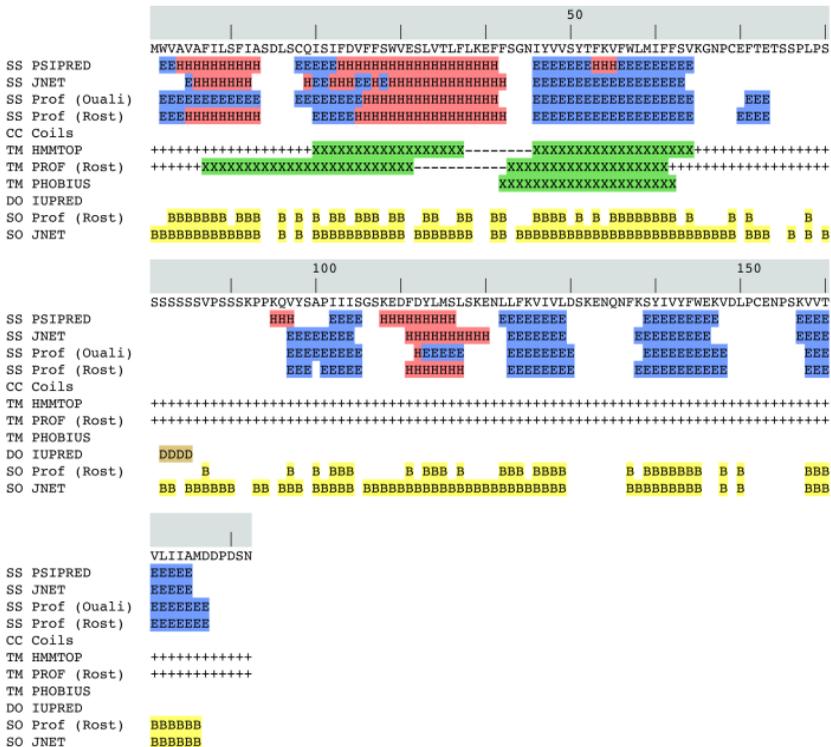


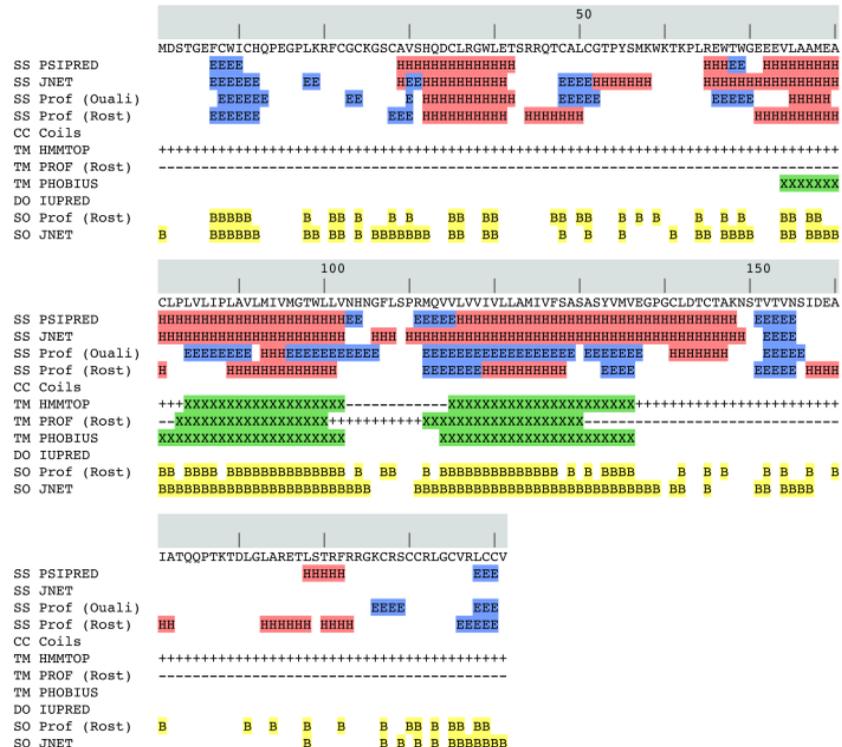
Supplemental Figure 1. Protein secondary structure prediction of RPHM21 and the ubiquitin ligase MK3. Both proteins showed two transmembrane regions. N-terminus and C-terminus "outside". (Quick2D; http://toolkit.tuebingen.mpg.de/quick2_d).

RPHM21 (from the M-type complete mitochondrial genome, GenBank: AB065374). MK3 of Murid herpesvirus 4 (NCBI Reference Sequence: NP_044852.1).

RPHM21



MK3



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 CC = Coiled Coils
 TM = Transmembrane ('+'=outside, '-'=inside)
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