



Fig. S1. Identification of V8 proteolytic fragments. (A) Limited proteolysis of UL11-StII with V8 protease. N-terminal sequence of the major proteolytic fragment is shown along with the molecular masses calculated from the sequence (MM_{calc}) and determined by mass spectrometry (MM_{MS}). (B) MALDI-TOF mass spectrometry analysis of the same digest. The x axis shows mass to charge ratio (m/z) in Da and the y axis shows relative abundance (%) scaled for the mass range in x. Molecular mass (MM_{MS}) of the major peak is shown along with the mass calculated from the sequence (MM_{calc}).