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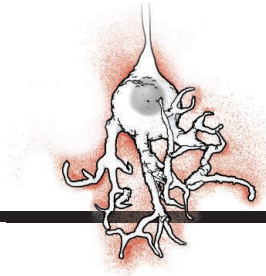

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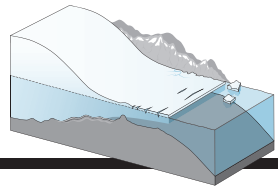

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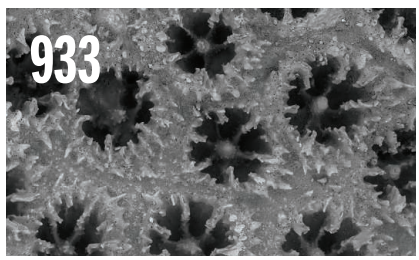

DEPARTMENTS

ON THE COVER

The Lassa virus glycoprotein in its trimeric, viral surface state (orange) bound to a fragment of a neutralizing antibody from a human survivor (white). A second, empty antibody binding site is illuminated. Bound antibody bridges different monomers in the trimer to prevent conformational changes required for infection. The crystal structure determined by Hastie *et al.* provides a template for vaccine development. See page 923.
Illustration: C. Bickel/Science; Coordinates: Kathryn Hastie and Erica Ollmann Saphire, The Scripps Research Institute

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