



**Supplementary information, Fig. S2 Cryo-EM data processing workflow of human telomerase holoenzyme. a** All datasets and particles used in this paper. **b** Masked 3D classification for the catalytic core. **c** Masked 3D classification for the biogenesis domain. **d** Data processing flowchart for the catalytic core. **e** Data processing flowchart for the catalytic core focusing on the H2A/H2B heterodimer. **f** Data processing flowchart for the biogenesis domain. **g** Data processing flowchart for the biogenesis domain focusing on stems p7-p8a of the hTR RNA.