



Figure 1. Preliminary Cryo-EM data of Spike protein in complex with neutralizing antibody. (a) Representative micrograph. (b) 2D class averages. (c) Heterogeneous refinement to filter out antibody bound 3D classes, particles from class 2 and 4 combine and refined by non-homogeneous refinement. (d) FSC curve. (e) Local resolution estimation of global map.