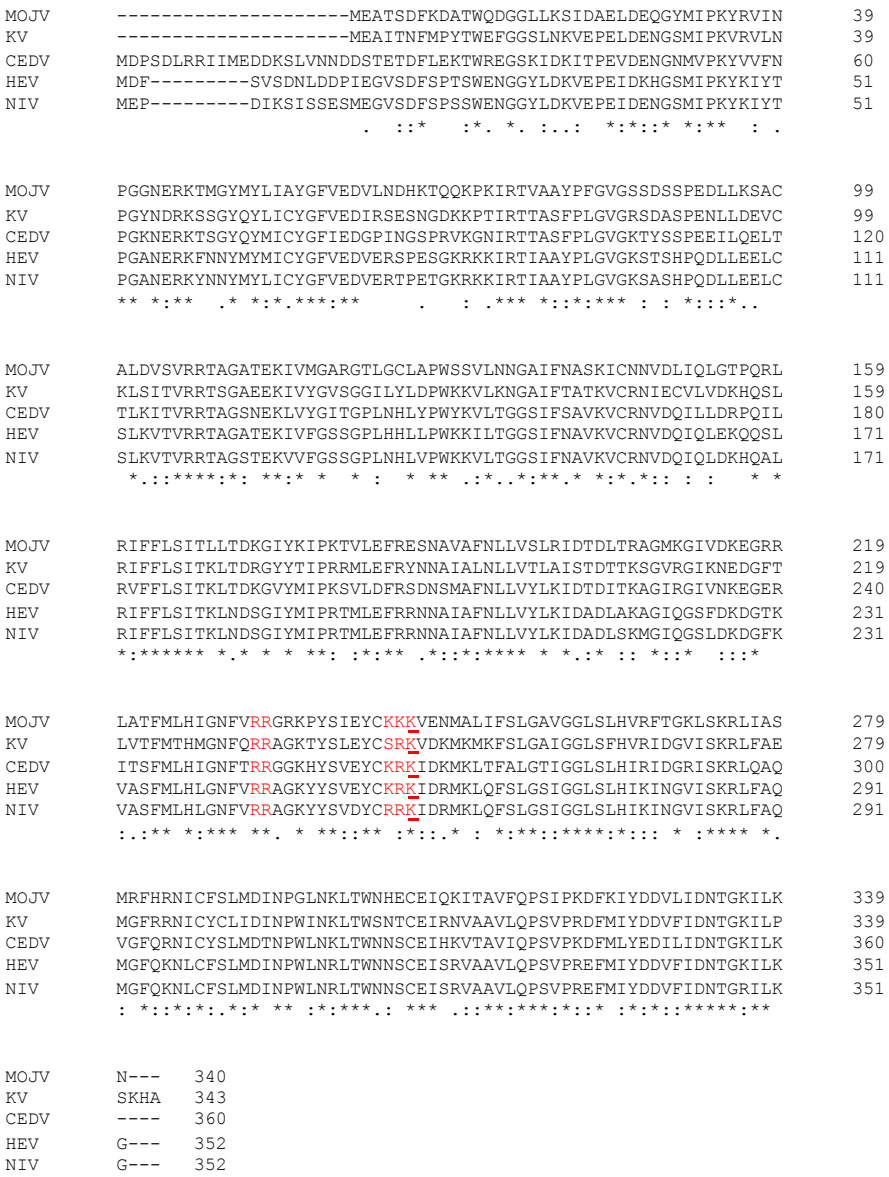
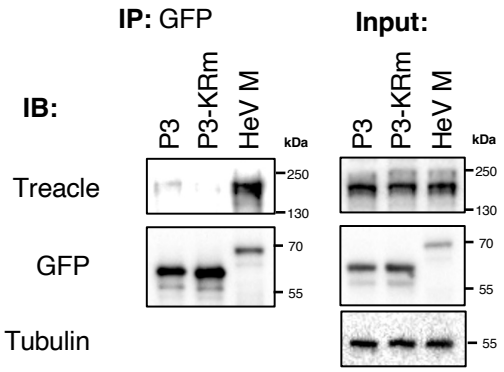
Figure S1



<u>Supplementary Figure S1</u> **Sequence alignment of henipavirus M proteins used in study.**

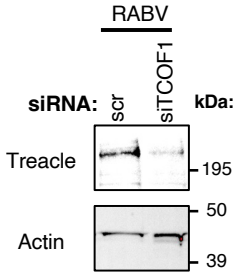
Alignment using Clustal Omega (version 1.2.4) of HeV M (Accession number: AEB21196.1), NiV M (AAY43914.1), CedV M (AFP87277.1), MojV M (AHM23775.1) and KV M (YP_009091836). Positions with an absolutely conserved residue (*); residues with strong similarity (:); Residues with weak similarity (.). Key residues of the bipartite NLS (two clusters of basic residues) are highlighted in red font; K258 residue (bold underlined) is absolutely conserved across all species.

Figure S2



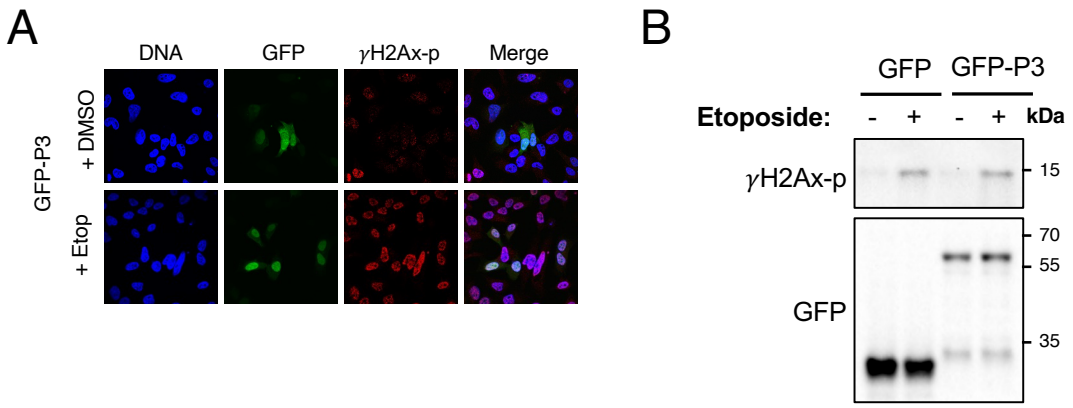
Supplementary Figure S2 **HeV M co-precipitates Treacle to a greater extent than P3**. HEK-293T cells were transfected to express the indicated GFP-fused proteins, prior to lysis and IP with GFP-Trap beads (as performed in Figs. 3A & 5A). IP and input samples were analysed by IB and probed with the indicated antibodies.

Figure S3



Supplementary Figure S3 **Confirmation of the effectiveness of siRNA knockdown**. HEK-293 cells were transfected with siRNA (scr or siTCOF1) for 48 h (see Materials and Methods for details) prior to infection by RABV (MOI 0.01). Cells were lysed at 48 h p.i. and analysed by IB using the indicated antibodies.

Figure S4



Supplementary Figure S4 **Expression of RABV P3 does not induce DNA-damage in HeLa cells**.

HeLa cells transfected to express GFP-P3 protein were treated without (DMSO) or with etoposide at 21 h p.t. for 3 h prior to (A) fixation and immunostaining for γH2AX phosphorylated at S139 (γH2AX-p), and imaging by CLSM (images representative of 10 fields of view taken from two independent experiments) or (B) lysed and analysed for IB analysis using the indicated antibodies.