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## **SUPPLEMENTARY MATERIALS**

## Identification of the initial nucleocapsid recognition element in the HIV-1 RNA packaging signal

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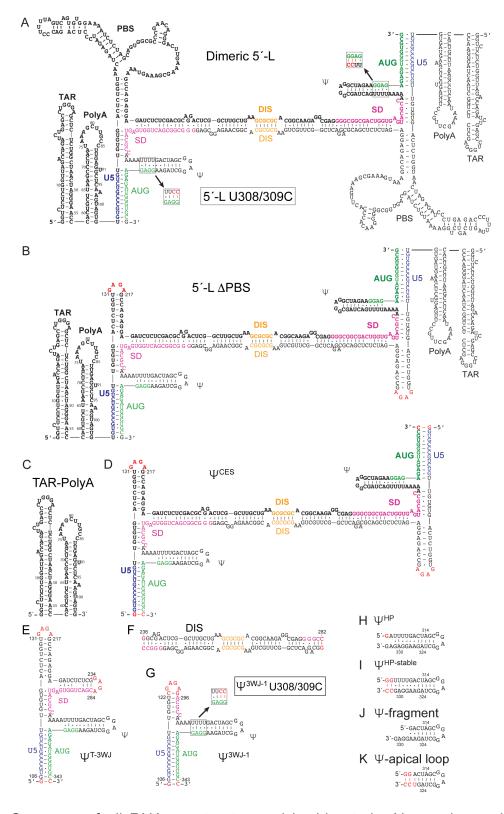
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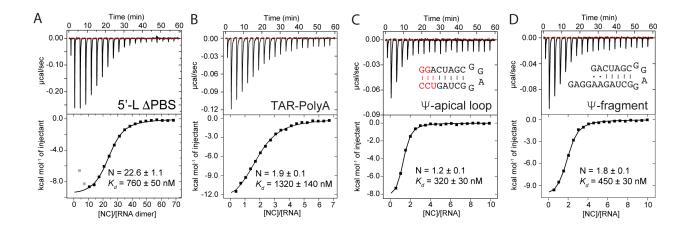
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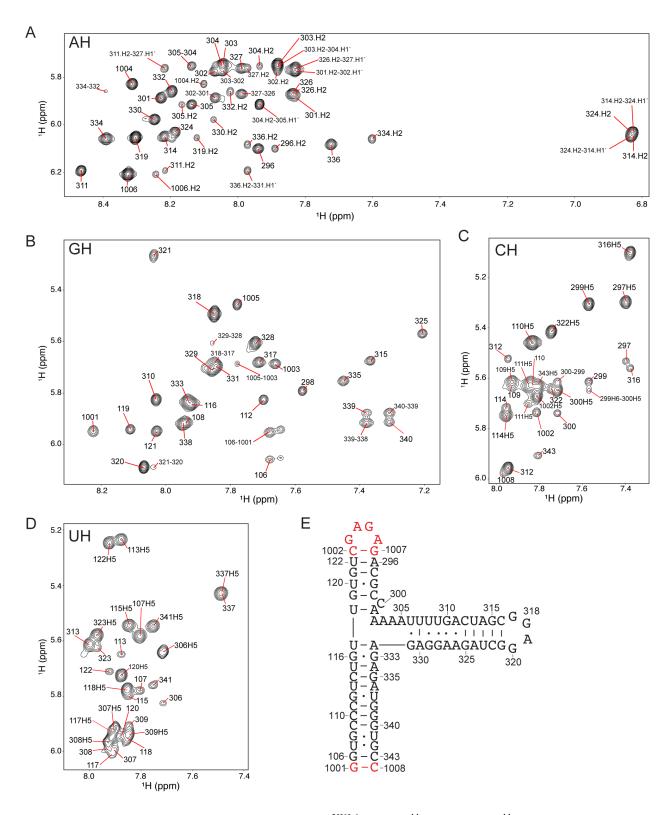
**CONTAINS SIX FIGURES AND FIGURE CAPTIONS** 



**Fig. S1.** Summary of all RNA constructs used in this study. Non-native residues are shown in red.



**Fig. S2.** ITC data of HIV-1 5´-L constructs. (*A*) 5´-L with the PBS region deleted. The gray data points due to endothermic binding were not included in data fitting. (*B*) TAR and PolyA hairpins; (*C*) the upper stem region and the apical loop from the Ψ-stem loop; (*D*) A construct with Ψ-apical loop and a single stranded "GGAG" region.



**Fig. S3.** Chemical shift assignments for  $\Psi^{3WJ-1}$ . For A<sup>H</sup>- (*A*) and G<sup>H</sup>- (*B*) spectra, intraresidue H8-H1′ cross-peaks are denoted by residue number; inter-residue H8-H1′ are indicated by (residue number of H8)-(residue number of H1′); Intra-residue H2-H1′ are

shown by (residue number).H2; Inter-residue H2-H1´ are labeled by (residue number of H2).H2-(residue number of H1´).H1´. For C<sup>H</sup>- (C) and U<sup>H</sup>- (D) spectra, intra-residue H6-H1´ cross-peaks are denoted by residue number; inter-residue H6-H1´ are indicated by (residue number of H6)-(residue number of H1´); Intra-residue H6-H5 cross-peaks are labeled by (residue number).H5; Inter-residue H6-H5 cross-peaks are indicated by (residue number of H6).H6-(residue number of H5).H5. (E) Secondary structure of  $\Psi^{3WJ-}$ 1. Non-native residues are shown in red and numbered 1001-1008.

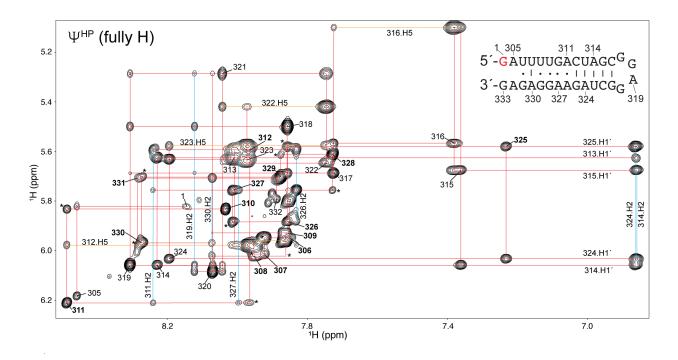
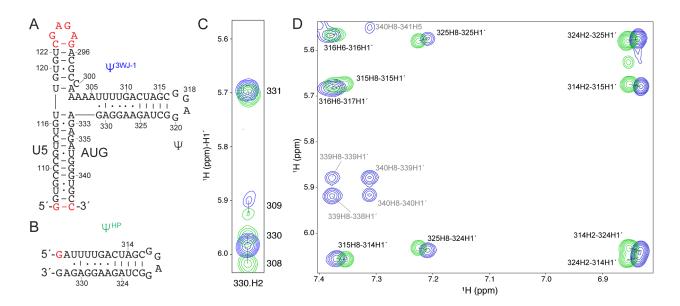
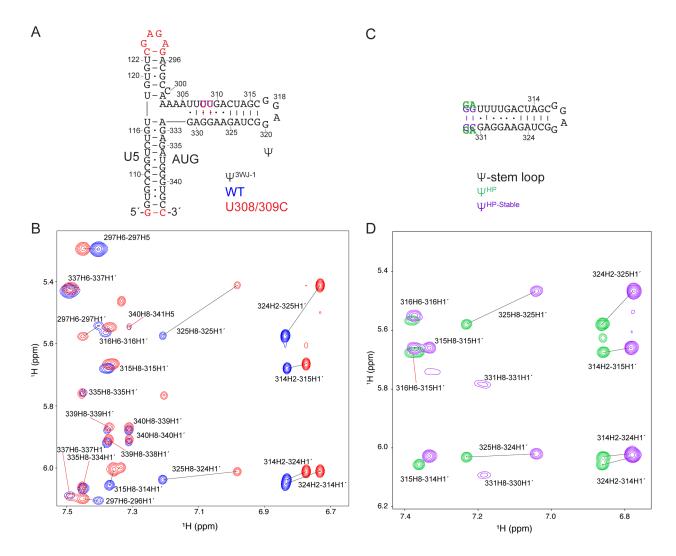


Fig. S4. Chemical shift assignments for  $\Psi^{HP}$ . Intra-residue H8/H6-H1′ cross-peaks are denoted by residue number. Adenosine H2 signals are indicated by blue lines; Pyrimidine H5 signals are labeled by yellow lines. Sequential H8/H6-H1′ cross-peaks for the [UUUUGAC]:[GAAGGAG] region are marked by stars.



**Fig. S5.**  $\Psi^{HP}$  is structurally similar to the same region in  $\Psi^{3WJ-1}$ . Secondary structure of  $\Psi^{3WJ-1}$  (*A*) and  $\Psi^{HP}$  (*B*). Non-native residues are shown in red. (*C*) 2D NOESY spectra overly of fully protonated  $\Psi^{HP}$  (green) and  $A^{2r}G^rU^r$ -labeled  $\Psi^{3WJ-1}$  (blue). (*D*) 2D NOESY spectra overly of fully protonated  $\Psi^{HP}$  (green) and fully protonated  $\Psi^{3WJ-1}$  (blue).



**Fig. S6.** Structural effects of stabilizing the [UUUU]:[GGAG] region. (*A*) Secondary structure of wild-type and G:U to G-C mutant of  $\Psi^{3WJ-1}$ . Non-native residues are shown in red. (*B*) 2D NOESY spectra overly of fully protonated wild-type (blue) and G:U to G-C mutant (red)  $\Psi^{3WJ-1}$ . (*C*) Secondary structure of  $\Psi^{HP}$  and  $\Psi^{HP-stable}$ . (*D*) 2D NOESY spectra overlay of fully protonated  $\Psi^{HP}$  (green) and  $\Psi^{HP-stable}$  (purple).