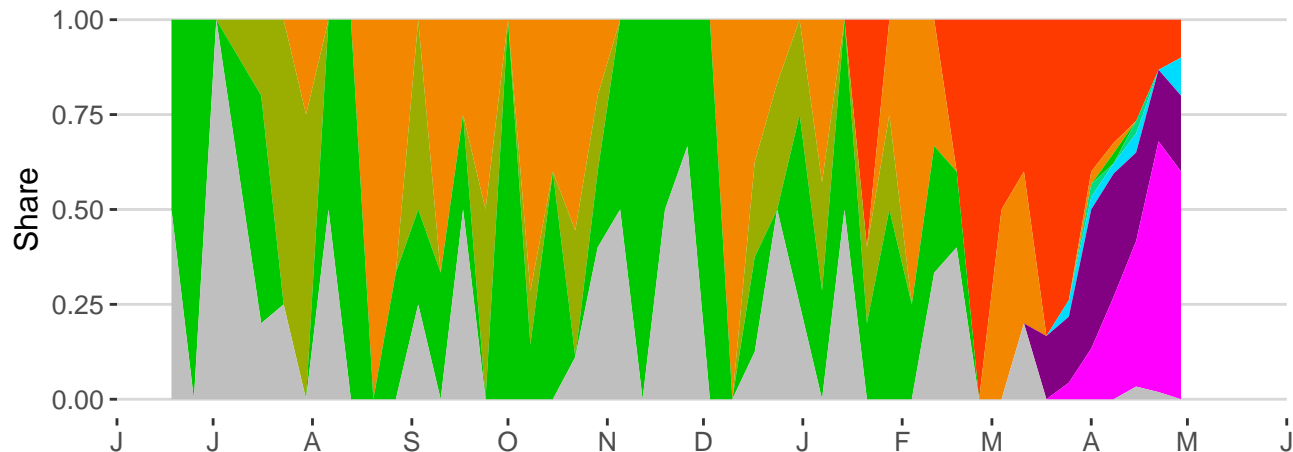


SPREAD OF SARS-CoV2 VARIANTS B.1.617.1 & B.1.617.2 AMONG CASES EXPORTED FROM INDIA (GISAID data Singapore & Japan)



Multinomial fit

