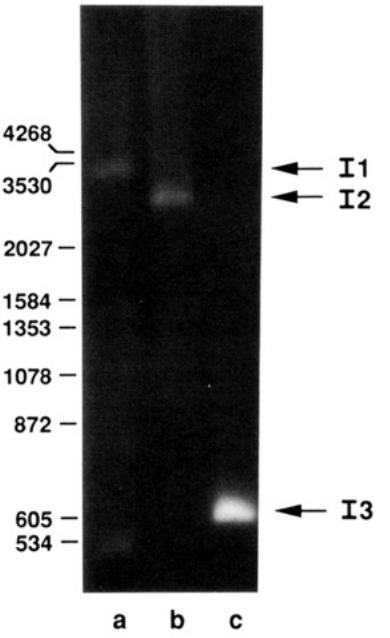

FIGURE 9: PCR analysis of intron size. Intron size was estimated by PCR amplification of genomic clones using primers near the intron/exon boundaries. The amplified product sizes were determined by mobility in agarose gel electrophoresis as compared to standards shown on the left. *Lane a*, intron 1 (I1) amplified from clone G-1.3; *lane b*, intron 2 (I2) amplified from clone G-2.2; *lane c*, intron 3 (I3) amplified from clone G-2.2. Calculated sizes for the PCR products were the following: intron 1, 3300 bp; intron 2, 2900 bp; intron 3, 630 bp. Subtraction of the distance between primers in the cDNA yielded net sizes for the introns as follows: intron 1, 2900 bp; intron 2, 2700 bp; intron 3, 400 bp.

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