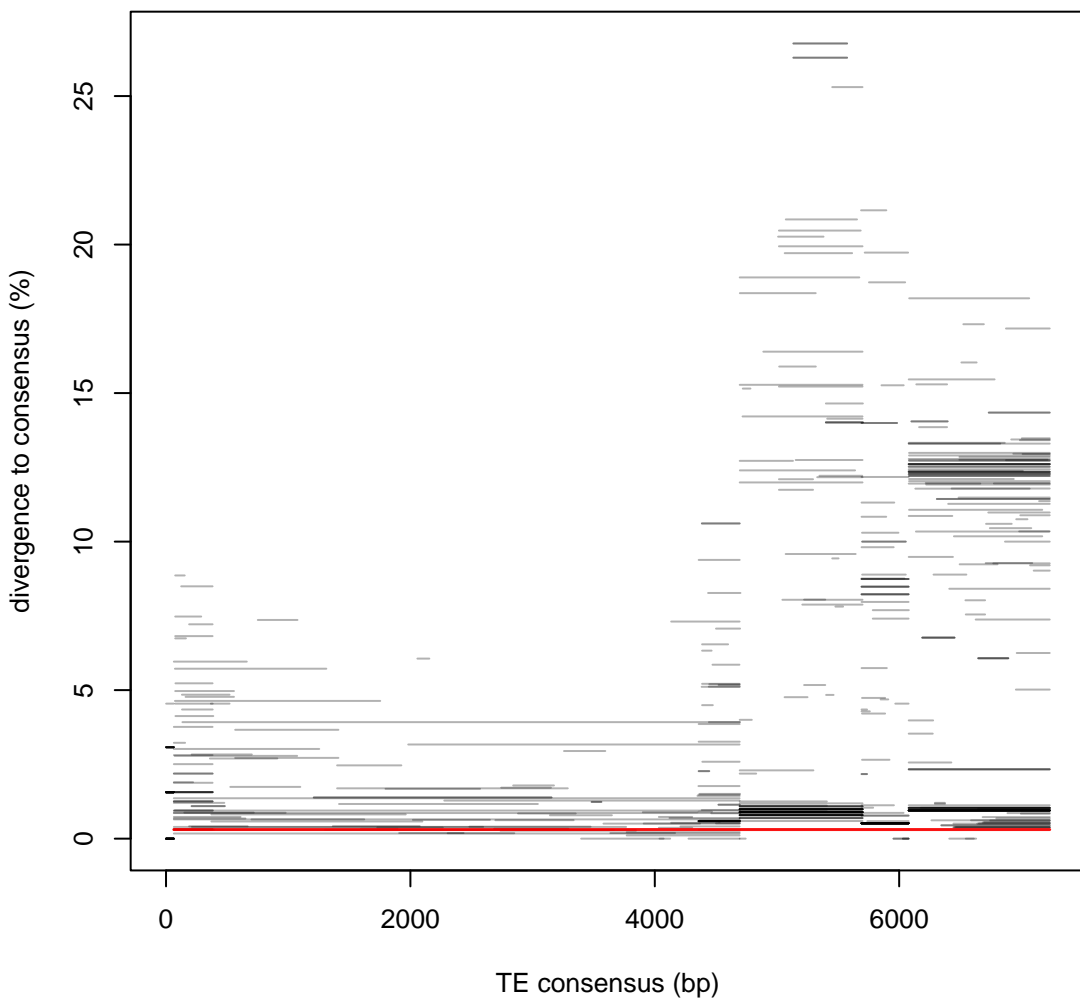
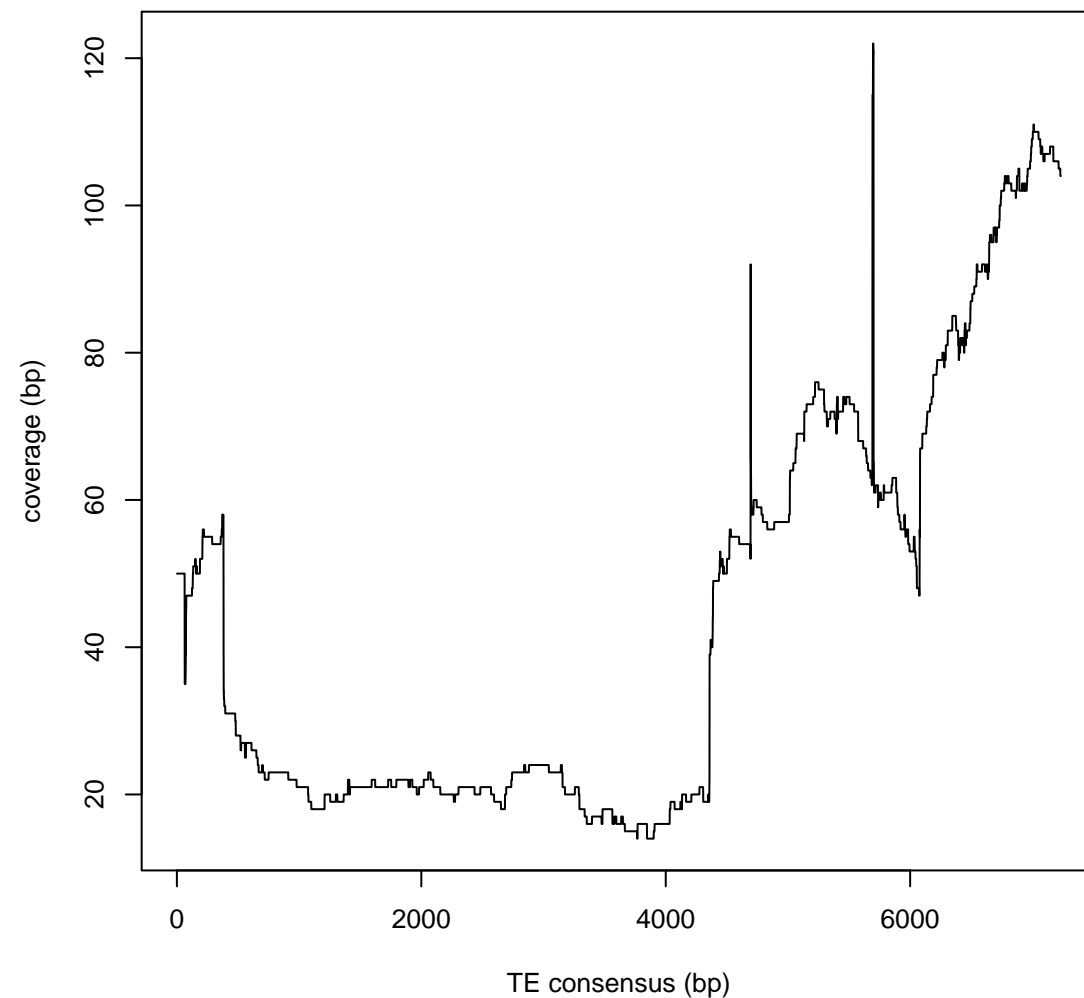


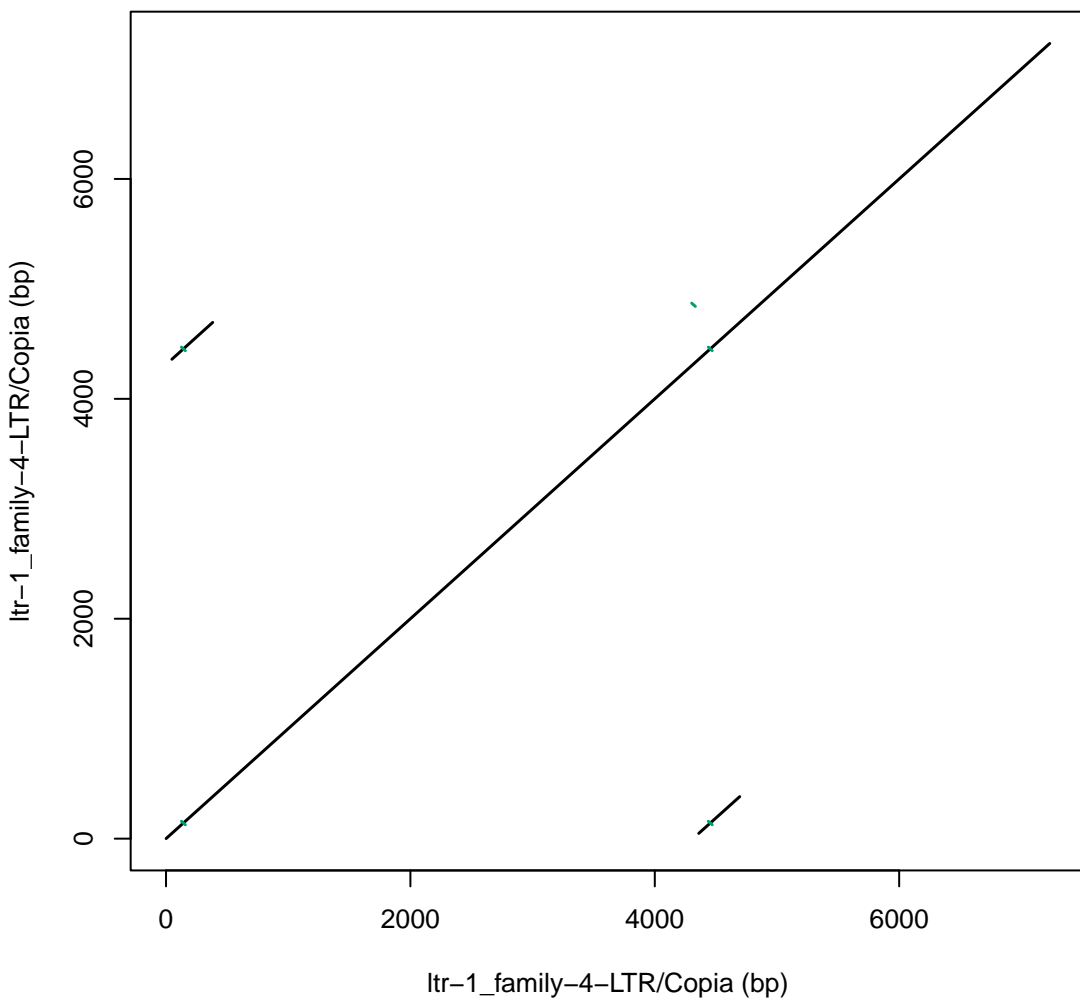
TE: Itr-1_family-4-LTR/Copia
consensus size: 7232bp; fragments: 512; full length: 1 (>=6508.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

