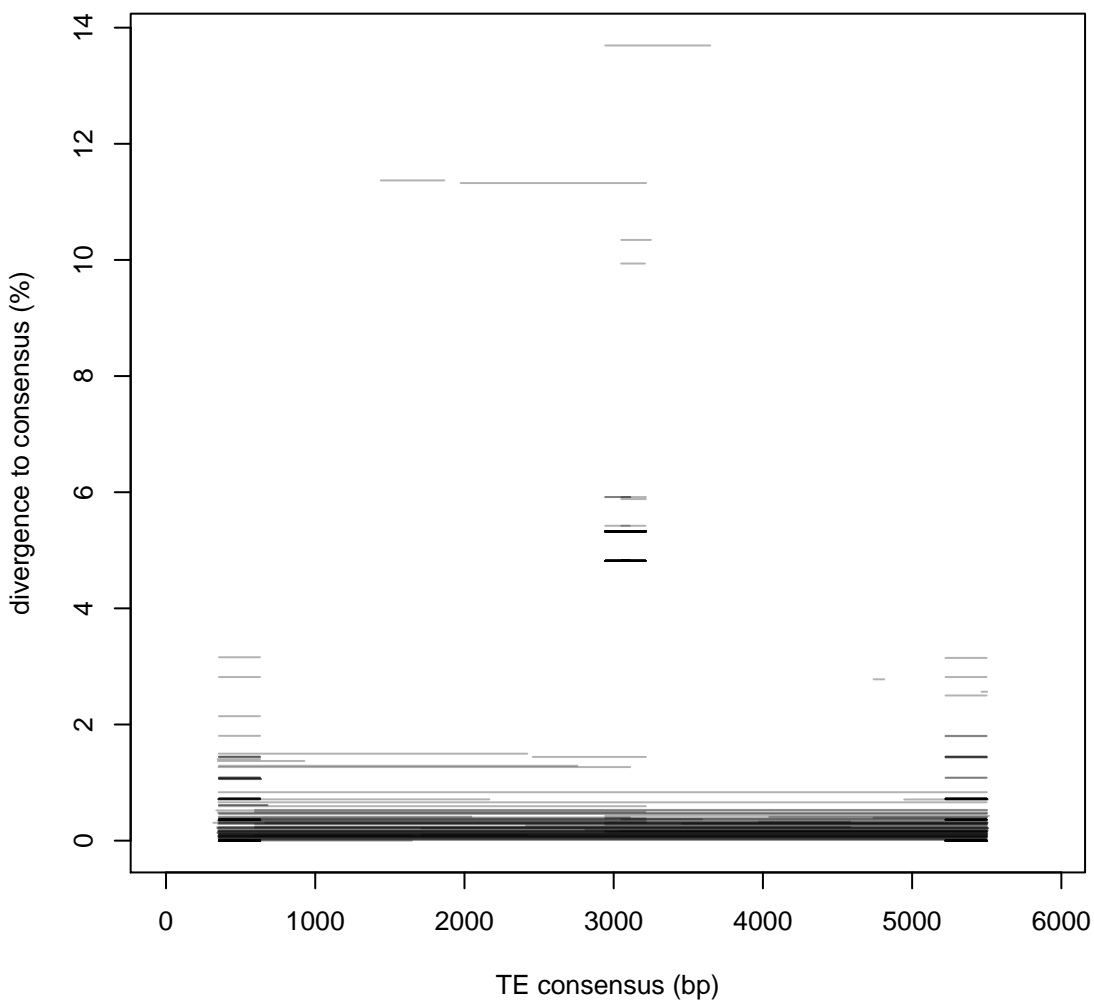
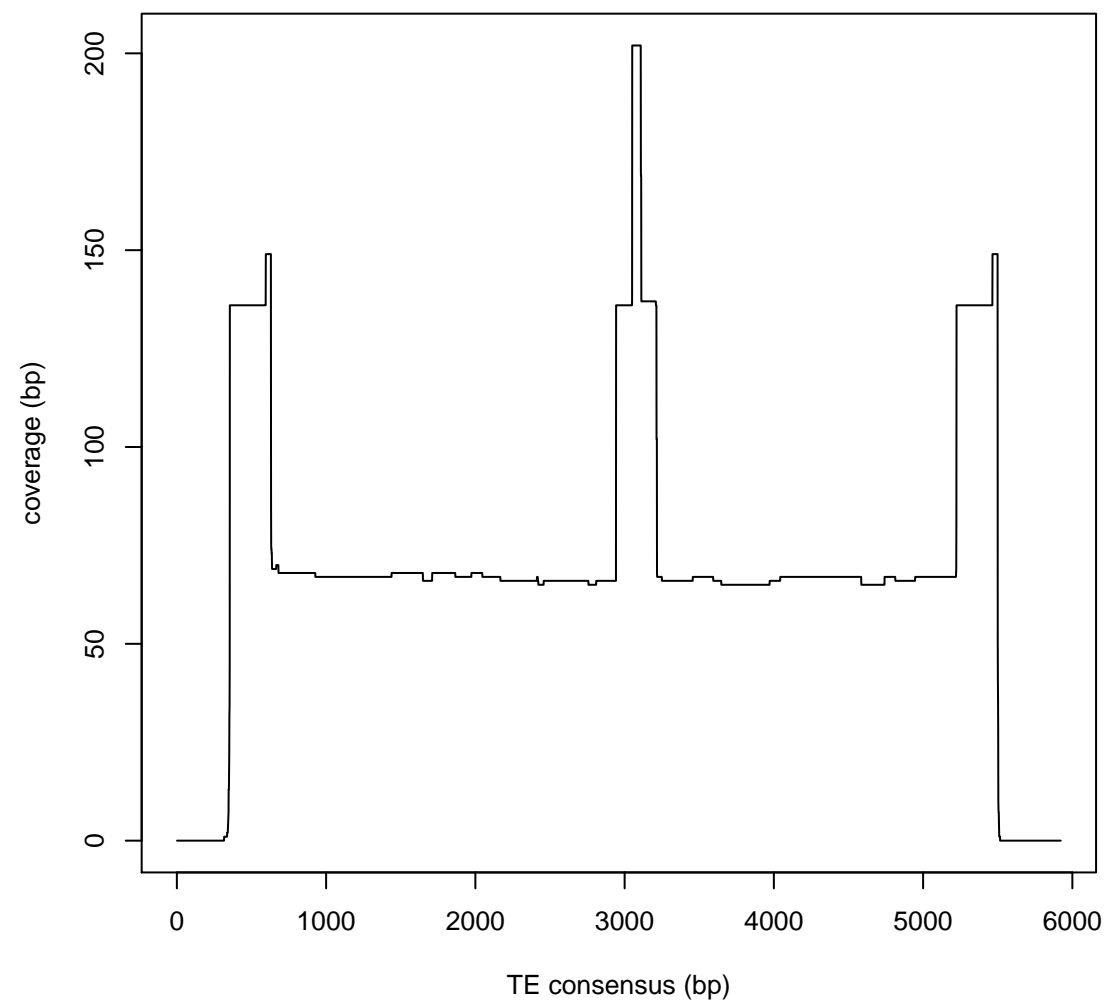


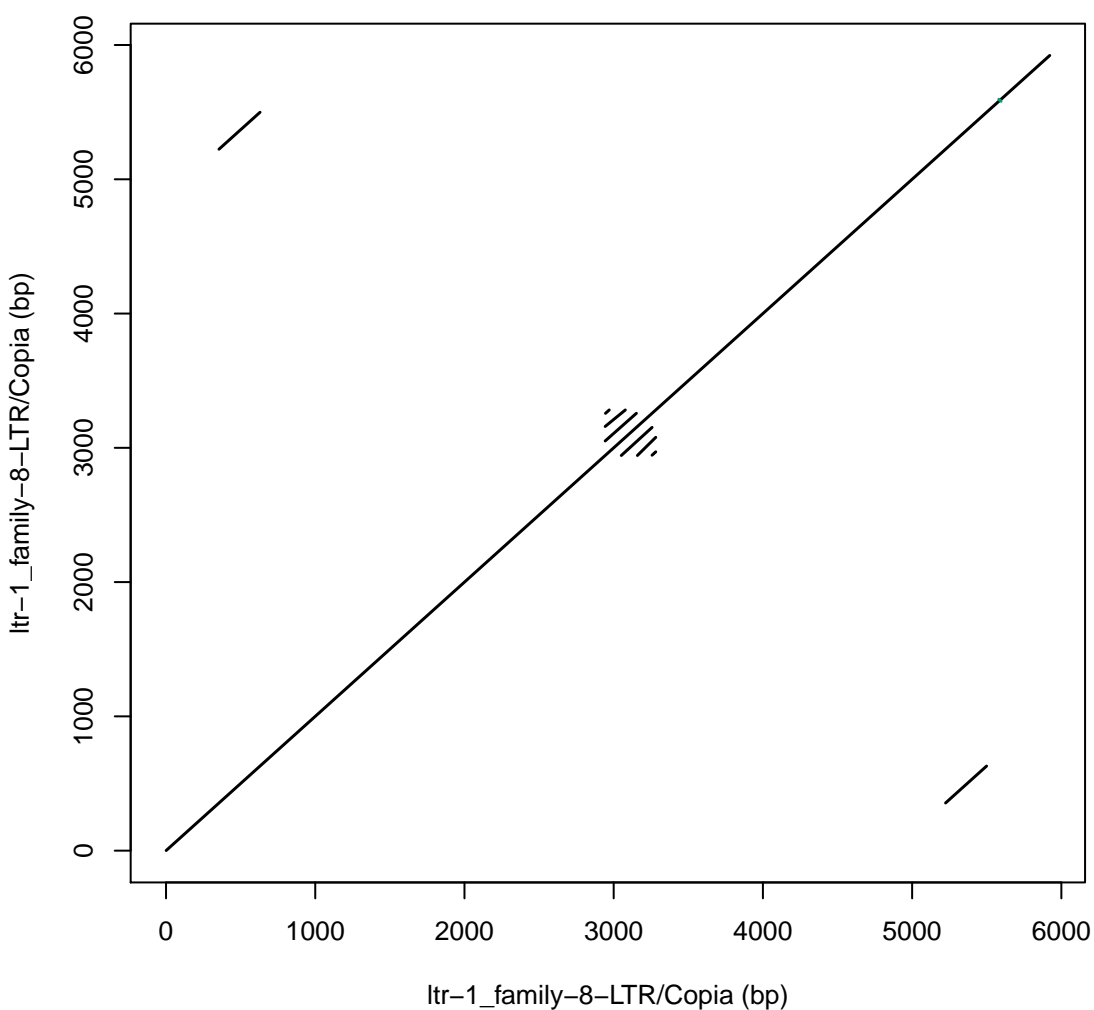
TE: Itr-1_family-8-LTR/Copia
consensus size: 5922bp; fragments: 381; full length: 0 (>=5329.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

