

S. Yan, M. Wu, D. J. Patel, N. E. Geacintov, and S. Broyde. 2003. Simulating structural and thermodynamic properties of carcinogen-damaged DNA. *Biophys. J.* 84:2137–2148.

The correct version of Table 5 is as follows:

**TABLE 5 Trajectory-averaged minor groove widths**

	10S (+)- <i>trans</i>	10R (-)- <i>trans</i>	B-DNA
C5–G22	8.4 (1.8)	7.5 (1.6)	7.2 (1.4)
G*6–G21	9.9 (1.4)	8.0 (2.0)	6.8 (1.4)
C7–T20	10.9 (1.0)	9.3 (1.6)	7.7 (1.2)
T8–A19	10.4 (0.6)	9.7 (0.8)	7.2 (1.1)
A9–G18	8.4 (1.3)	8.5 (1.4)	5.2 (1.4)
C10–C17	7.0 (1.4)	8.2 (1.3)	6.9 (1.5)
C11–G16	6.5 (1.8)	6.6 (1.8)	7.3 (1.5)

Minor groove width is calculated from the indicated interstrand phosphate-to-phosphate distance for each structure over the 1.5–3-ns time frame, using MD Toolchest (Ravishanker et al., 2004, *MD Toolchest*, Wesleyan University, Middletown, CT). Standard deviations are given in parentheses. All distances are in Å. Minor groove widths appear to be larger in solution than in crystals (Wang et al., 2002, *Biophys. J.* 83:382–406).

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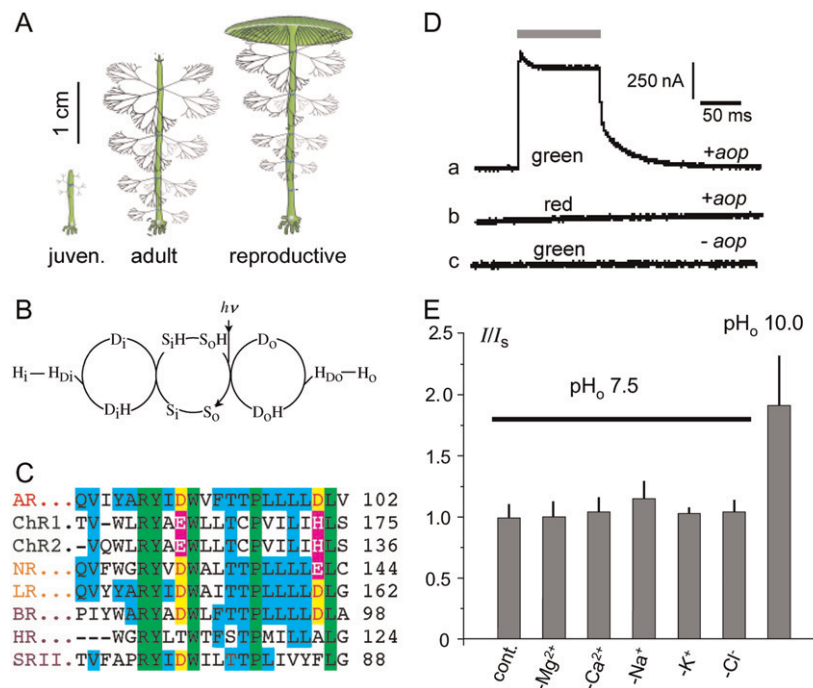
A. P. Liu and D. A. Fletcher. 2006. Actin polymerization serves as a membrane domain switch in model lipid bilayers. *Biophys. J.* 91:4064–4070.

Due to a copy editing error, the lipid composition of the giant unilamellar vesicles described in the Materials and Methods section on page 4065 was incorrectly written as “2%:1%:30%:0.6%” DPPC/DOPC/Chol/BODIPY TMR PIP<sub>2</sub>. The lipid composition should read “2:1:30%:0.6%” DPPC/DOPC/Chol/BODIPY TMR PIP<sub>2</sub>, indicating a 2:1 ratio of DPPC/DOPC with 30% and 0.6% of total lipids being Chol and BODIPY TMR PIP<sub>2</sub>, respectively. The BioFAST version of the article has the correct composition.

doi: 10.1529/biophysj.106.0900151

Satoshi P. Tsunoda, David Ewers, Sabrina Gazzarrini, Anna Moroni, Dietrich Gradmann, and Peter Hegemann. 2006. H<sup>+</sup>-pumping rhodopsin from the marine alga *Acetabularia*. *Biophys. J.* 91:1471–1479.

Due to printer error, Fig. 1 printed incorrectly. Here is the correct image:



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