



*Figure 1 (A) Micrograph (left) and 2D class averages (right) of the STING polymer bound to both cGAMP and one of our newly designed STING agonists - UT009. Long polymers are indicated by red arrows. (B) Chemical structure of UT009. (C) Cryo-EM structure of the STING polymer bound to both cGAMP and UT009 at 2.9 Å resolution.*