

Figure S1: Time course of measles genotypes detected in Taiwan between 2015 and 2020. The vertical axis indicated the number of each genotype detected, and the horizontal axis indicated the epidemic week of each year.

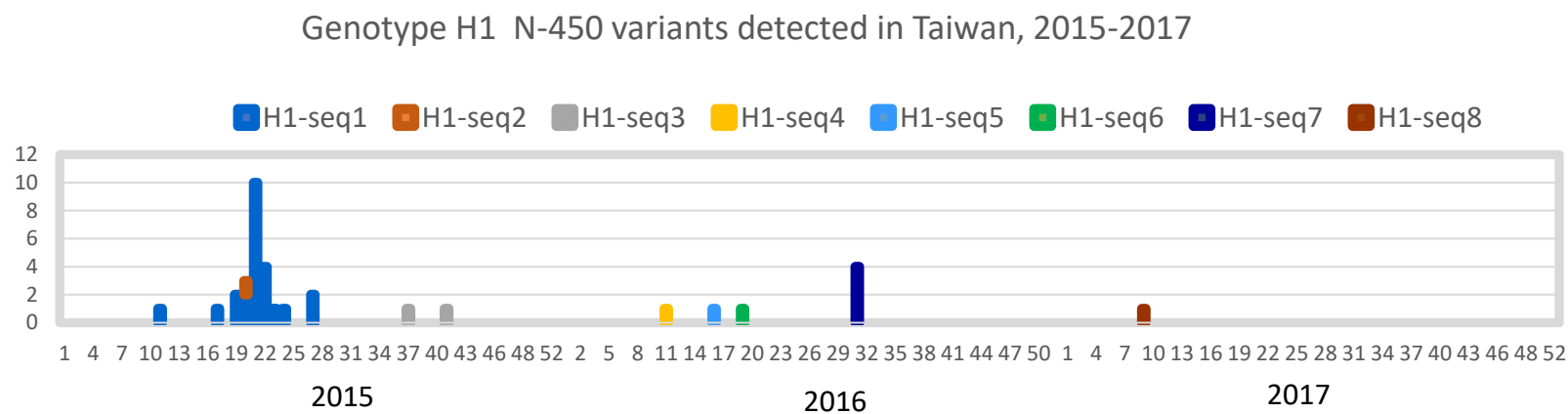


Figure S2: Time course of 8 measles genotype H1 variants detected in Taiwan between 2015 and 2017. The vertical axis indicated the number of each variant detected, and the horizontal axis indicated the epidemic week of each year.

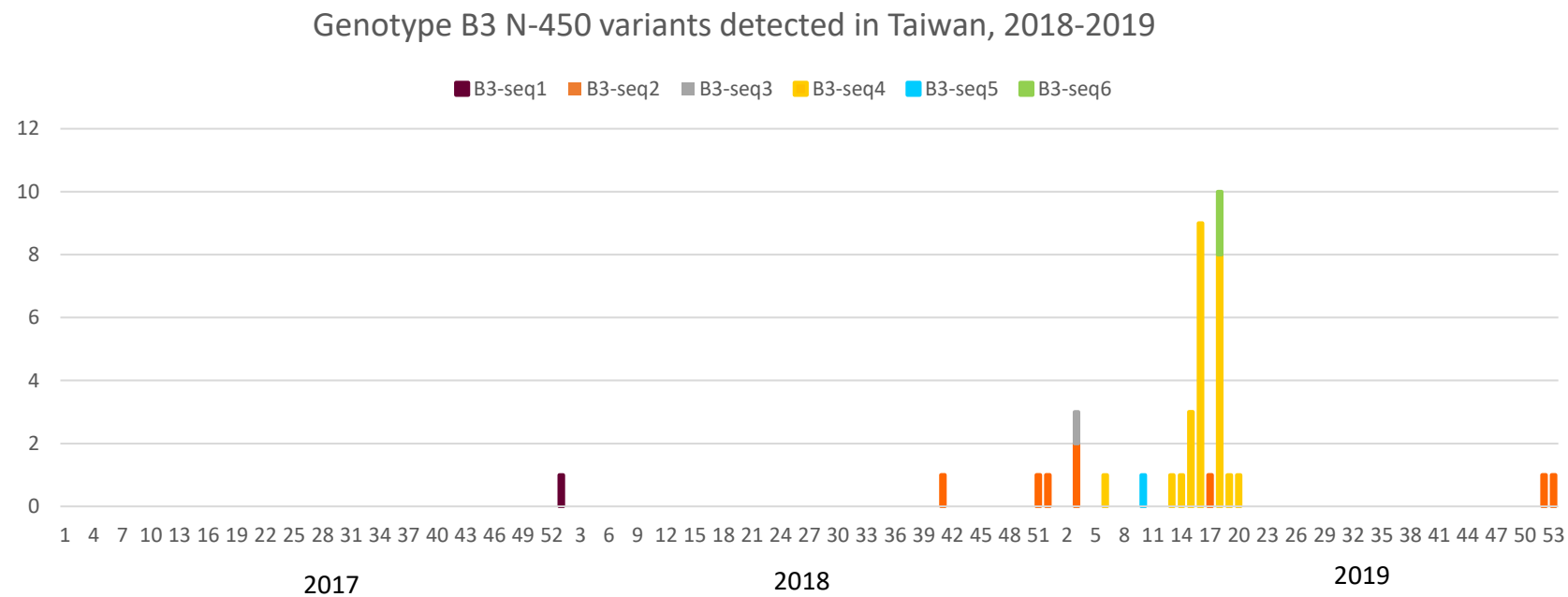


Figure S3: Time course of 6 genotype B3 variants detected in Taiwan between 2018 and 2019. The label of vertical axis and horizontal axis are described as in the legend of Figure S2.

