



S9 Fig. Structural mapping of additional Yamagata-lineage ‘clade-defining’ trunk substitutions on influenza B polymerase complex. Substitutions not located at potential PB1/PA or PB1/vRNA interface regions are highlighted as spheres coloured by emergence in Yamagata-lineage clade 2 (orange) or clade 3 (red) viruses. See Fig 6 legend for details.