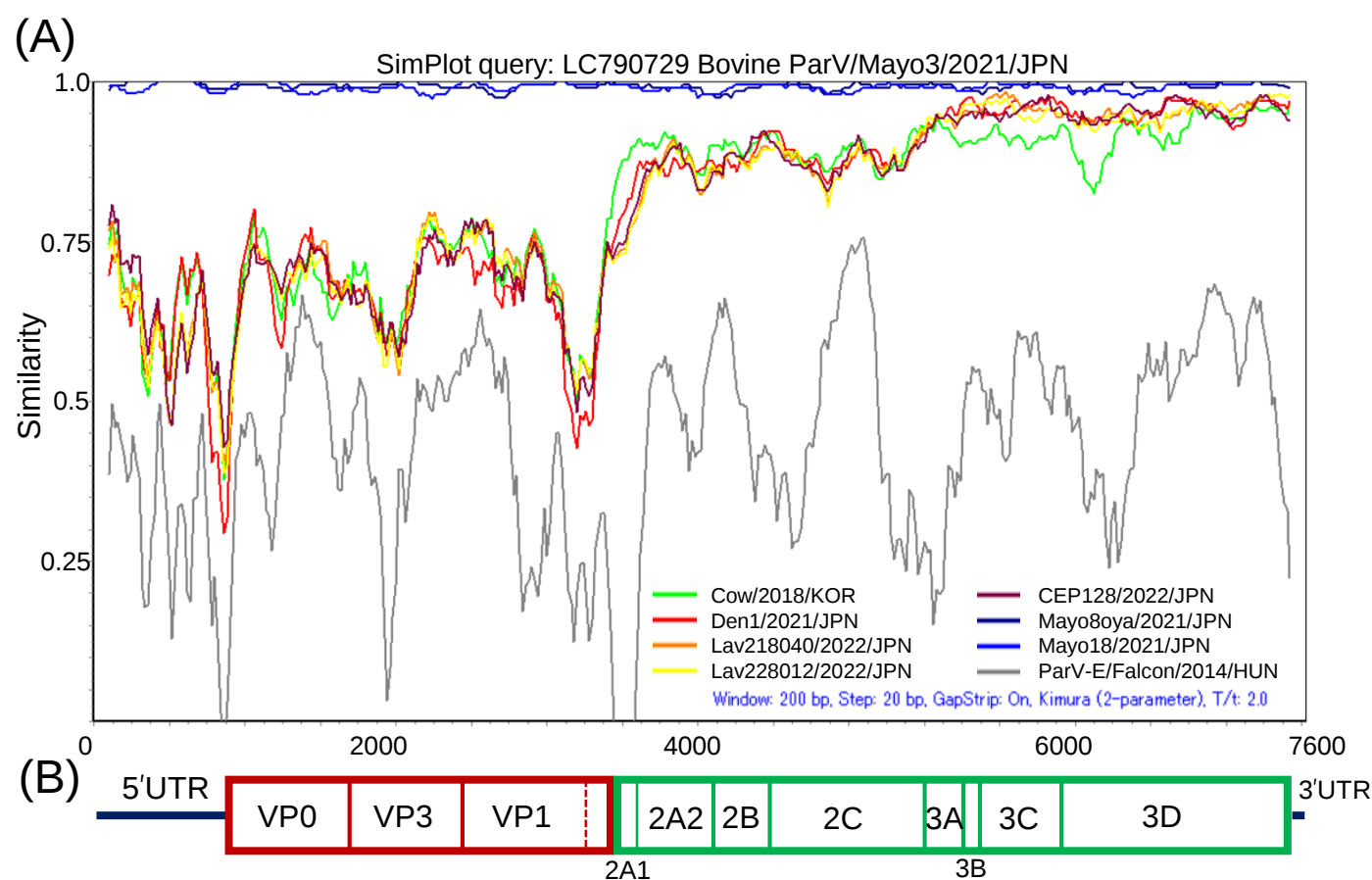




Supplementary Fig. Oba et al.

Supplementary Fig. Similarity plot analysis (A) and genome structure of the bovine parechovirus (B). Similarity plots of the complete genome sequences of Bo_ParV/cow/2018/KOR (green curve), Bo_ParV/Den1/2021/JPN (red curve), Bo_ParV/Lav218040/2022/JPN (orange curve), Bo_ParV/Lav228012/2022/JPN/ (yellow curve), Bo_ParV/CEP128/2022/JPN (purple curve), Bo_ParV/Mayo8oya/2021/JPN (dark blue curve), Bo_ParV/Mayo18/2021/JPN (light blue curve), ParV-E/Falcon/2014/HUN (gray curve), and Bo_ParV/Mayo3/JPN/2021 as query sequences, with a sliding window of 200 nt and a moving step size of 20 nucleotides.