SUPPLEMENTARY FIGURE 1

A

Flow diagram of data processing pipeline followed to give reported structure. All steps used RELION4.0.

B

Local variation of resolution in reported structure

C

Fourier shell correlation plot for final structure
Orthogonal views of superposition of the closed Human RAD17-RFC 9-1-1 and the two published closed confirmations of the Yeast Rad24-RFC 9-1-1 structures with subunits (or equivalents) coloured as for figure 1.
SUPPLEMENTARY FIGURE 3

Montage of features in the experimental map corresponding to adenine nucleotides bound to each subunit of the clamp loader. In all cases volumes suggestive of binding of ATPγS are evident.