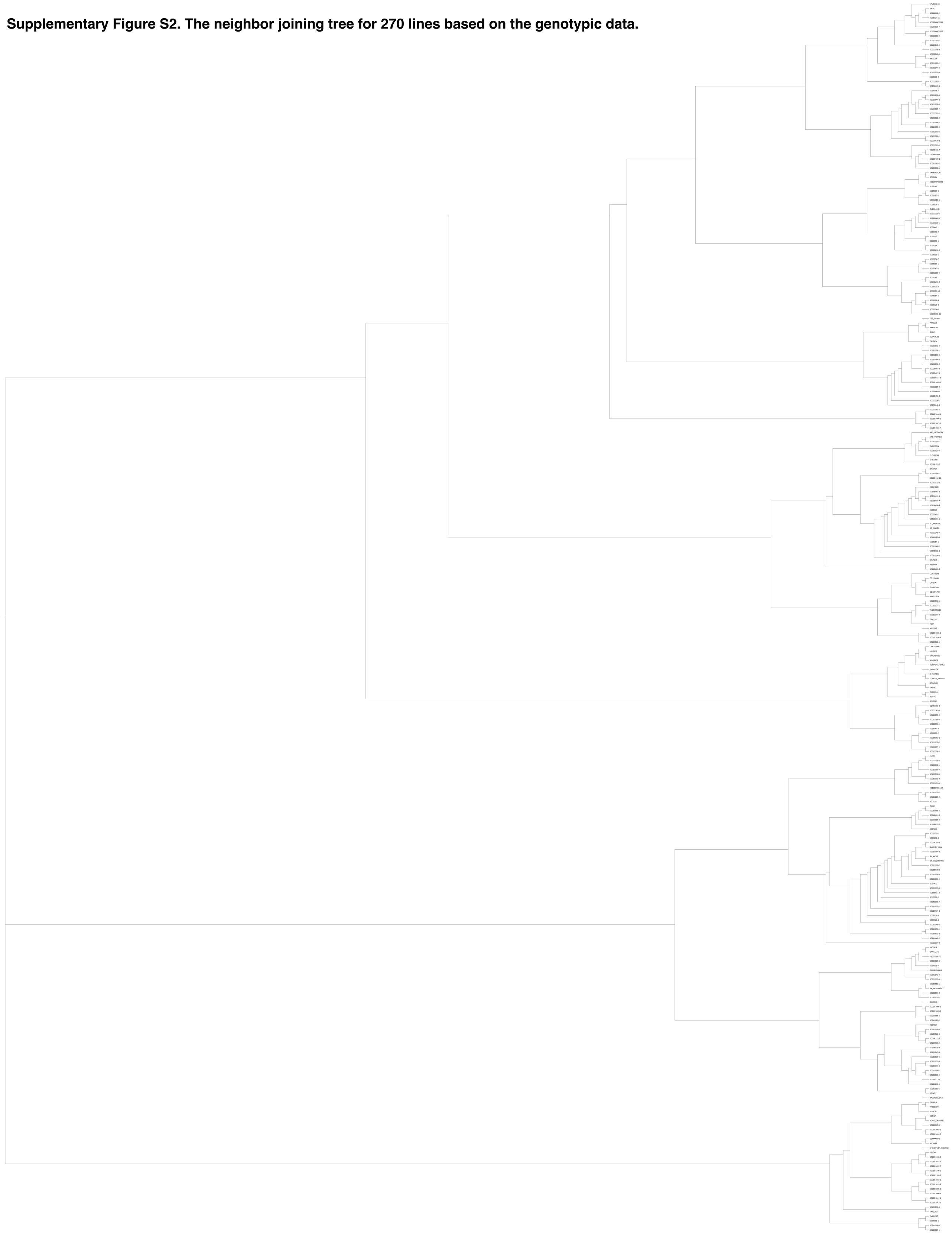Supplementary Figure S2. The neighbor joining tree for 270 lines based on the genotypic data.