

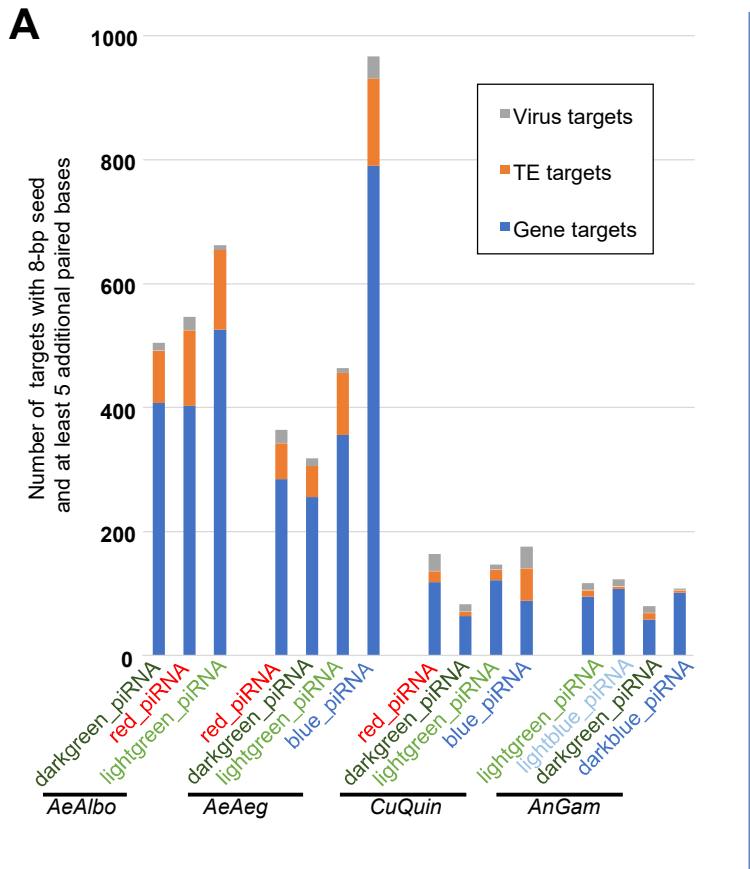


Figure S8. Predictions of gene mRNAs, transposons and virus transcripts potentially targeted by the four piRNAs in the Mosquito Conserved piRNA Cluster Locus (MCpiRCL). (A) Total counts of predicted targets from the four mosquito species. (B) For each of the *AeAlbo* piRNAs from its MCpiRCL, the top two mRNA, transposon and virus target sites is displayed, ranking based on the degree of base-pairing. (C) Top targets for the *AeAeg* piRNAs from its MCpiRCL. (D) Top targets for the *CuQuin* piRNAs from its MCpiRCL. (E) Top targets for the *AnGam* piRNAs from its MCpiRCL.

C

AeAeg MCPiRCL predicted targets

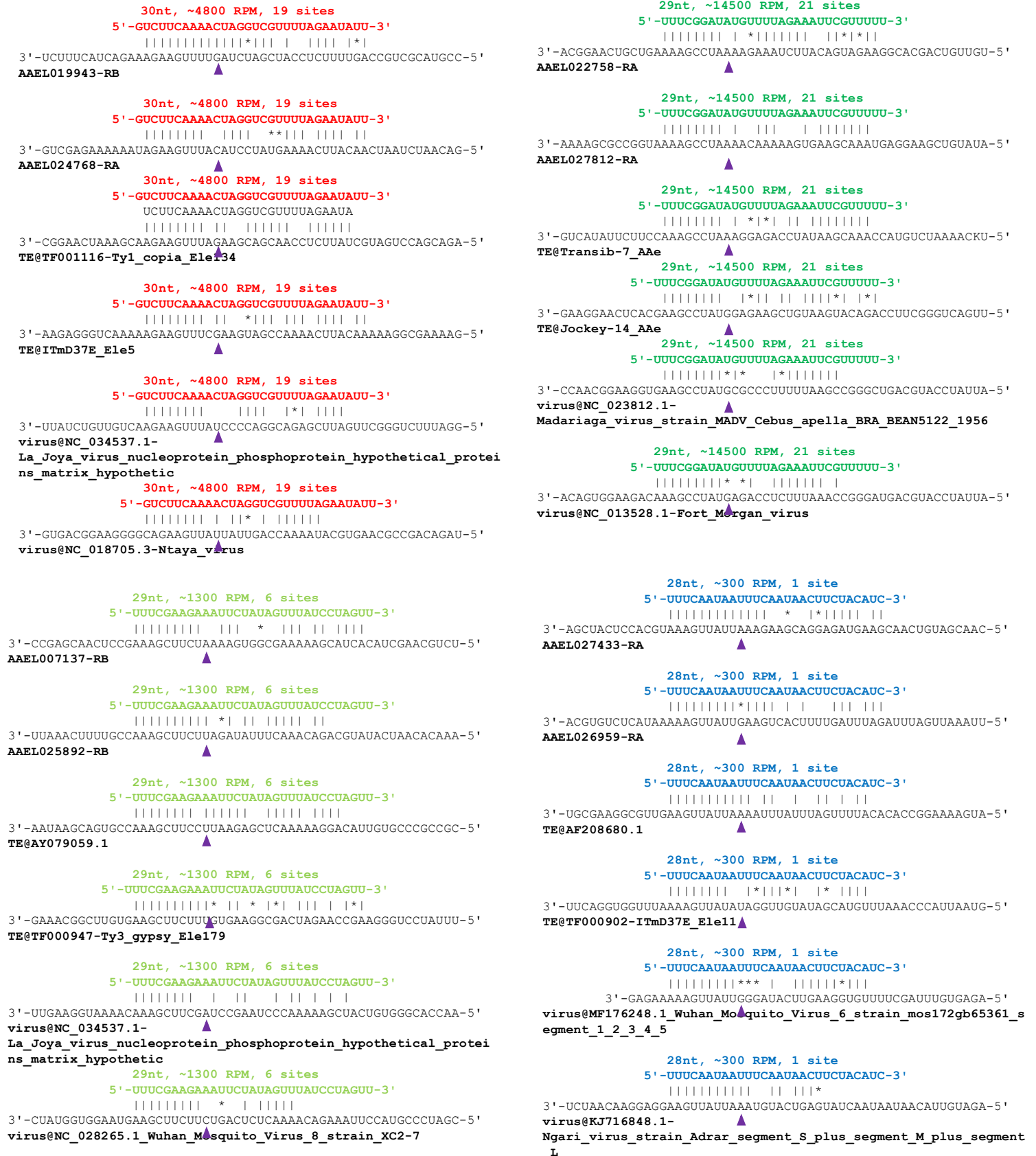


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D *CuQuin* MCpiCRL predicted targets

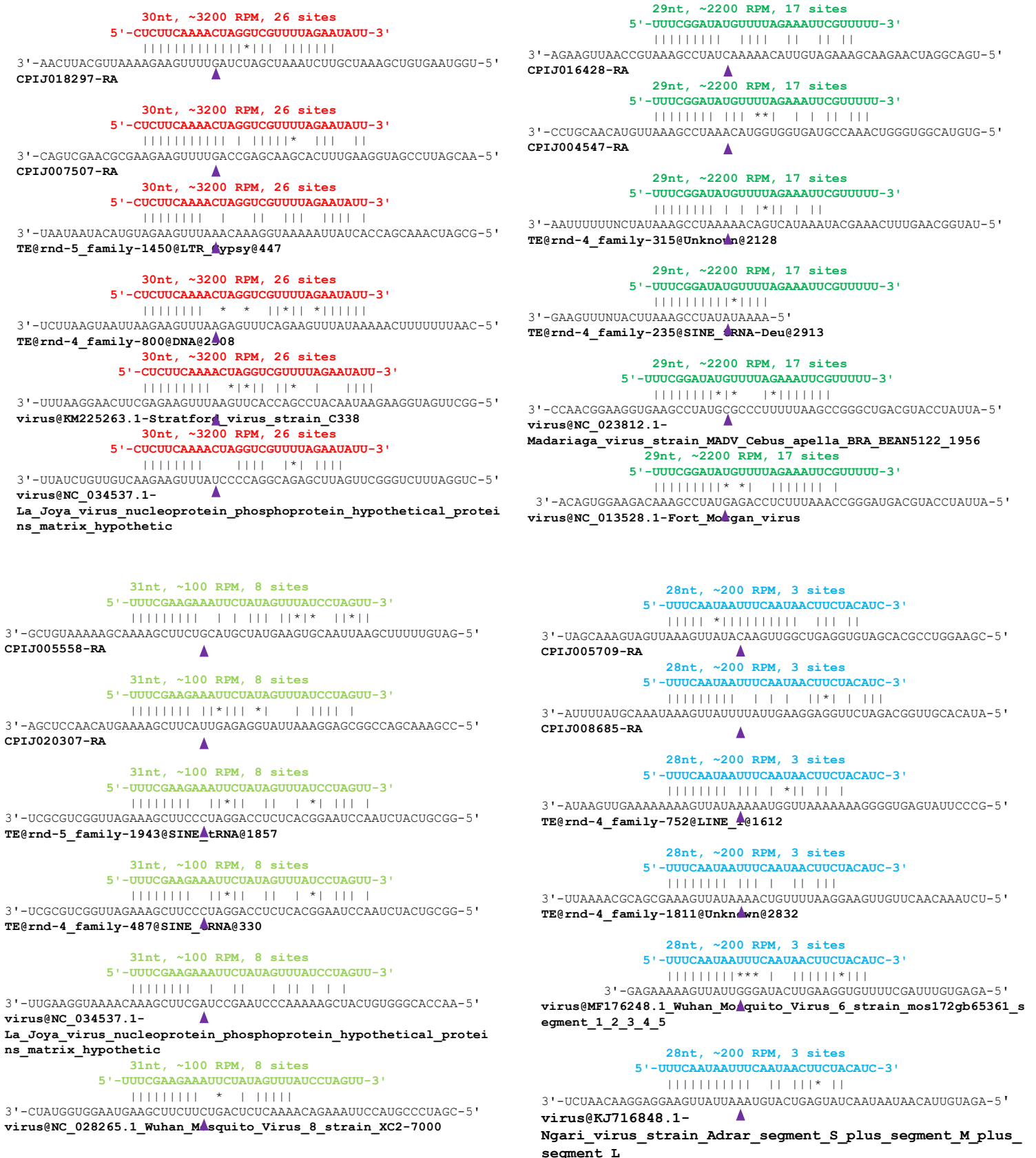


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AnGam MCPiCRL predicted targets

E

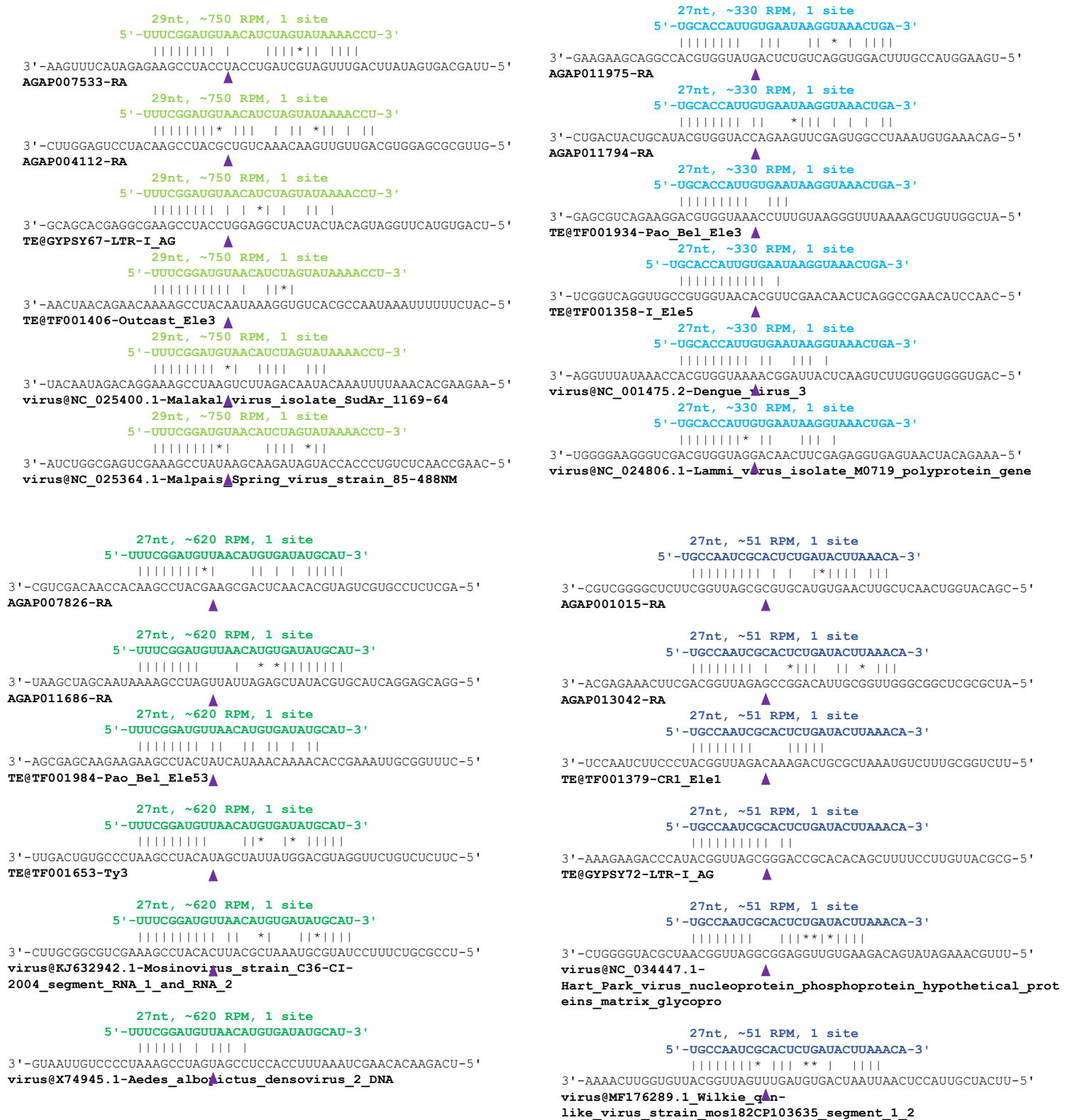


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