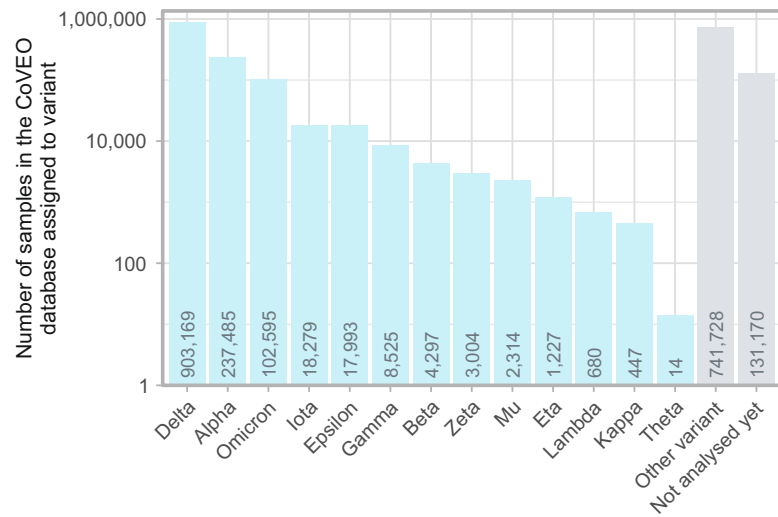**Supplementary Figure 1.** The number of samples in the CoVEO database. **a.** The number of good-quality SARS-CoV-2 samples with a human host assigned to different variants in the CoVEO database. **b.** The relationship between the total number of samples assigned to a specific variant and the number of co-infection samples containing the variant. The straight black line represents a linear dependence with a slope of 1 on a log-log graph. R represents the Pearson-correlation coefficient, and the corresponding p-value is derived from a two-sided t-test (n=13 variants).