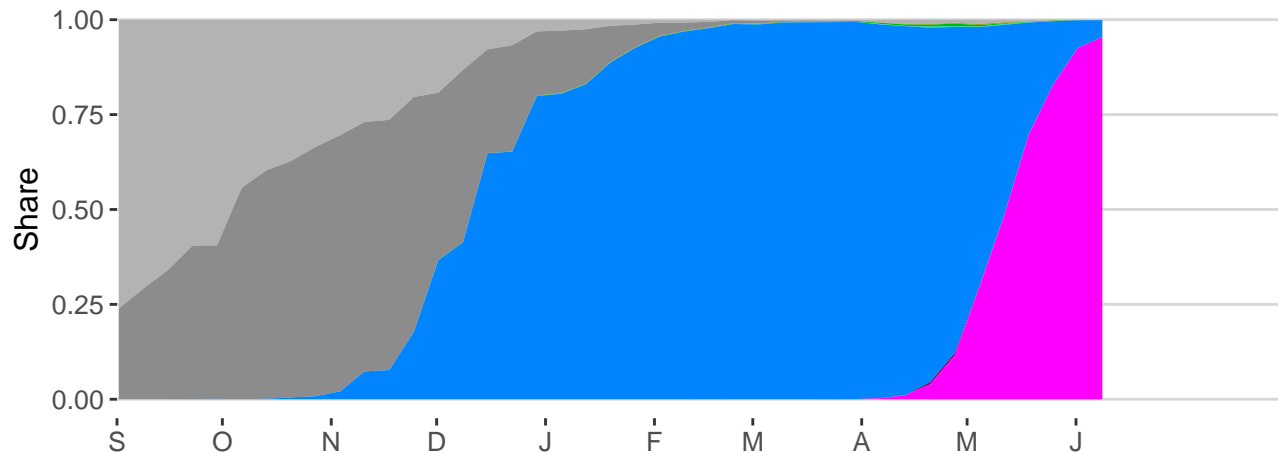


SPREAD OF SARS-CoV2 VARIANTS B.1.617.1 & B.1.617.2 IN ENGLAND (Sanger Institute baseline surveillance data)



Multinomial fit

