

Data collection and processing

A single wavelength anomalous diffraction experiment was carried out at the CuK α wavelength on crystals cryoprotected in mother liquor plus 25% 2-methyl-2,4-pentanediol; clear diffraction was observed to at least 1.8 Å resolution. We indexed, integrated and scaled the data using D*TREK²⁵, identified heavy atom sites with SOLVE²⁶, and carried out refinement with CNS²⁷ to obtain a model with final R_{xtal} and R_{free} values of 17.8% and 22.8%, respectively. The model contains all RNA atoms except A82, for which electron density was not observed, along with the hypoxanthine ligand. For additional experimental details and references, see Supplementary Information.

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Competing interests statement The authors declare that they have no competing financial interests.

Correspondence and requests for materials should be addressed to R.T.B. (robert.batey@colorado.edu). The atomic coordinates and structure factors have been deposited in the RCSB Protein Data Bank under accession number 1U8D.

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corrigendum

The genome of *Cryptosporidium hominis*

Ping Xu, Giovanni Widmer, Yingping Wang, Luiz S. Ozaki, Joao M. Alves, Myrna G. Serrano, Daniela Puiu, Patricio Manque, Donna Akiyoshi, Aaron J. Mackey, William R. Pearson, Paul H. Dear, Alan T. Bankier, Darrell L. Peterson, Mitchell S. Abrahamsen, Vivek Kapur, Saul Tzipori & Gregory A. Buck

Nature **431**, 1107–1112 (2004).

The GenBank accession number was supplied incorrectly as AAEL000000. The sentence should have read: ‘The sequences reported in this paper have been deposited in GenBank under the project accession number AAEL000000000’. In addition, the received date should have been 14 March 2004. □