A)

B)

S14 Fig. Peak viral concentration and overall posterior viral concentration trajectories in terms of genome equivalents per ml. Posterior peak viral concentration distribution for symptomatic (red) and asymptomatic (blue) individuals (A) and for all individuals combined (B). Undelrying data are available at
https://github.com/gradlab/CtTrajectories/tree/main/output/params_df_split.csv[10] (A) and
https://github.com/gradlab/CtTrajectories/tree/main/output/params_df_combined.csv[10] (B)