

Corrections

INAUGURAL ARTICLE, ENVIRONMENTAL SCIENCES. For the article “Simulation of Sahel drought in the 20th and 21st centuries,” by I. M. Held, T. L. Delworth, J. Lu, K. L. Findell, and T. R. Knutson, which appeared in issue 50, December 13, 2005, of *Proc. Natl. Acad. Sci.*

USA (**102**, 17891–17896; first published December 1, 2005; 10.1073/pnas.0509057102), the numerical scales corresponding to the color bars in Figs. 2, 4, and 5 appeared incorrectly, due to a printer’s error. The corrected figures and their legends appear below.

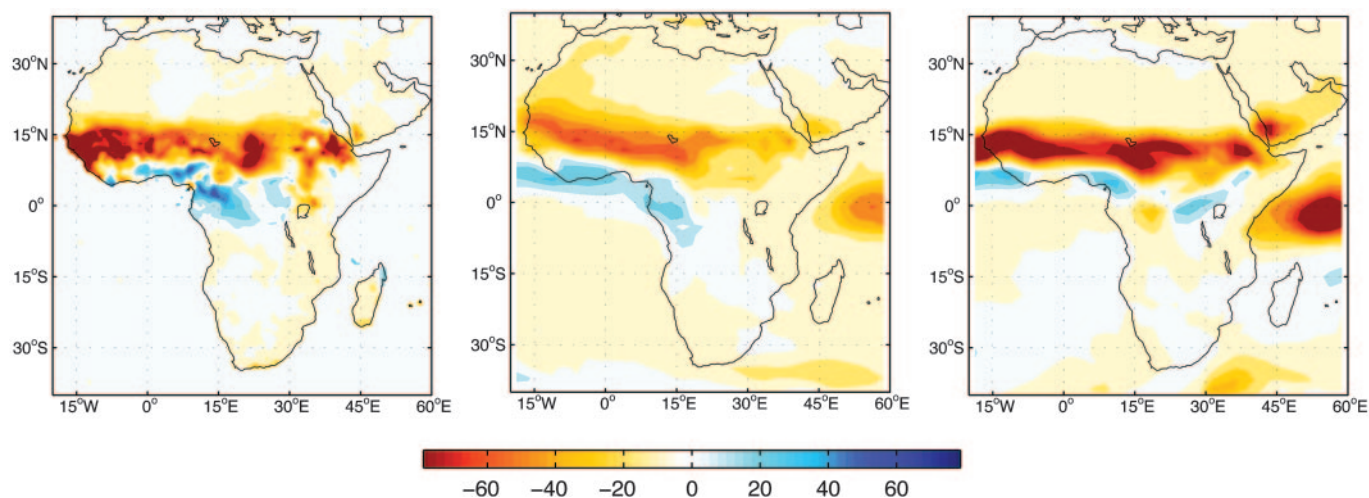


Fig. 2. Observed and modeled rainfall trends. (*Left*) The linear trend from 1950 to 2000 in the observed (CRU) July–August–September rainfall over land, in mm/month per 50 years. Blue areas correspond to a trend toward wetter conditions, and brown areas toward a drier climate. (*Center*) The linear trend for the eight-member ensemble mean of CM2 but plotted over both land and ocean. (*Right*) Linear trend for an ensemble mean of 10 simulations with the atmospheric/land component of CM2.0 running over observed sea surface boundary conditions.

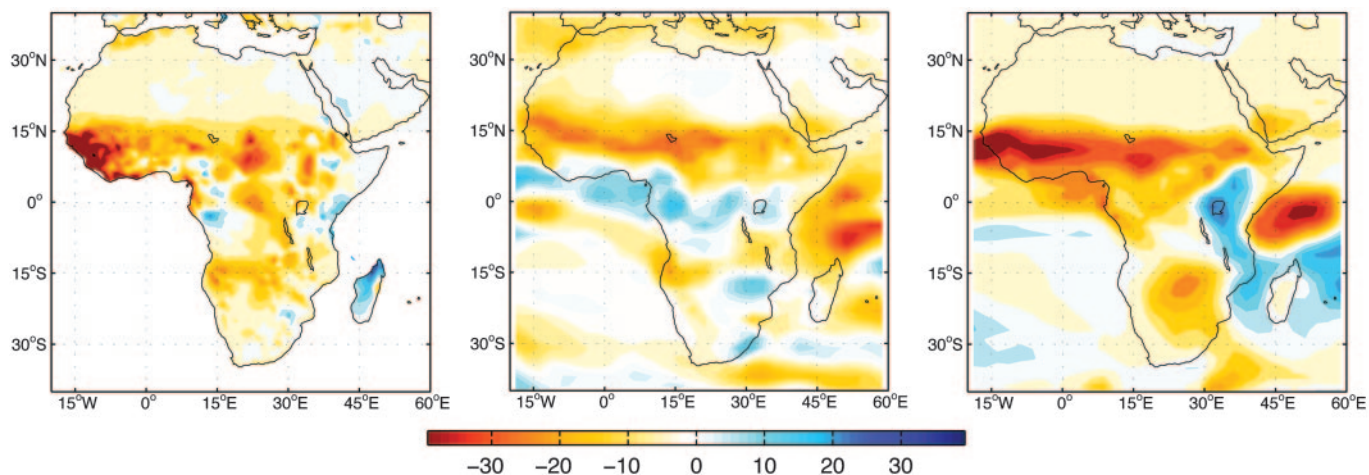


Fig. 4. As in Fig. 2 but for the annual mean precipitation. The CM2 ensemble mean trend in *Center* has been multiplied by 2.

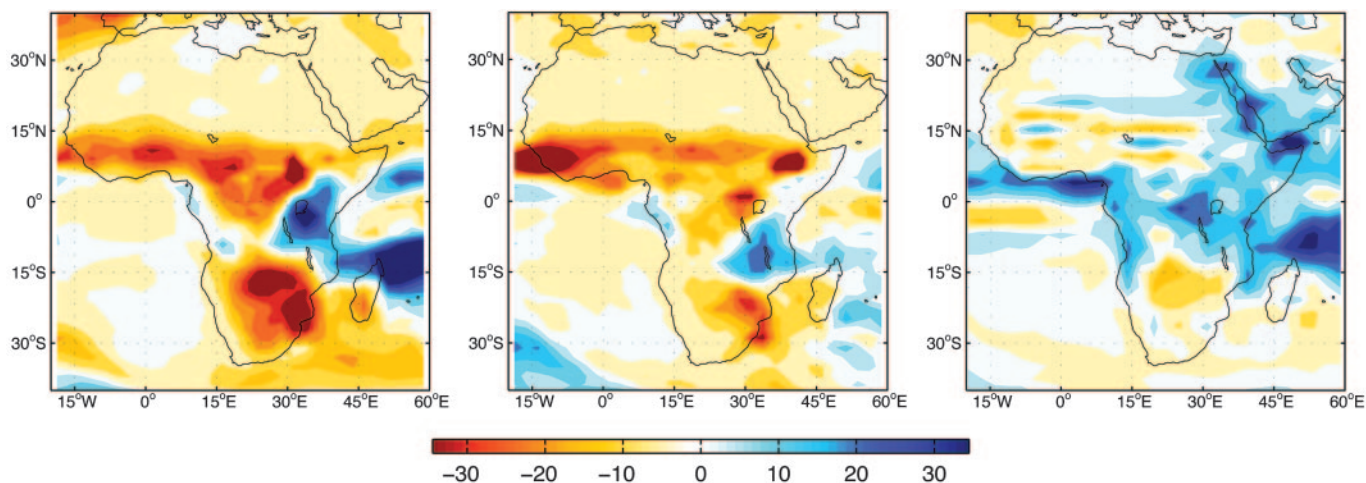


Fig. 5. The annual mean precipitation response of three atmospheric models to a uniform warming of ocean temperatures. (Left) The atmospheric component of CM2.0. (Center) A model developed at National Aeronautics and Space Administration's Global Modeling and Assimilation Office (J. Bacmeister, personal communication). (Right) The CAM3 model developed at the National Center for Atmospheric Research (J. Kiehl, personal communication).

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BIOCHEMISTRY. For the article “Multiple modes of interaction of the deglycosylation enzyme, mouse peptide *N*-glycanase, with the proteasome,” by Guangtao Li, Xiaoke Zhou, Gang Zhao, Hermann Schindelin, and William J. Lennarz, which appeared in issue 44, November 1, 2005, of *Proc. Natl. Acad. Sci. USA* (**102**, 15809–15814; first published October 25, 2005; 10.1073/pnas.0507155102), the authors note that the horizontal labels for His-6-mPNGase and His-6-p97 in Fig. 5 are transposed. The corrected figure and its legend appear below.

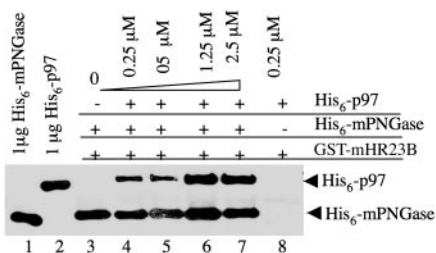


Fig. 5. mHR23B, mPNGase, and p97 form a complex. GSH-agarose beads (8 μ l) containing 0.25 μ M bound GST-mHR23B were incubated with 0.25 μ M His-6-mPNGase in the presence of 0, 1, 2, 5, or 10 molar excess of His-6-p97. The samples were eluted in 45 μ l of SDS-loading buffer and subjected to SDS/PAGE, electrotransferred, and then blotted with anti-His mAb.

www.pnas.org/cgi/doi/10.1073/pnas.0510409102

CELL BIOLOGY. For the article “Abnormal centrosome amplification in cells through the targeting of Ran-binding protein-1 by the human T cell leukemia virus type-1 Tax oncoprotein,” by Jean-Marie Peloponese, Jr., Kerstin Haller, Akiko Miyazato, and Kuan-Teh Jeang, which appeared in issue 52, December 27, 2005, of *Proc. Natl. Acad. Sci. USA* (**102**, 18974–18979; first published December 19, 2005; 10.1073/pnas.0506659103), the authors note that the following statement should be added to the Acknowledgments: “We thank Dr. Patrizia Lavia (National Research Council Center of Evolutionary Genetics, University of Rome) for the generous gift of the RanBP1HA plasmid.”

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