Interaction of the mammalian endosomal sorting complex required for transport (ESCRT) III protein hSnf7-1 with itself, membranes, and the AAA⁺ ATPase SKD1.

Yuan Lin, Lisa A. Kimpler, Teresa V. Naismith, Joshua M. Lauer, and Phyllis I. Hanson

The plasmid used to express FLAG-hSnf7N (residues 1–116) in this paper had an unintended missense mutation changing serine at residue 2 to cysteine. We found that this cysteine was palmitoylated. Changing it back to serine decreased the amount of hSnf7N associated with membranes from all for the mutant fragment containing cysteine to approximately half for the wild-type fragment containing serine. The images of FLAG-hSnf7N in Figs. 7B and 8 represent average cells expressing mutant (Cys-2) hSnf7N, whereas for wild-type (Ser-2) hSnf7N, these images correspond to cells expressing high levels of protein. All other plasmids are as indicated, and the conclusions of the paper remain unchanged.

Role of the N-terminal domain of the human DMC1 protein in octamer formation and DNA binding.

Takashi Kinebuchi, Wataru Kagawa, Hitoshi Kurumizaka, and Shigeyuki Yokoyama

Due to an inadvertent error, the wrong image was presented in Fig. 4C. Fig. 4C should appear as shown below. The figure legend and text remain unchanged.

Dynamic changes in histone H3 lysine 9 methylation.
IDENTIFICATION OF A MITOSIS-SPECIFIC FUNCTION FOR DYNAMIC METHYLATION IN CHROMOSOME CONGRESSION AND SEGREGATION.

Kirk J. McManus, Vincent L. Biron, Ryan Heit, D. Alan Underhill, and Michael J. Hendzel

The concentration of a drug that was employed, adenosine dialdehyde, was erroneously reported as 25 mM. The concentration that was employed was actually 250 μM.

Conditional deletion of hypothalamic Y2 receptors reverts gonadectomy-induced bone loss in adult mice.

Susan J. Allison, Paul Baldock, Amanda Sainsbury, Ronaldo Enriquez, Nicola J. Lee, En-Ju Deborah Lin, Matthias Klugmann, Matthew During, John A. Eisman, Mei Li, Lydia C. Pan, Herbert Herzog, and Edith M. Gardiner

Dr. Klugmann’s name was misspelled in the author line. The correct spelling is shown above.

The first structure from the SOUL/HBP family of heme-binding proteins, murine P22HBP.

Jorge S. Dias, Anjos L. Macedo, Gloria C. Ferreira, Francis C. Peterson, Brian F. Volkman, and Brian J. Goodfellow

Column 2, first line: The concentrations should read "(4.0 μM) or hemin (3.5 μM) . . ."
PTP-PEST couples membrane protrusion and tail retraction via VAV2 and p190RhoGAP.

Sarita K. Sastri, Zenon Rajfur, Betty P. Liu, Jean-Francois Cote, Michel L. Tremblay, and Keith Burridge

In Fig. 5, panel D was inadvertently omitted and is shown below. The figure legend is correct as it appears.

**FIGURE 5**

D

PTP-PEST -/-

Actin

PTP-PEST -/- + VAV2 RS

VAV2

Actin

FIGURE 5
Correction of pulmonary abnormalities in Sftpd$^{-/-}$ mice requires the collagenous domain of surfactant protein D.

Paul S. Kingma, Liqian Zhang, Machiko Ikegami, Kevan Hartshorn, Francis X. McCormack, and Jeffrey A. Whitsett

Fig. 5: An incorrect image was used for the panel A inset. The correct image is shown below.

**FIGURE 5**
Interaction of the mammalian endosomal sorting complex required for transport (ESCRT) III protein hSnf7-1 with itself, membranes, and the AAA⁺ ATPase SKD1. VOLUME 280 (2005) PAGES 12799-12809

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