Source data for figures in “**M segment-based minigenomes and virus-like particle assays as an approach to assess the potential of tick-borne *Phlebovirus* genome reassortment”** by Rezelj et al, 2018.

1. Measurements of FFLuc and *Renilla* luciferase to assess the effectiveness of the HRTV minigenome system. Results represent n=3 experimental repeats.

**Tab 1: Fig 1**

1. Measurements of FFLuc and *Renilla* luciferase in UUKV, HRTV and SFTSV M segment-based minigenomes using UUKV, HRTV or SFTSV N and L proteins. Results represent n=3 experimental repeats (\*, P > 0.05).

**Tab 2: Fig 2A**

**Tab 3: Fig 2B**

**Tab 4: Fig 2C**

1. Measurements of FFLuc and *Renilla* luciferase to assess activity of virus-like particles containing S or L segment minigenomes using UUKV, HRTV or SFTSV glycoproteins. Results represent n=4 experimental repeats (\*, P > 0.05)

**Tab 4: Fig 3A**

**Tab 5: Fig 3B**

**Tab 6: Fig 3C**

1. Measurements of FFLuc and *Renilla* luciferase to assess activity of wild-type or mutated UUKV or SFTSV M segment-based minigenomes. Results represent n=3 experimental repeats (\*, P > 0.05).

**Tab 7: Fig 4E**

**Tab 8: Fig 4F**

**Tab 9: Fig 4G**

**Tab 10: Fig 4H**