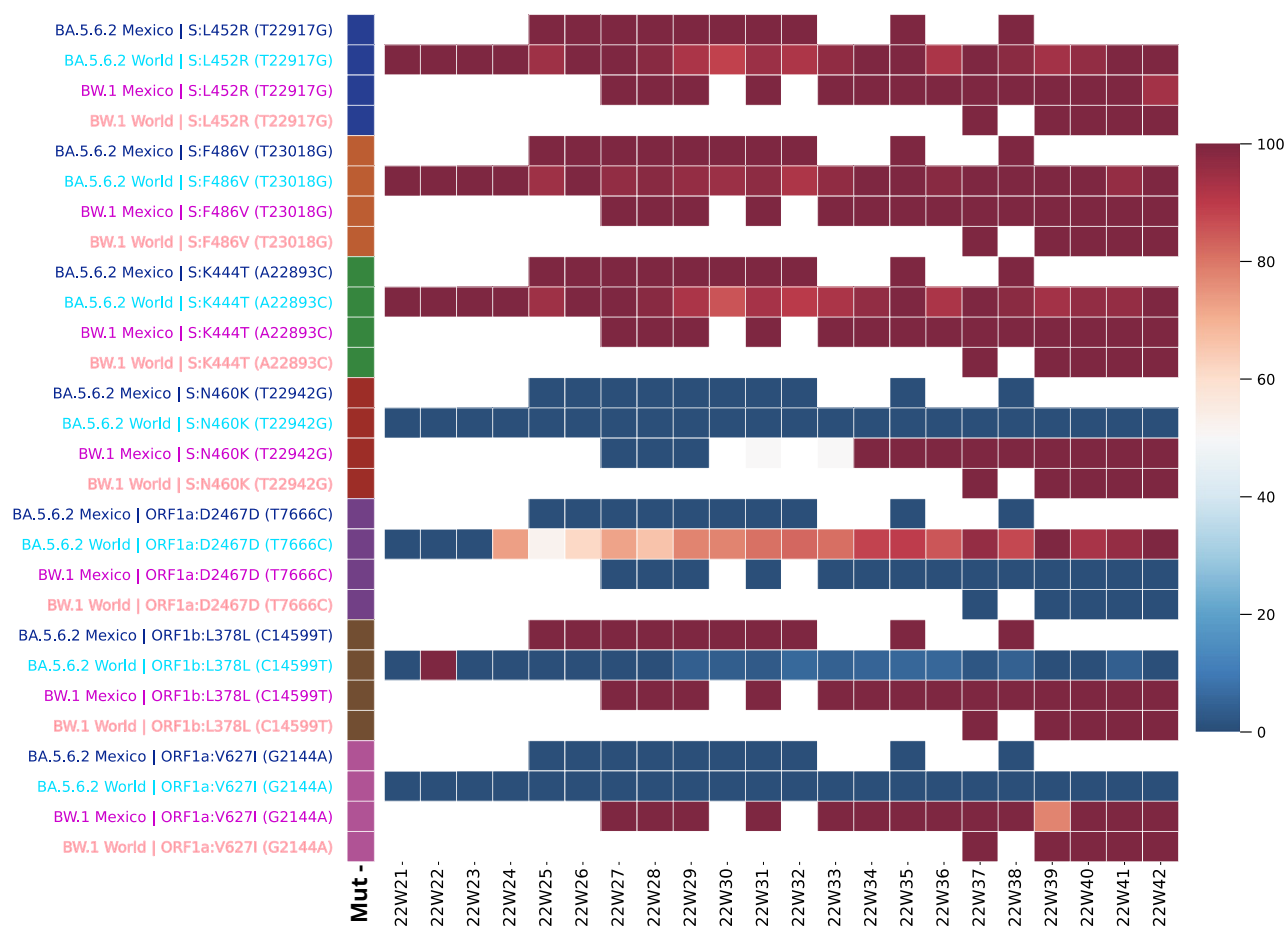


Title: SARS-CoV-2 BW.1, a fast-growing Omicron variant from southeast Mexico bearing clinically relevant escape mutations

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Supplementary Figure 1. Heatmap showing the longitudinal evolution of key mutations in genomes from Mexico and the rest of the world (y axis) from variants BA.5.6.2 and BW.1 per week (x axis). Numbers in each cell represent the percentage of sequences in that month that bear each mutation. Blank spaces represent an absence of samples.