*In vitro* selection of Remdesivir resistance suggests evolutionary predictability of SARS-CoV-2

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We have included the data used to generate the figures and relevant statistic in their own file. Ie the data for Fig 3D is in a subfolder Fig3D&E within the folder Figure 3.

All supplementary figure data has been included. Found in a folder labeled supplementary data.

All datasets have also been included.