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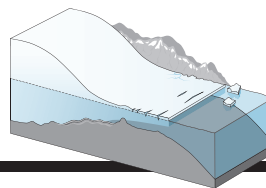
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RESEARCH ARTICLE SUMMARY; FOR FULL TEXT:

[dx.doi.org/10.1126/science.aal3345](https://doi.org/10.1126/science.aal3345)

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[dx.doi.org/10.1126/science.aah6849](https://doi.org/10.1126/science.aah6849)



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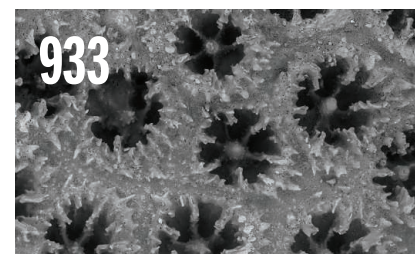
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The Lassa virus glycoprotein in its trimeric, viral surface state (orange) bound to a fragment of a neutralizing antibody from a human survivor (white). A second, empty antibody binding site

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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