



Fig.S2 The frequency distributions (ordinate) of sequence identities (abscissa) of MITEs in the silkworm and triatomine bug. The percentage identity is based on BLAST search, using the consensus from each species as the query. The sequences shorter than 100 bp were excluded. The blue or purple lines represent the tendency estimated by a simple moving average of the percentage identity scores. The name of each family is also shown.