

HIV co-opts a cellular antiviral mechanism, activation of stress kinase PKR by its RNA, to enable splicing of *rev/tat* mRNA

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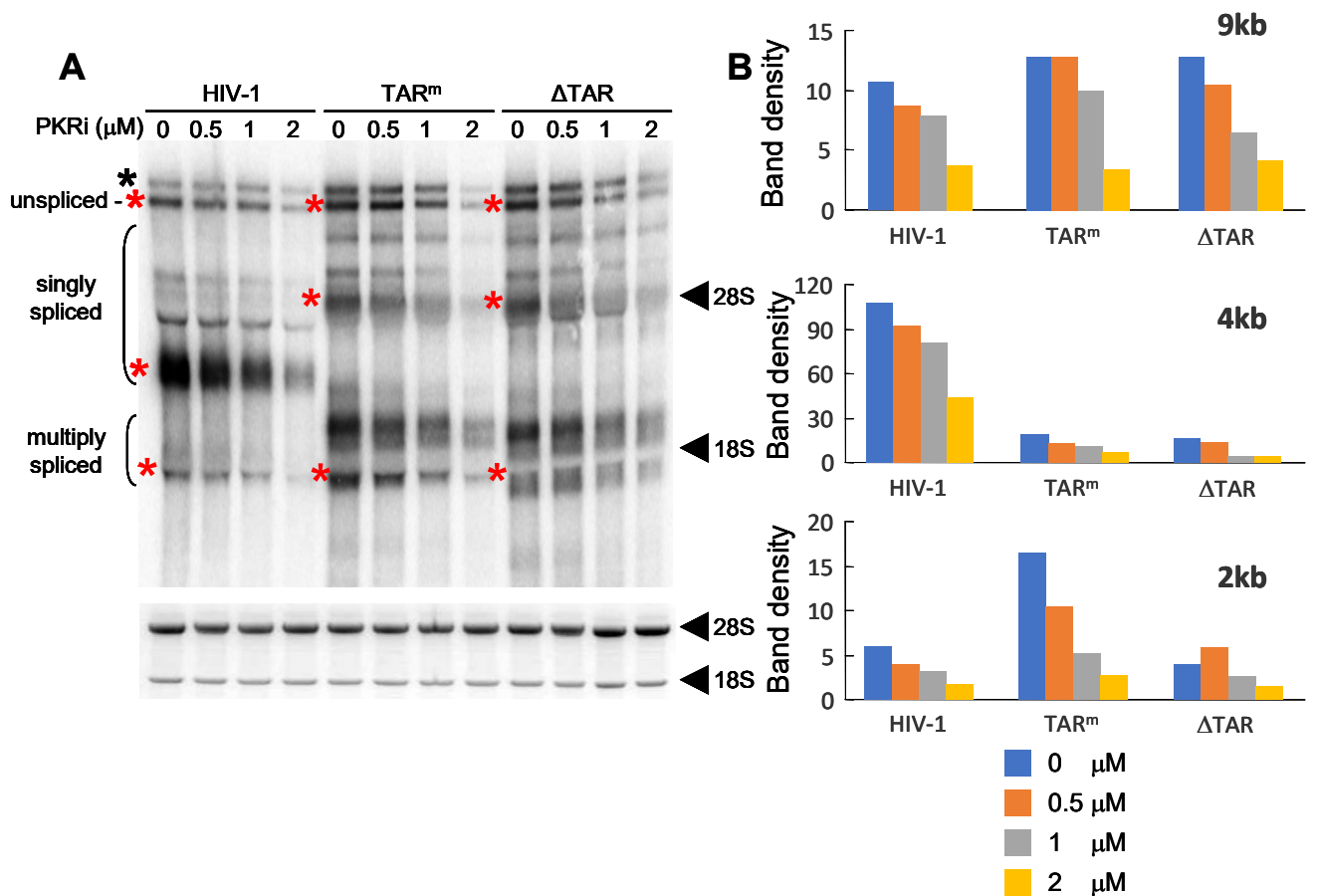
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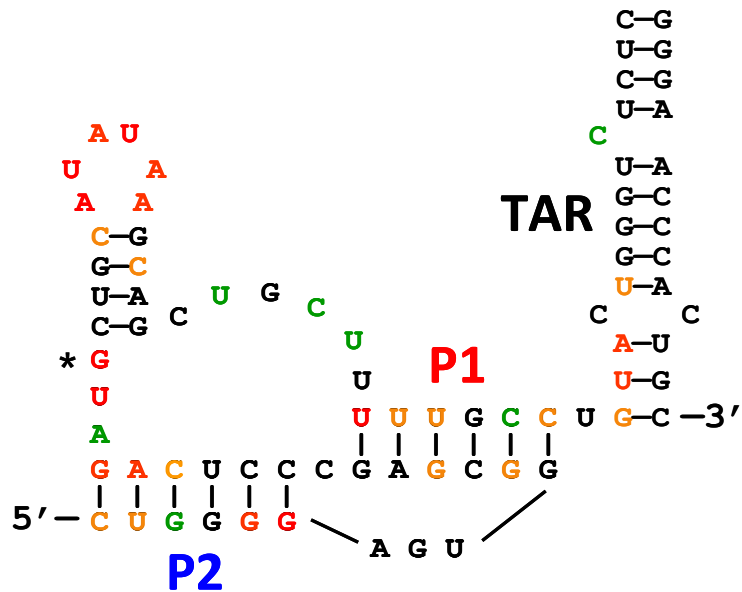
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Supplementary Figures



Additional file 1: Fig. S1 PKR inhibitor impairs expression of all size classes of HIV mRNA. **A** Representative mRNA species from each mRNA class in Fig. 1C (unspliced, 9 Kb; singly spliced, 4 Kb; multiply spliced, 2 Kb) are denoted by red asterisks. The shift in spliced mRNA banding pattern between the HIV-1 wt and the HIV-rtTA variants (TAR^m and ΔTAR) is caused by sequence differences (rtTA replaced Nef, different 3'UTR). **B** The denoted species from each mRNA class were quantified using ImageJ software (<https://imagej.nih.gov/>) (PKRi, μM).



Additional file 1: Fig. S2 Chemical reactivity of HIV pseudoknot stems P1 and P2. Secondary structure upstream of TAR and lower part of the TAR stem are shown for NL4-3 HIV-1 RNA. G labeled (*) is C in HIV-1 LAI. Chemical reactivity by SHAPE is shown, colors denote reactivity as high (red), moderate (orange), low (green) and little or none (black) [40].