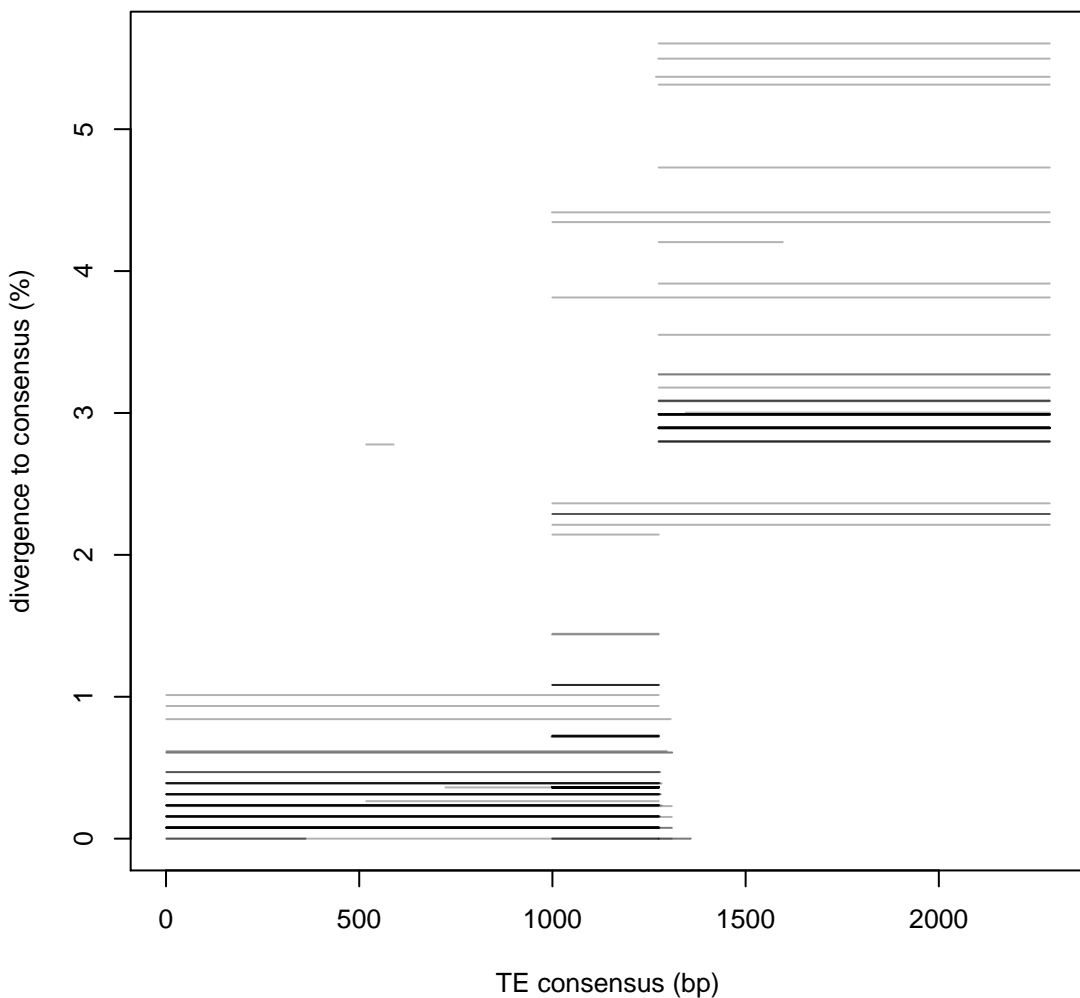
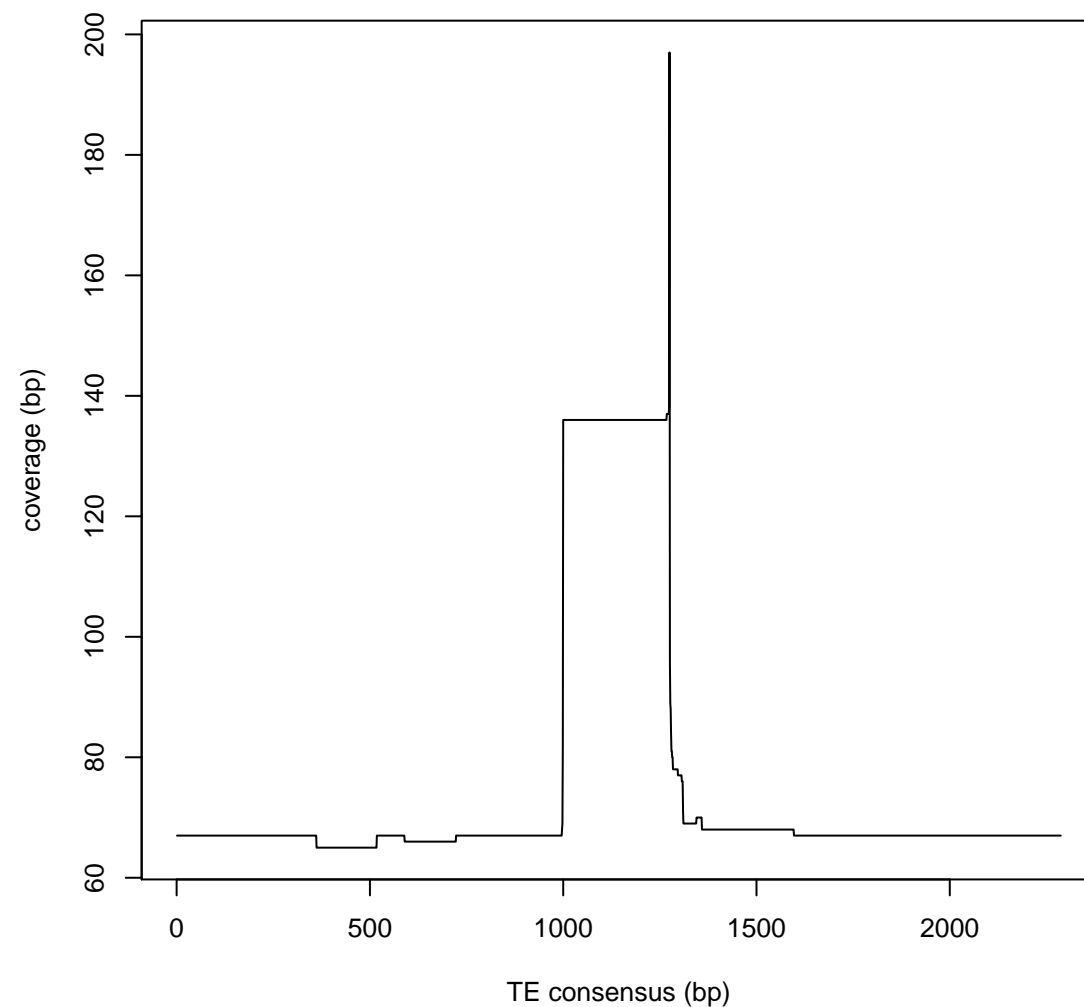


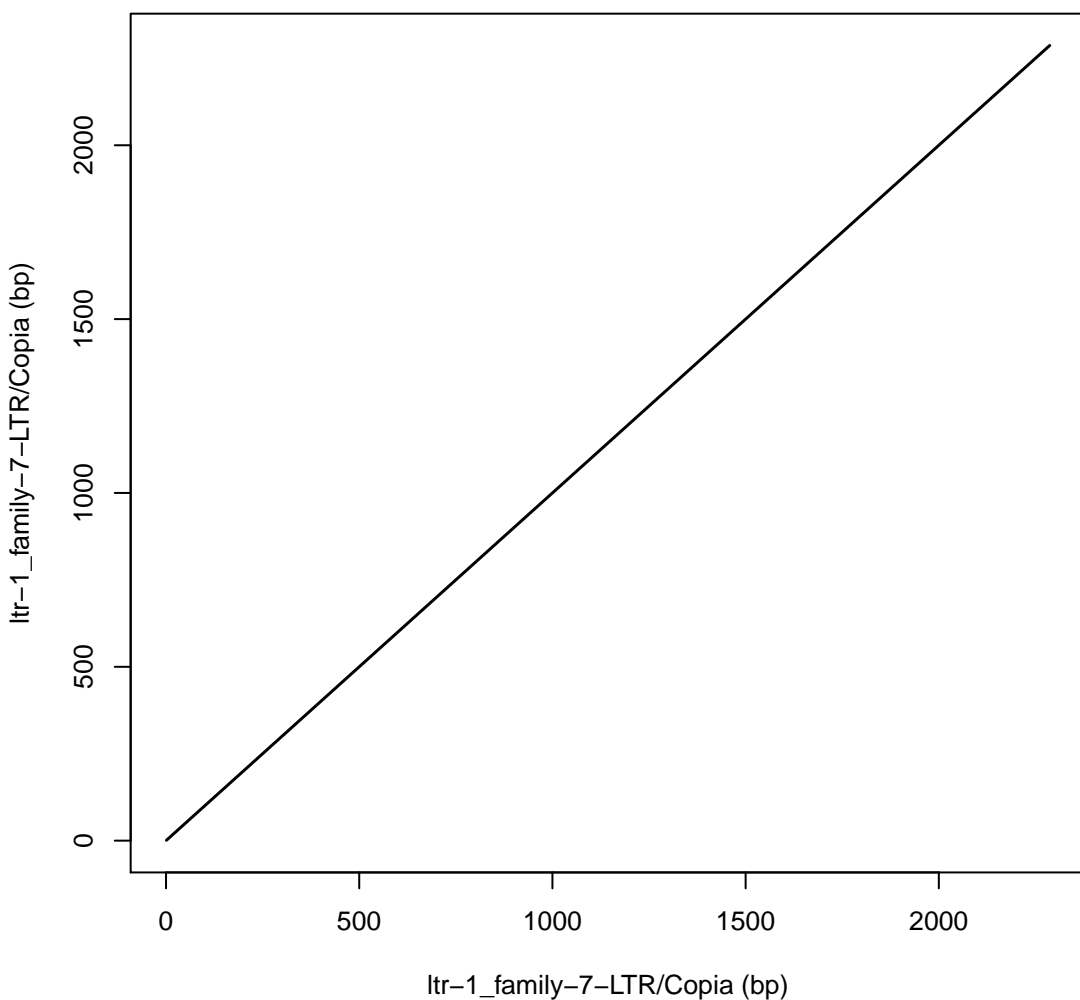
TE: Itr-1_family-7-LTR/Copia
consensus size: 2287bp; fragments: 201; full length: 0 (>=2058.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

