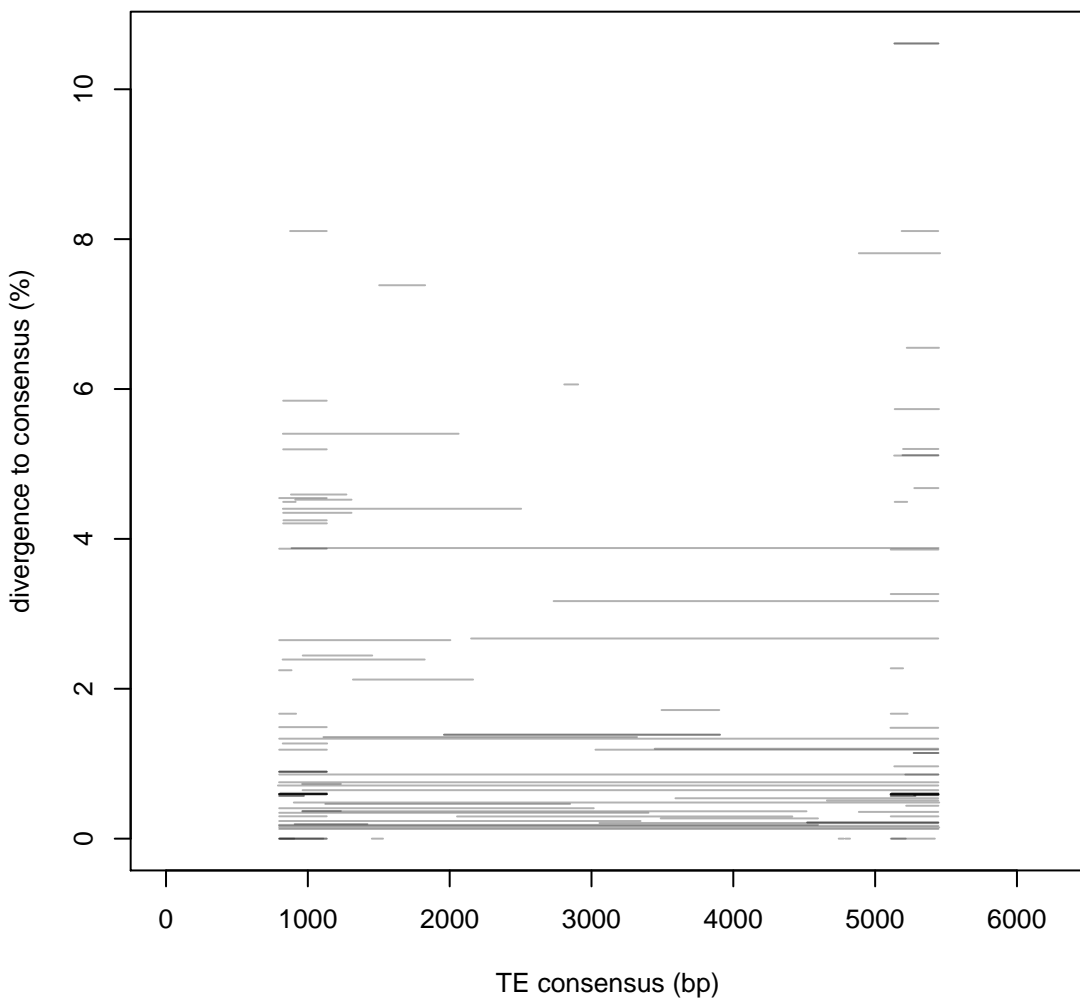
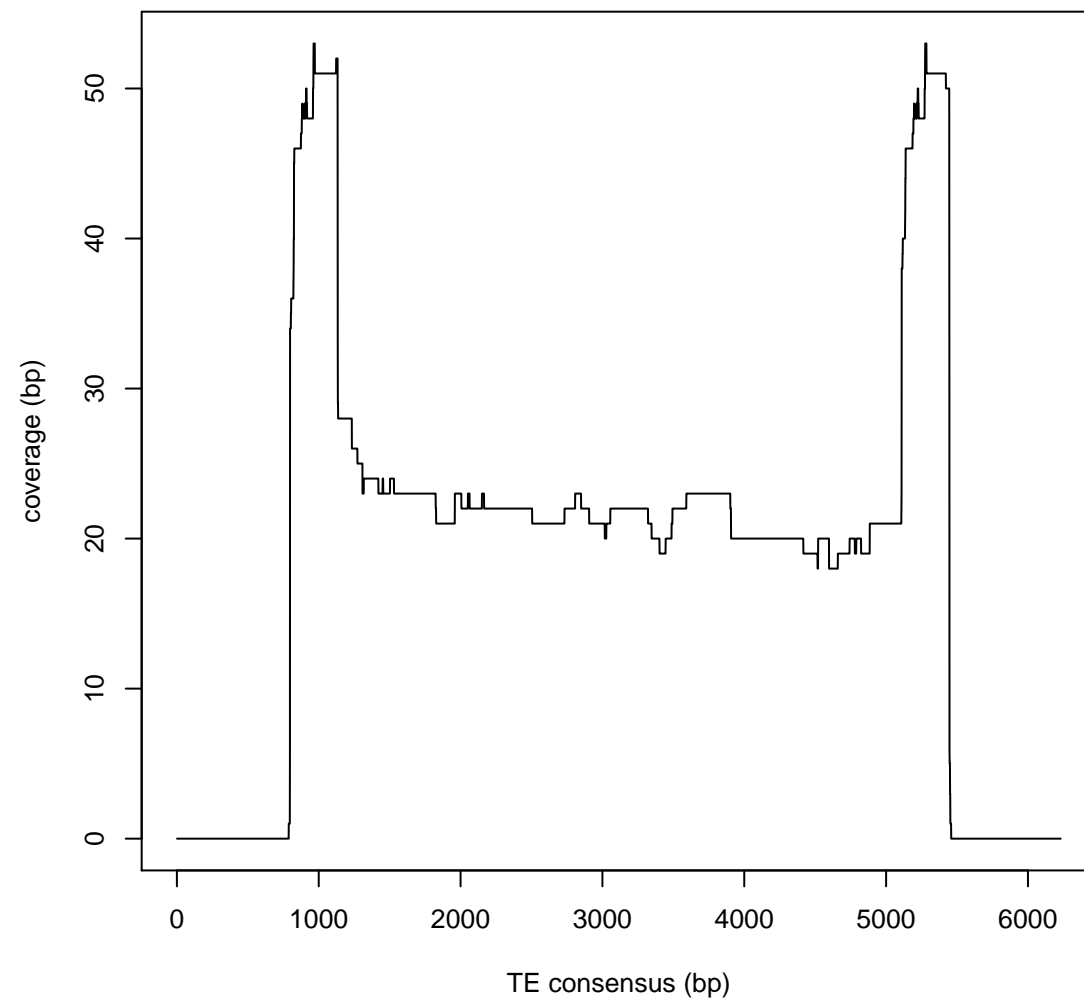


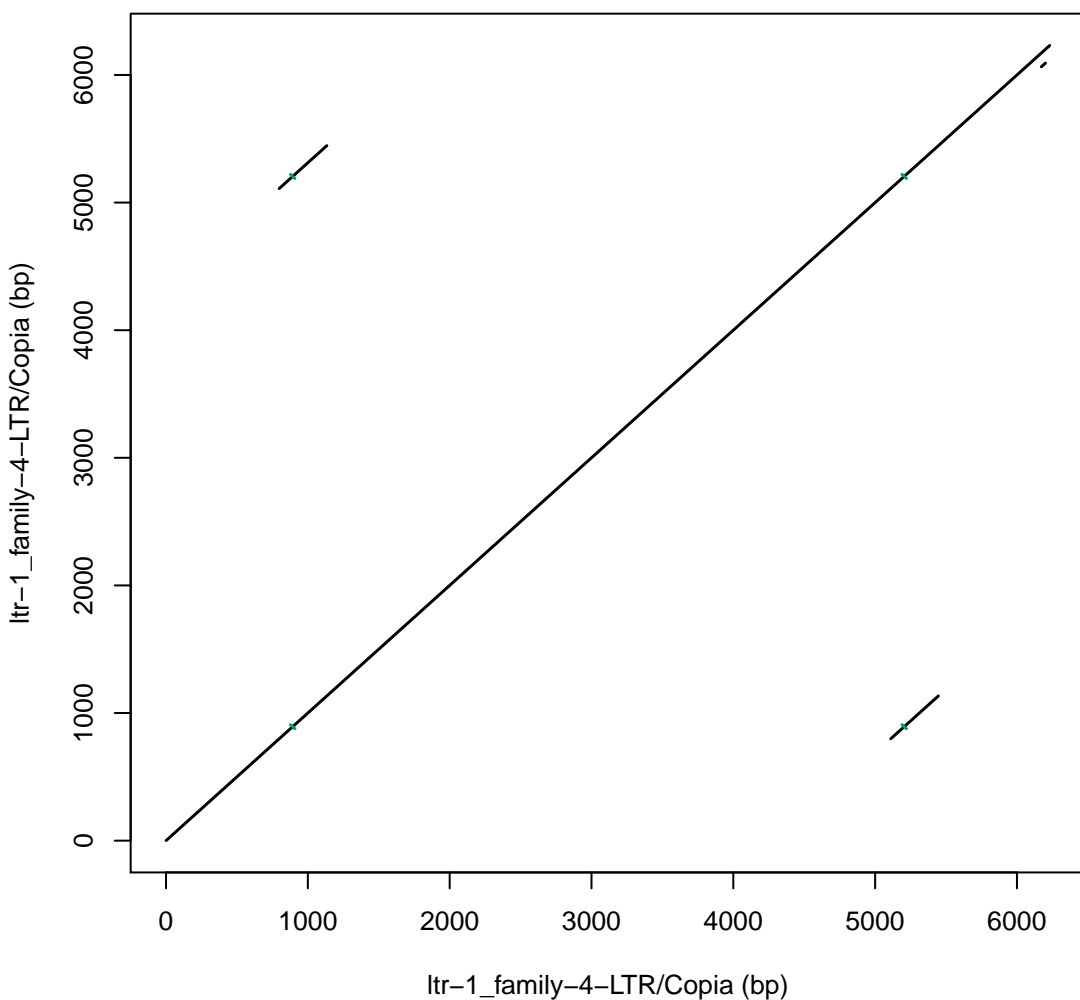
TE: Itr-1_family-4-LTR/Copia
consensus size: 6231bp; fragments: 117; full length: 0 (>=5607.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

