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Supplementary Information accompanies the paper on *Nature's* website (♦ <http://www.nature.com>). A stereo representation of figure 3 and the FSC plot to assess the resolution of the reconstruction are provided.

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Competing interests statement The authors declare competing financial interests: details accompany the paper on *Nature's* website (♦ <http://www.nature.com/nature>).

Correspondence and requests for materials should be addressed to M.v.H. (e-mail: m.vanheel@ic.ac.uk). The atomic coordinates of the *E. coli* RF2 structure fitted into the cryo-EM map have been deposited in the Protein Data Bank (e-mail: <http://www.rcsb.org>) with the accession code 1ML5. The cryo-EM map has been deposited in the 3D EM data base (e-mail: http://www.ebi.ac.uk/msd/MSDProjects/IIMS3D_EM.html) with the accession code EMD-1005.

corrigenda

The genome sequence of *Schizosaccharomyces pombe*

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In this Article, the author Andreas Düsterhöft was mistakenly omitted: his name and affiliation (footnote 6) should have been inserted between M. Fuchs and C. Fritzc in the author list. In addition, the name of L. Cerutti (in the last line of the author list) was misspelled. On p874 in the penultimate sentence of the 'Intergene regions' section, "tamendly oriented genes" should read "divergently oriented genes." □

Probing the free-energy surface for protein folding with single-molecule fluorescence spectroscopy

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The upper limit on the polypeptide reconfiguration time (τ_0) was inadvertently calculated using $(\sigma_{app} - \sigma_0)^2$, instead of $(\sigma_{app}^2 - \sigma_0^2)$, as given in the formula in the text (page 745, right column). The correct upper limit is therefore 0.2 ms. This results in a lower limit on the free energy barrier (Δ) of $2k_B T$, corresponding to an activation entropy of $+3k_B$ (page 746, right column), and an upper limit on the pre-exponential factor ($2\pi\tau_0$) of 1 ms. This mistake does not affect any of the conclusions. We thank Taekjip Ha for bringing it to our attention. □