

# Supplementary

## Materials

### **Integrating Conformational Dynamics and Perturbation-Based Network Modeling for Mutational Profiling of Binding and Allostery in the SARS-CoV-2 Spike Variant Complexes with Antibodies: Balancing Local and Global Determinants of Mutational Escape Mechanisms**

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Supplemental Materials include scripts for computation of distance fluctuations from MD simulations.