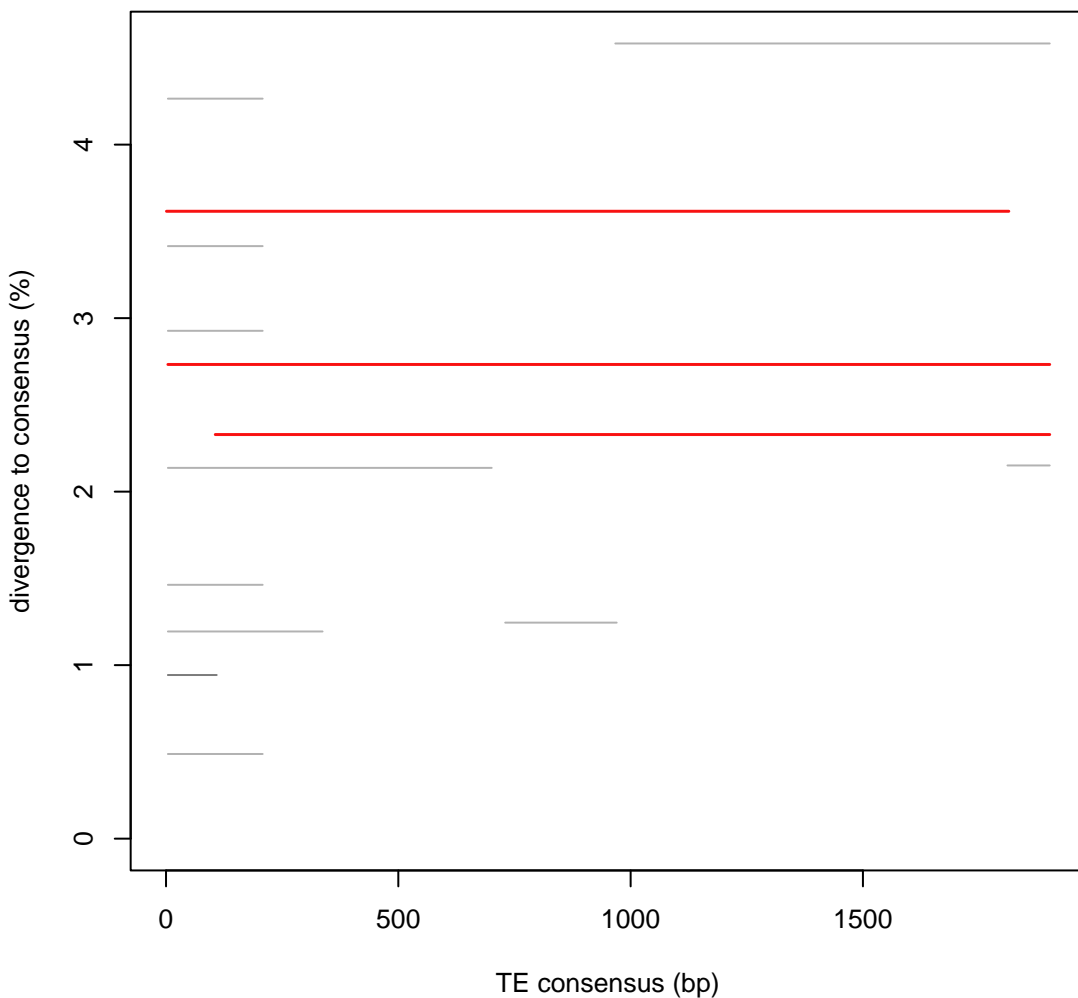
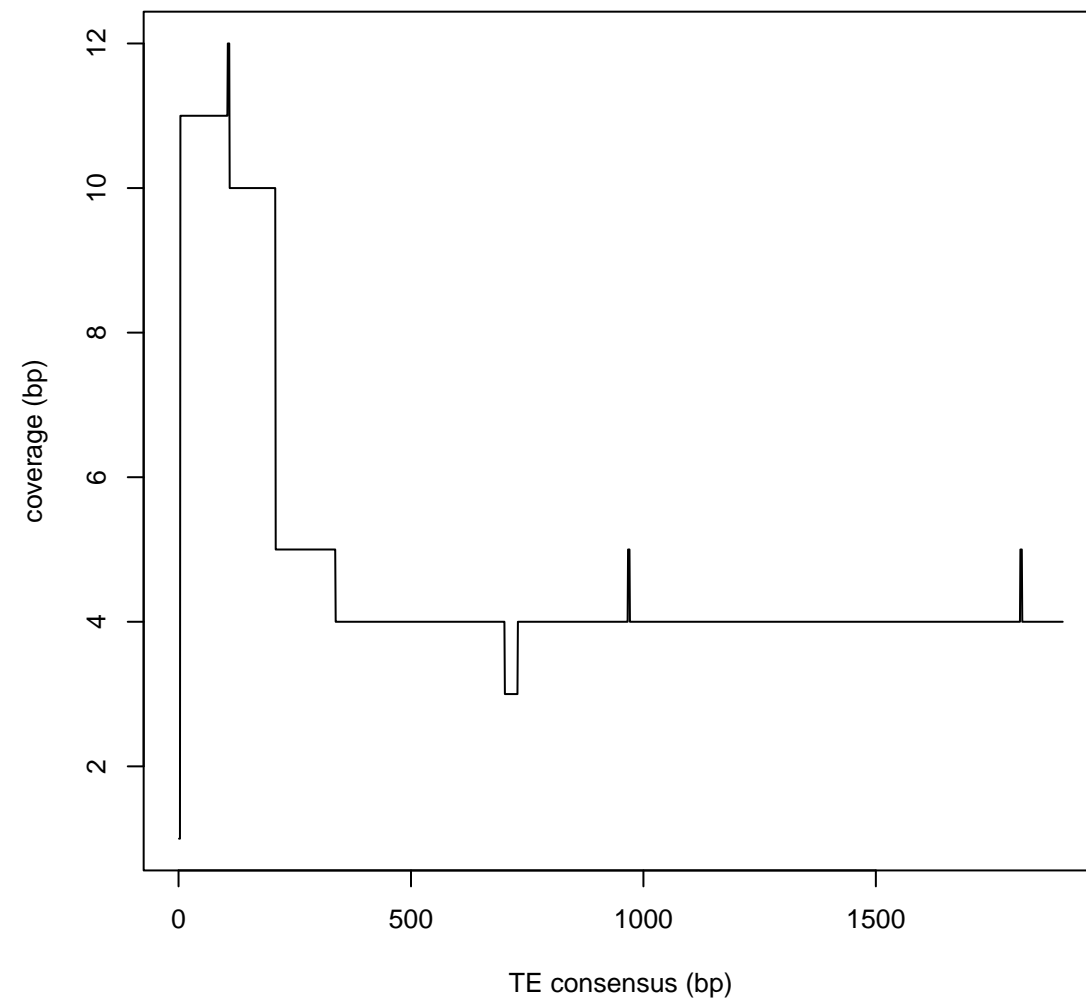


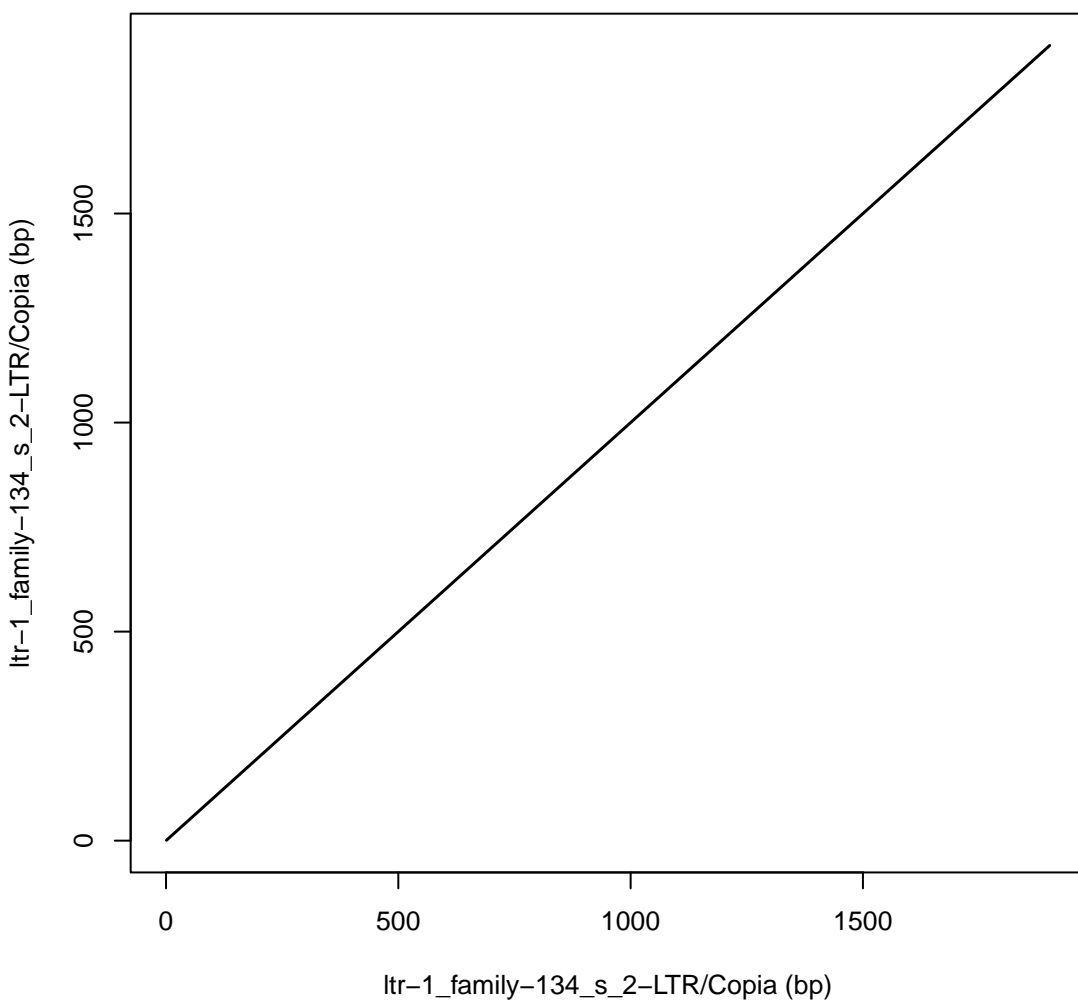
TE: ltr-1_family-134_s_2-LTR/Copia
consensus size: 1902bp; fragments: 15; full length: 3 (>=1711.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

