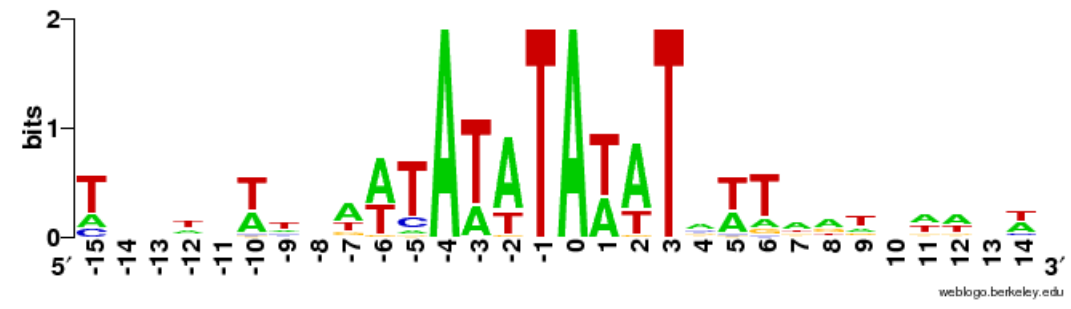
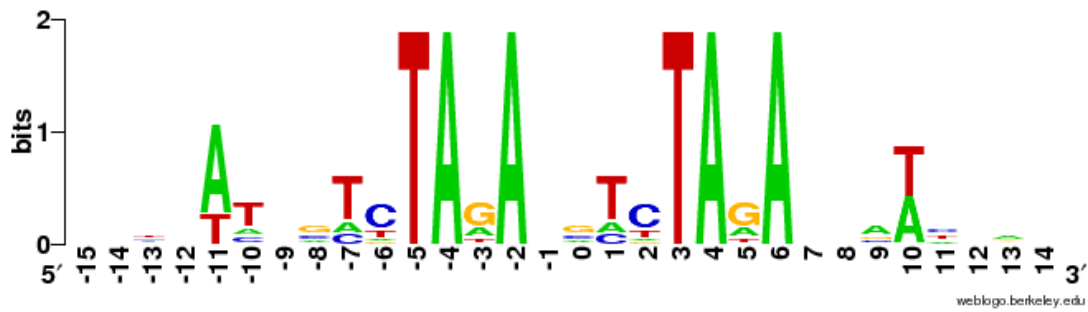


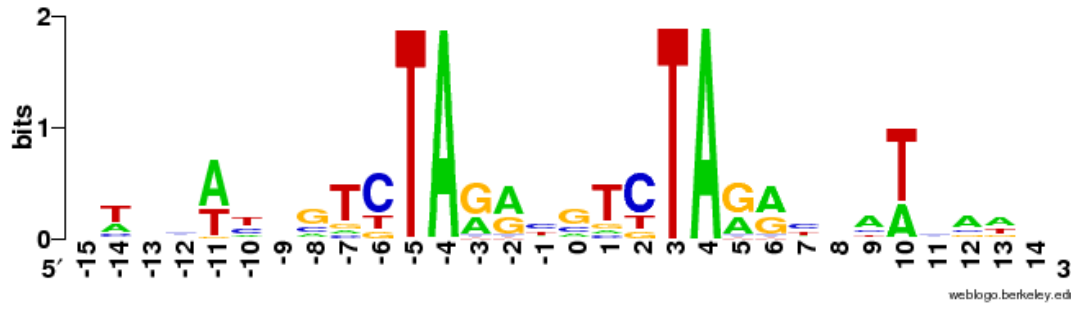
Sola1_NA_BM



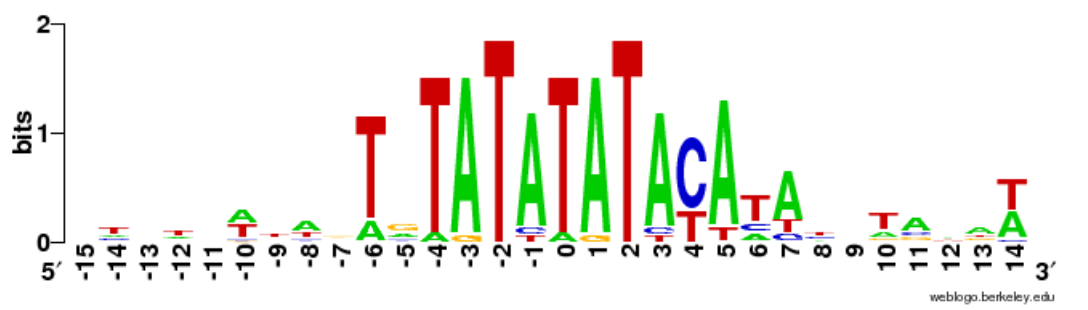
Sola1_NA_RP



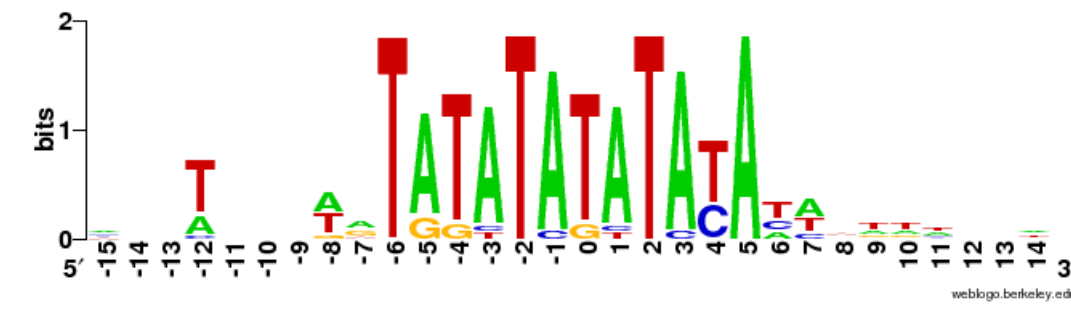
hAT1_NA_BM



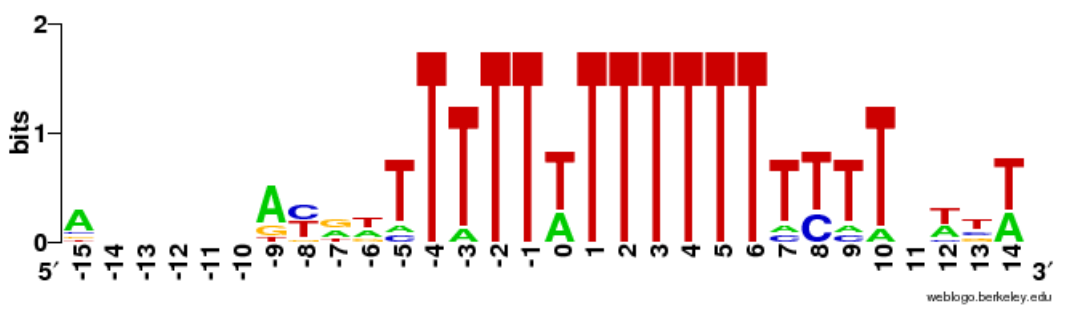
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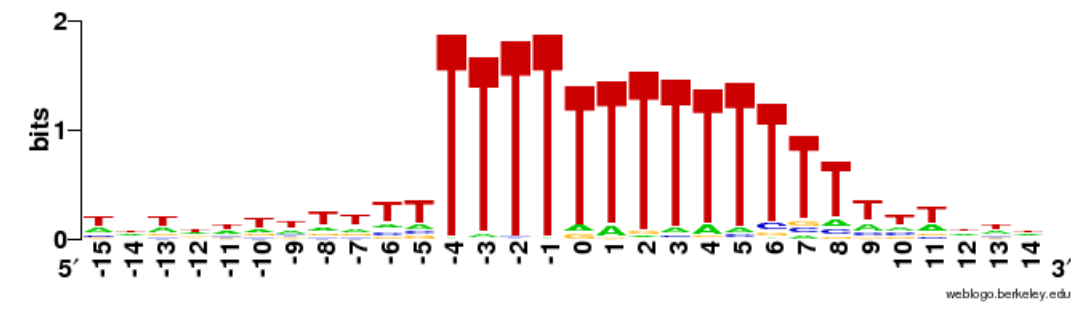
Ginger2_NA_BM



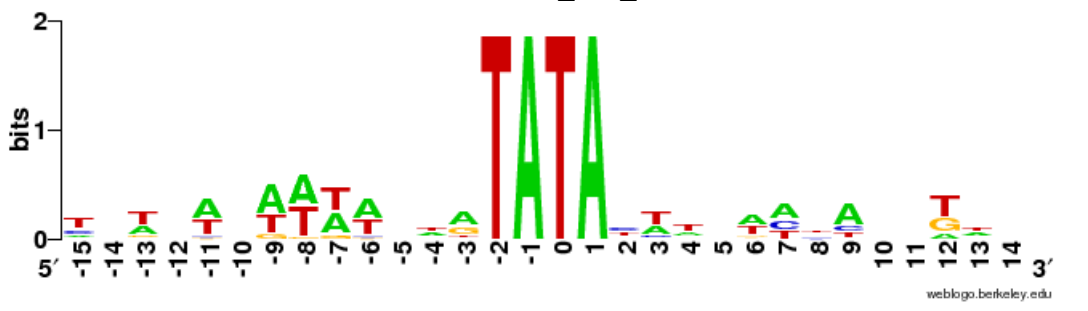
Ginger2_NA_RP



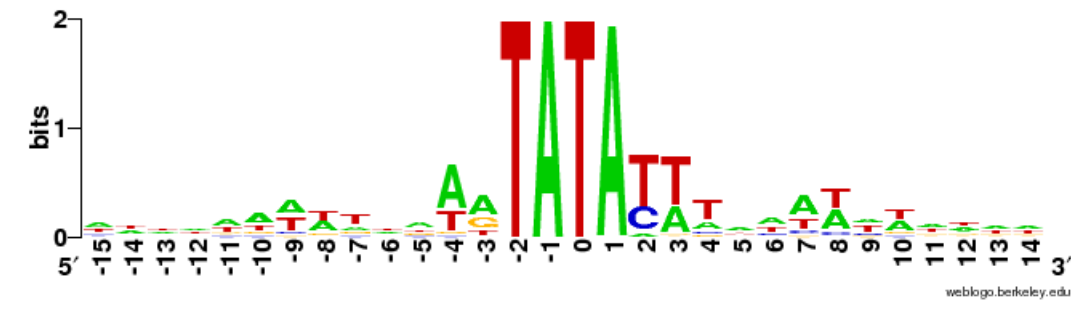
Novel_NA_BM



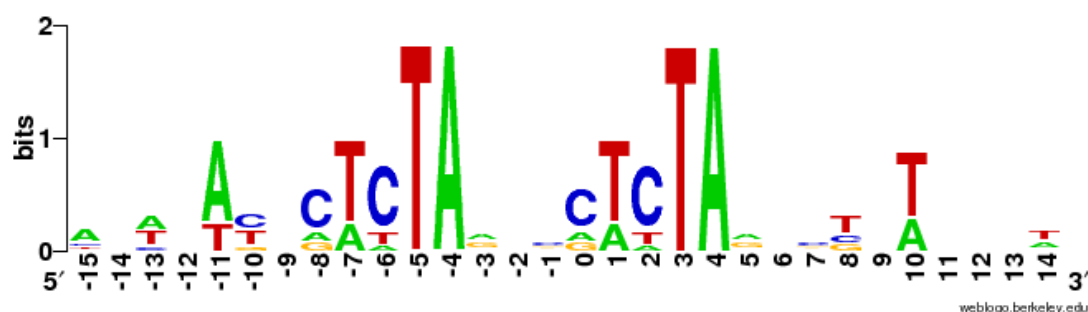
Novel_NA_RP



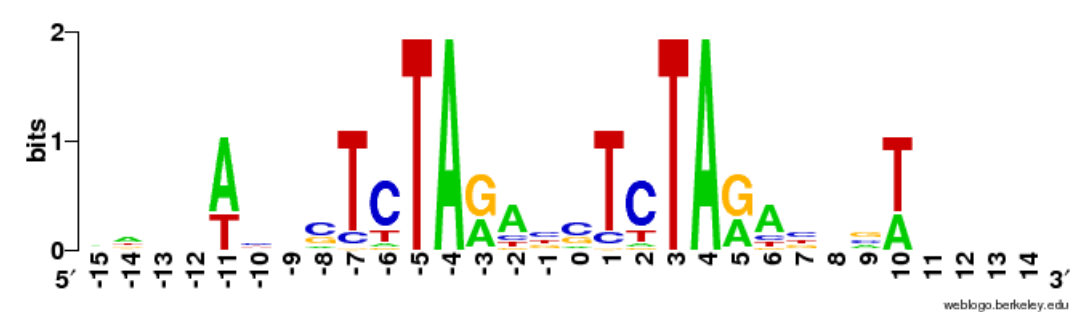
pogo_NA_BM



pogo_NA_RP



hAT2_NA_BM



hAT2_NA_RP

Fig.S3 Sequence logo of the regions flanking MITEs insertions. The 15 nt upstream and downstream of all full-length copies of these families in the silkworm and triatomine bug are presented in each logo.

The vertical axis is a measure of sequence information, which has a maximum value of 2 and is proportional to the level of sequence conservation at each position.