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Salk's outlook
The genome of *Methanococcus jannaschii* is the first one of the Archaea domain that has been completed by whole-genome shotgun sequencing at The Institute for Genomic Research. This provides the first opportunity for large-scale comparisons of the genetic components from all principal domains—Eukaryotes, Bacteria, and Archaea. See the article on page 1058 and the News story on page 1043.

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BTK as a Mediator of Radiation-Induced Apoptosis in DT-40 Lymphoma B Cells

Corticofugal Modulation of Time-Domain Processing of Biosonar Information in Bats
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**On the Web**

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