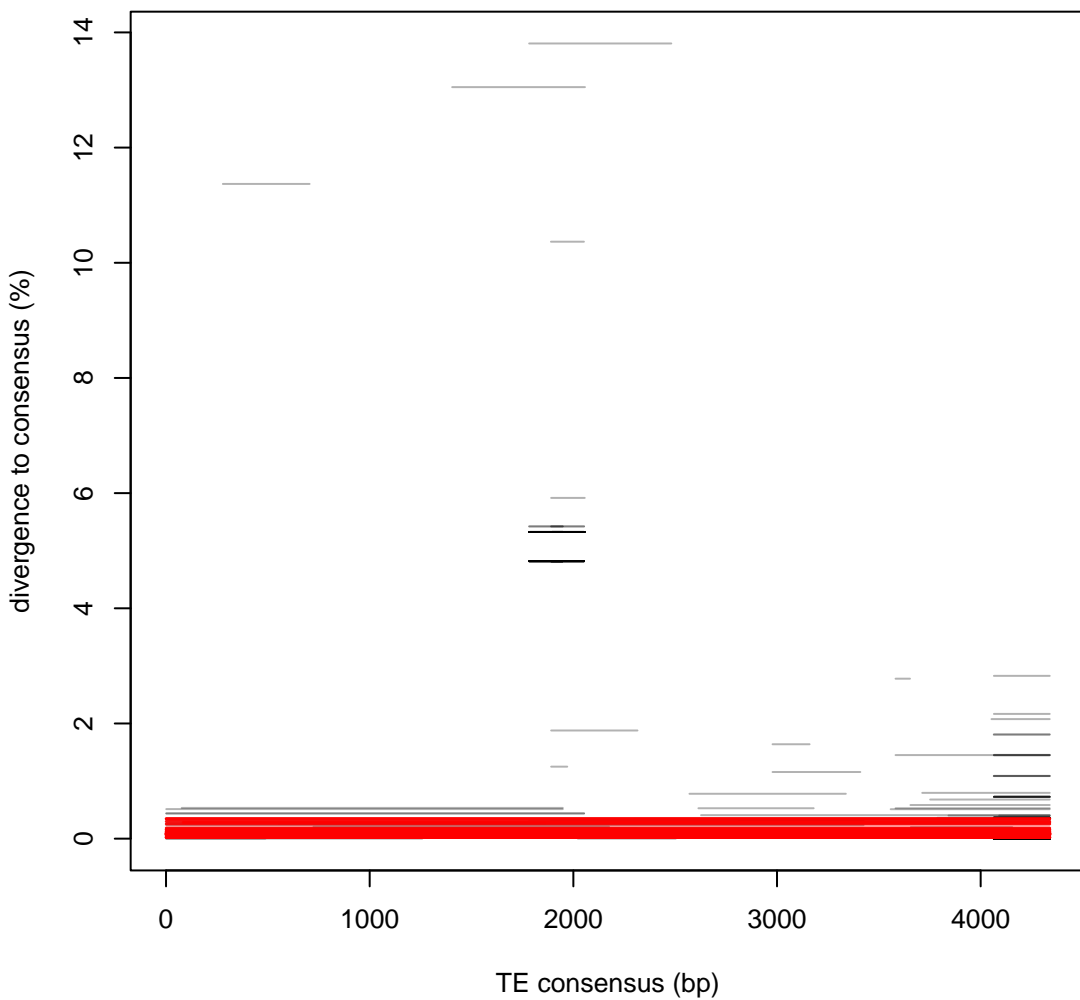
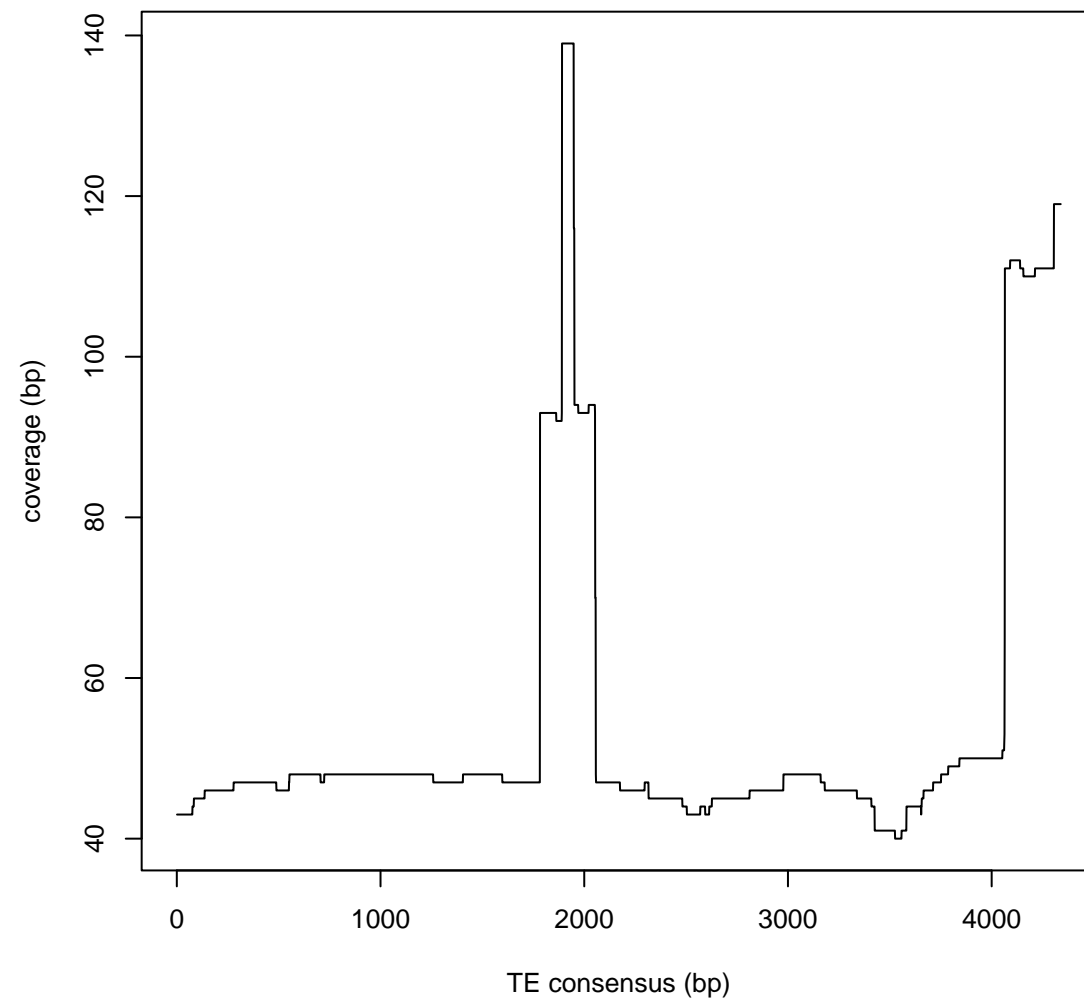


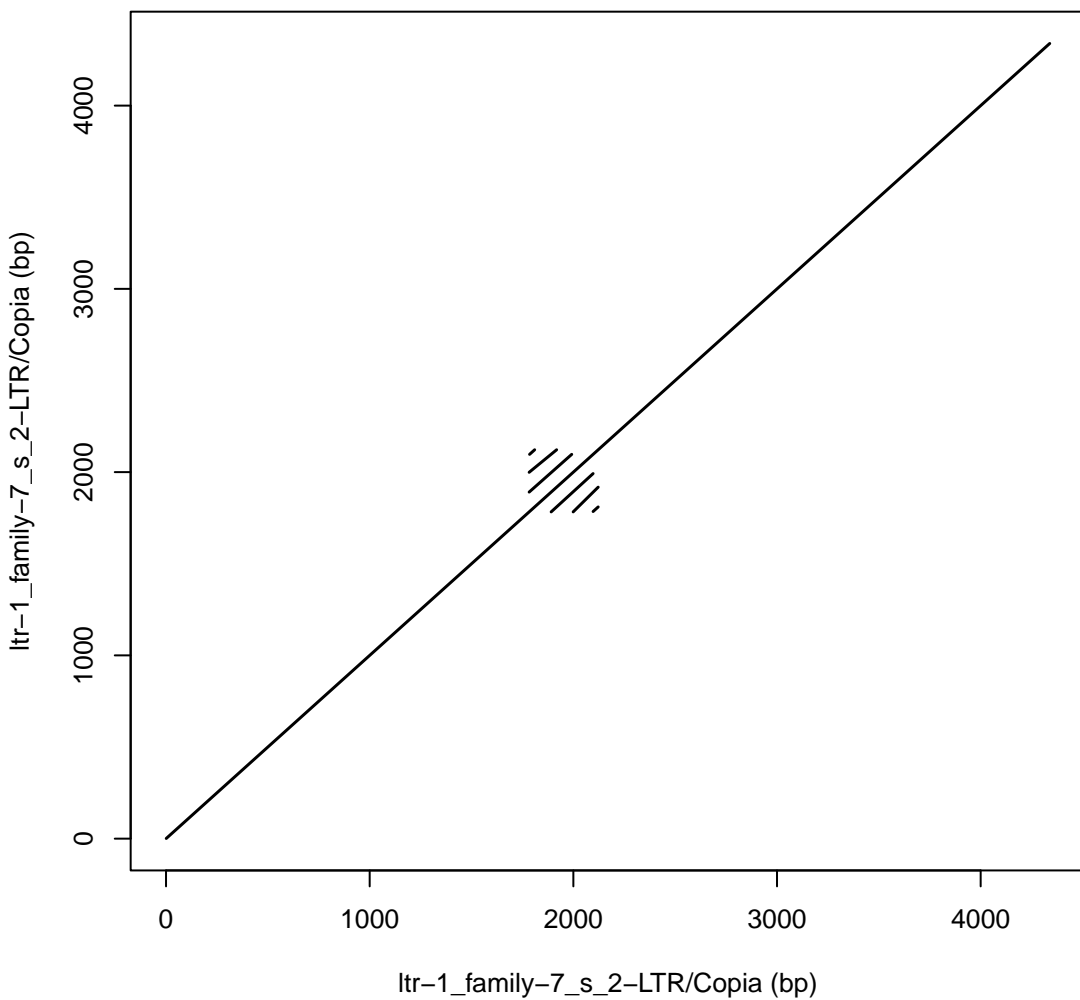
TE: ltr-1_family-7_s_2-LTR/Copia
consensus size: 4339bp; fragments: 234; full length: 34 (>=3905.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

