

Figure 1: Isolation of hDUS2(1-340) – tRNA-Val complex. (a) Mechanism of cross-linking between hDUS2 and 5-BrUrd-modified RNA. (b) Western blot analysis of cross-linking between 5-BrUrd-modified tRNA-Val-CAC. (c) Peptide-oligo adduct of hDUS2-tRNA-Val-CAC complex characterized by bottom-up proteomics. (d) Scheme for isolation of hDUS2-tRNA complex by mechanism-based crosslinking. (e) RNA-protein complex isolation by HiTrap Q anion exchange chromatography. (f) RNA-protein complex isolation by Superdex 200 size exclusion chromatography. (g) SDS-PAGE analysis of purified hDUS2 (1-340) and purified hDUS2-tRNA complex.

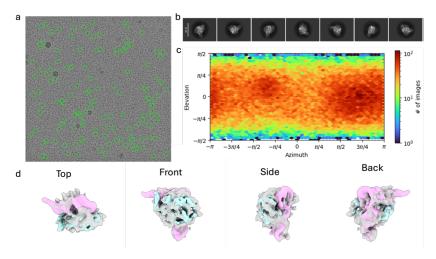


Figure 2: Preliminary Cryo-EM imaging results. **(a)** Representative micrographs (green circle = 136 Å). **(b)** Best 2D classes. **(c)** 3D map orientation distribution. **(d)** Cryo-EM maps docked with PDB 4tna (pink) and 4xp7 (teal).