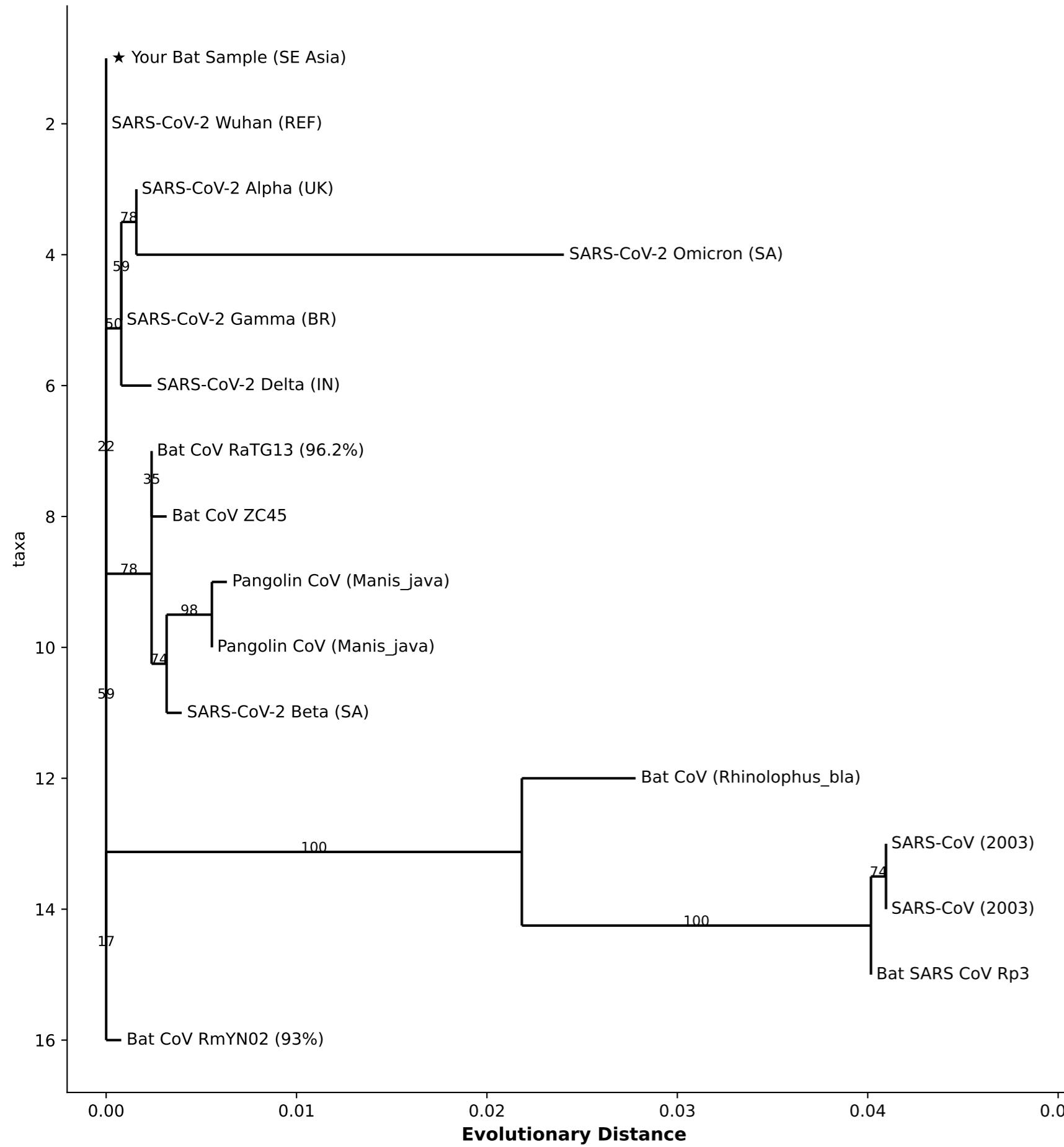


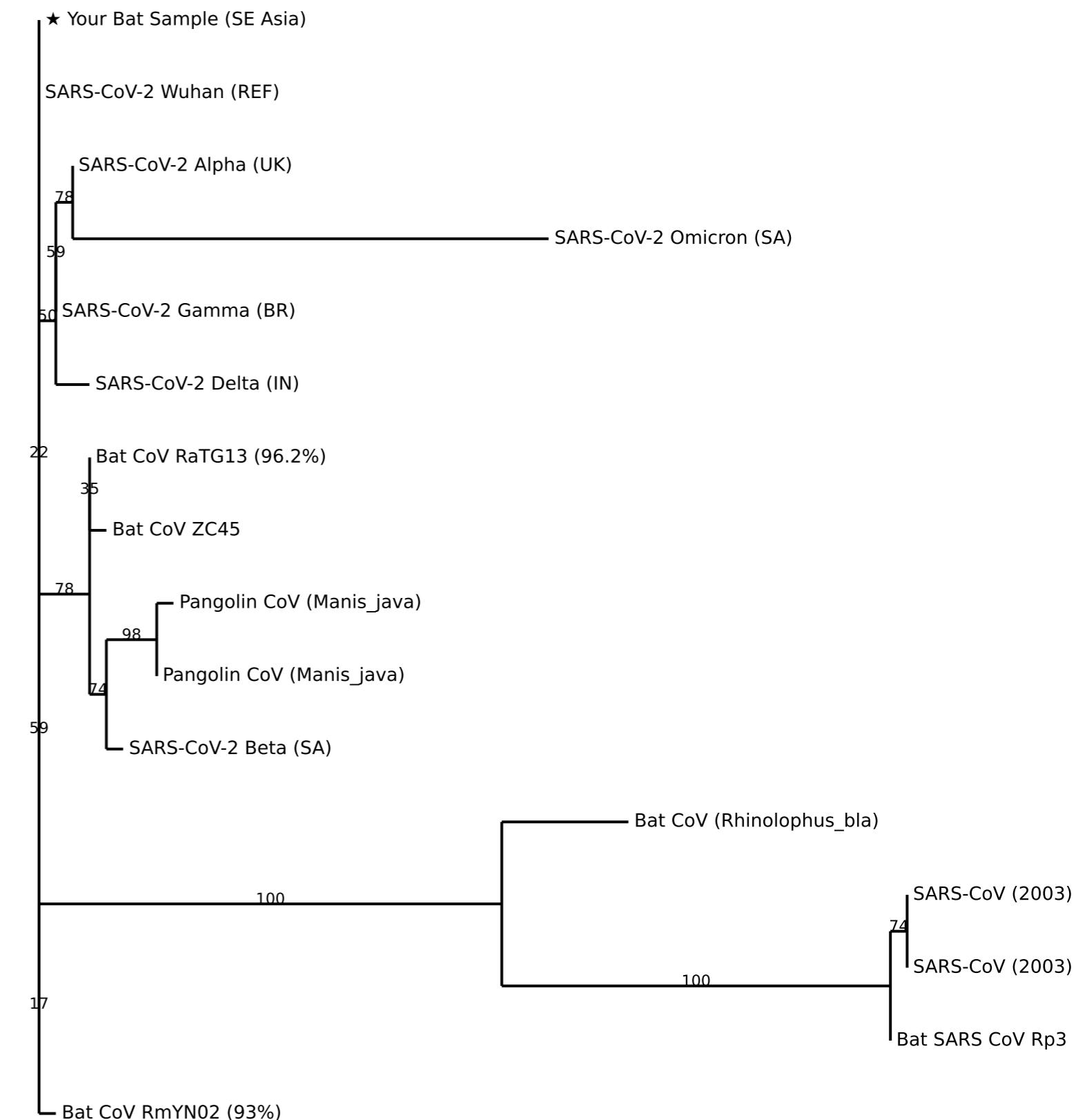
SARS-CoV-2 Evolutionary Analysis

Spike Protein Phylogeny (16 Sequences)

**A. Phylogenetic Tree - Rectangular Layout
16 Coronavirus Spike Proteins**



**B. Phylogenetic Tree - Linear Layout
For Selection Analysis**



Your Sample (n=1) SARS-CoV-2 variants (n=6) Bat coronaviruses (n=5) Pangolin CoVs (n=2) SARS-CoV 2003 (n=2)