



Supplementary Figure S2: The proportion of genetic subgroups of all SARS-CoV-2 viruses analyzed by Next Generation Sequencing (NGS) from Oslo and Viken, i.e. the counties of recruitment, per month during the study recruitment period, among sequences with >70% coverage. All subgroups with less than 5 occurrences are categorized as «Others», while «B» og «B.1» includes virus that were not allocated to a subgroup. Source: Surveillance data from the Norwegian Institute of Public Health (NIPH)