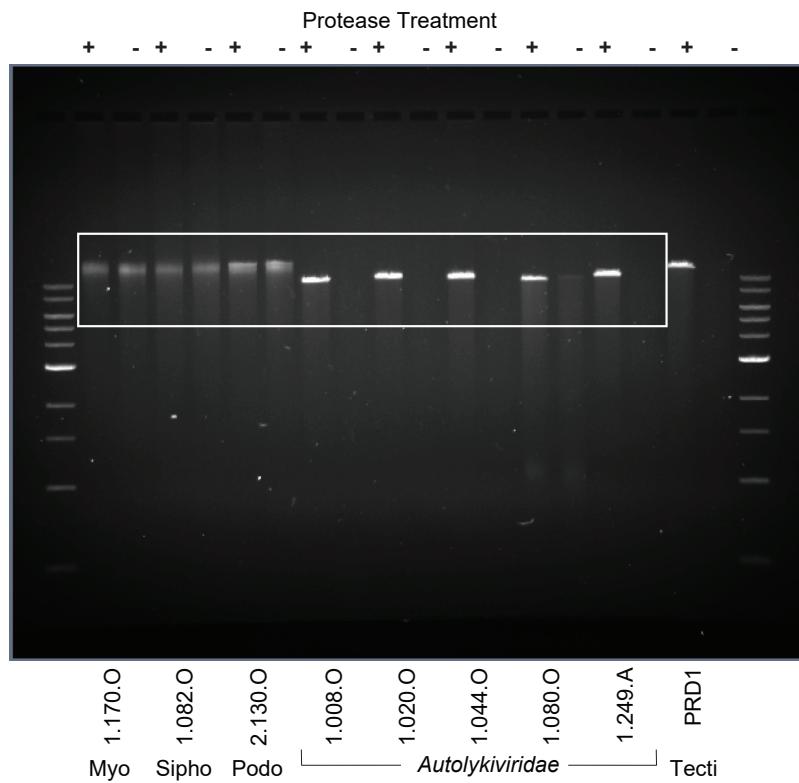


SUPPLEMENTARY INFORMATION

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Supplementary Figure 1. Comparison of tailed virus and autolykivirus genome recovery with and without protease treatment. Protease-treated sample load volumes normalized to 50 ng, equal volumes of untreated partner samples in adjacent lanes. White outline indicates cropped region shown in Fig. 3c.