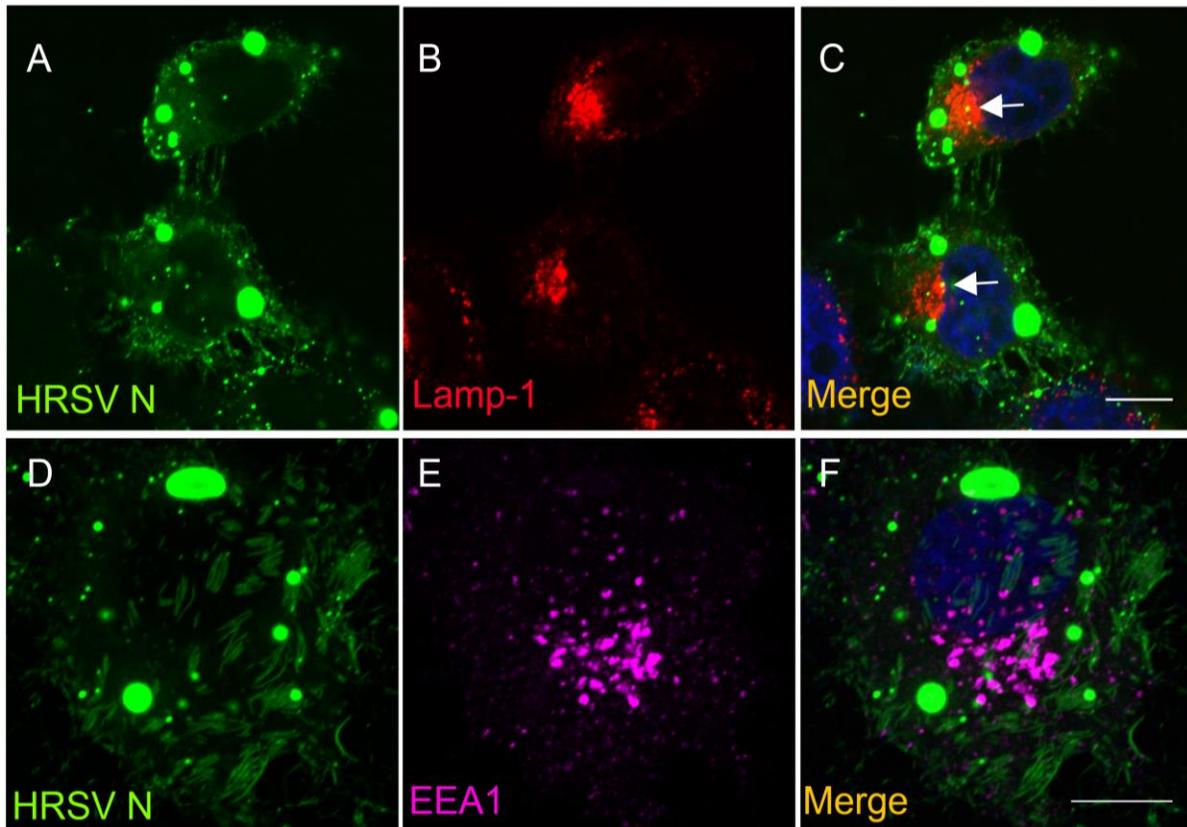


are representative of three independent experiments. (D) graph of the quantity of IBAGs counted in inclusion bodies in A3.01 and HEp-2-infected cells. Magnification 63x. Scale bars = 10 μ m. To the (C) and (D) graphs was counted at least 10 fields of three independent experiments from single focal plane of Z-stacks imaging acquisition. The statistical method used was Student's T-test, $p^* < 0.05$, $p^{**} < 0.01$, and $p^{***} < 0.001$.

Supplementary Figure 2



Supplementary figure 2. Co-localization of the HRSV N protein with Lamp-1 and EEA1 in HEp-2 cells. (A-F), HEp-2 cells, were infected with HRSV, and 48 hpi the cells were subjected to immunofluorescence. (A-C) co-localization of HRSV N (A) in green, Lamp-1 (B) in red, and the merge (C). Arrows point to the co-localizations. (D-F), co-localization between HRSV N (D) in green, EEA1 (E) in magenta, and the merge (F). The images were acquired in a Zeiss 780 Confocal, and are representative of a single focal plane of three independent experiments. Scale bar =10 μ m.