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Experimental electron density map of the P4-P6 domain of the Tetrahymena thermophila group I intron contoured at 1.4 standard deviations above the mean. The 2.8 angstrom crystal structure of the 160-nucleotide domain reveals the close packing of RNA helices likely to be a common feature of ribozymes, the ribosome, and the spliceosome. See pages 1678 and 1696 and the related Perspective on page 1676. [Image: J. H. Cate and J. A. Doudna]

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Small-Conductance, Calcium-Activated Potassium Channels from Mammalian Brain


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