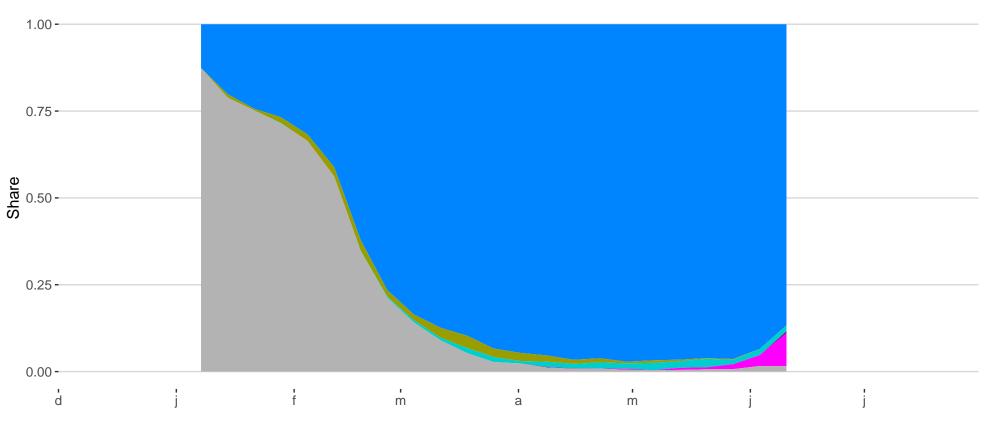
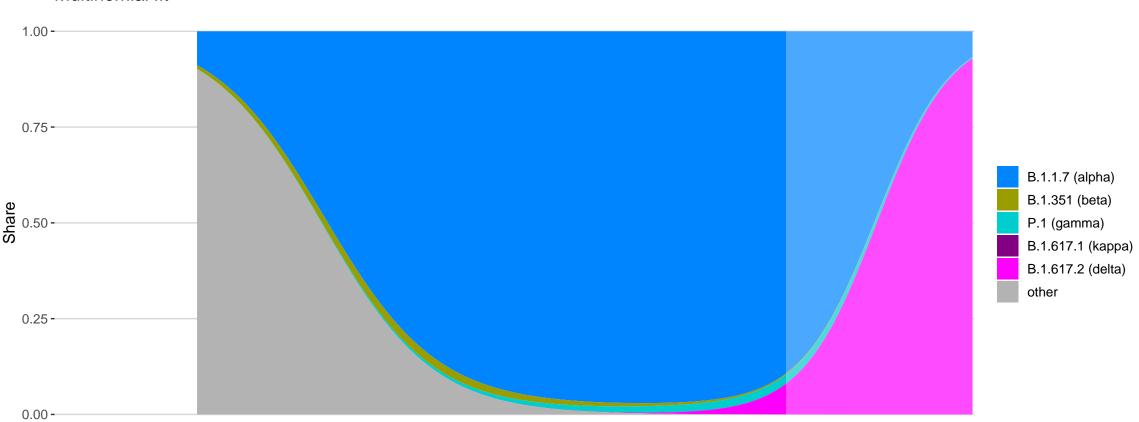
Spread of SARS-CoV2 variants of concern in Netherlands (baseline surveillance))



Multinomial fit



m

a

m