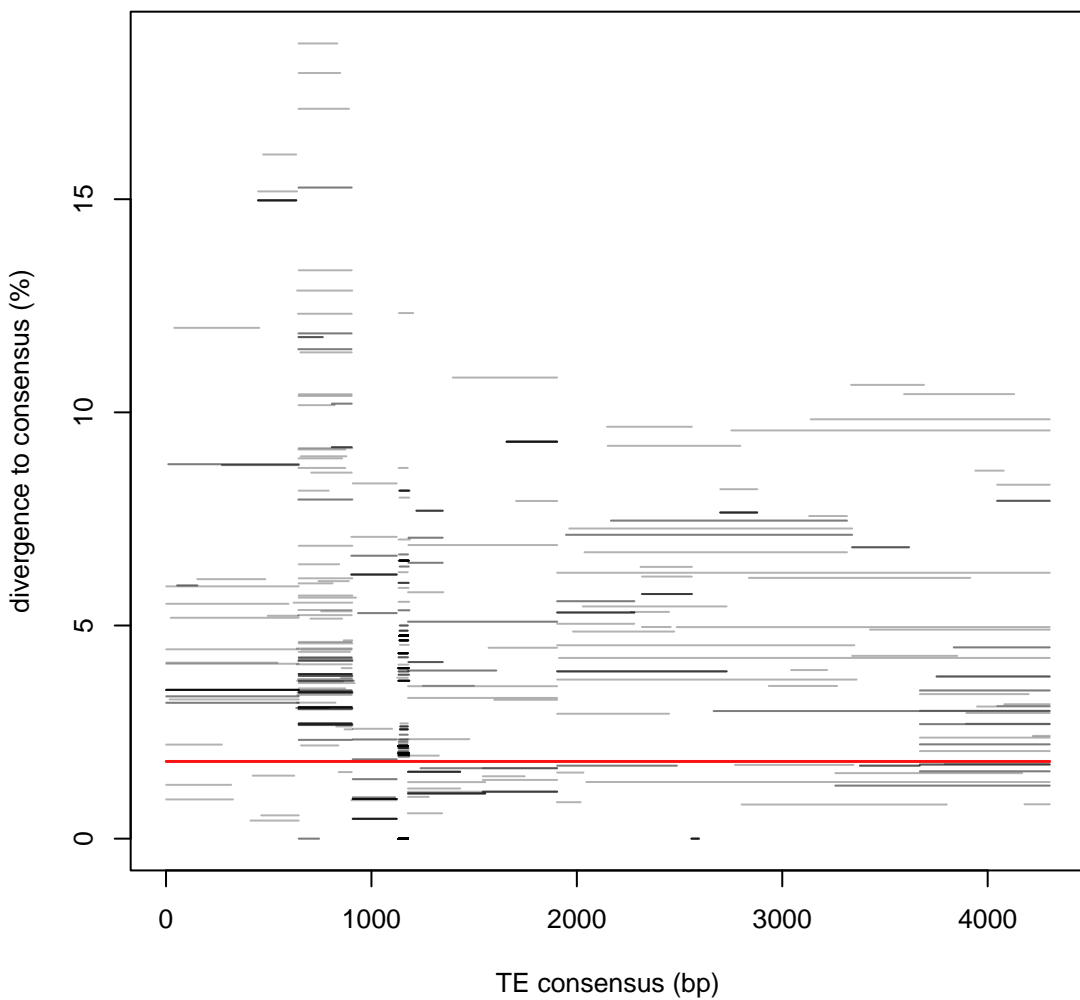
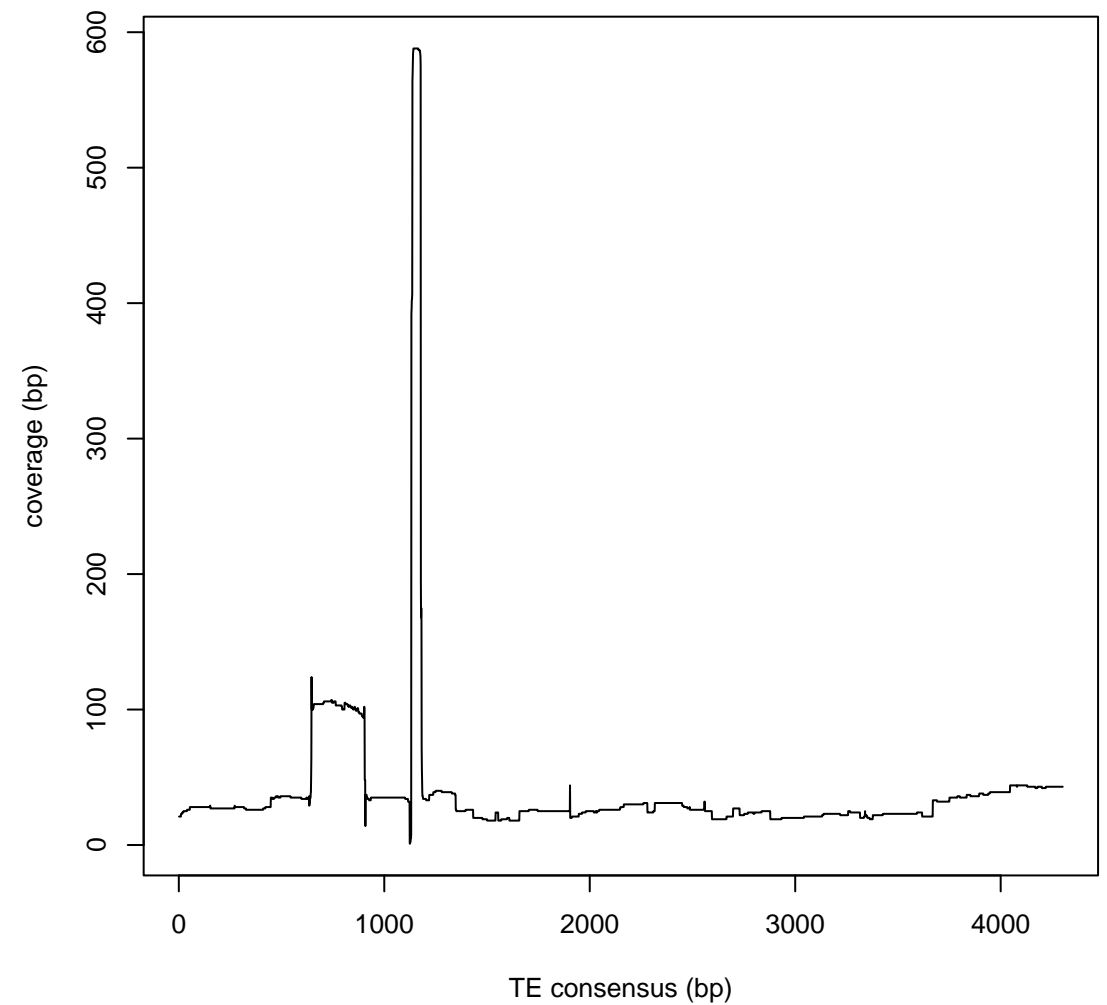


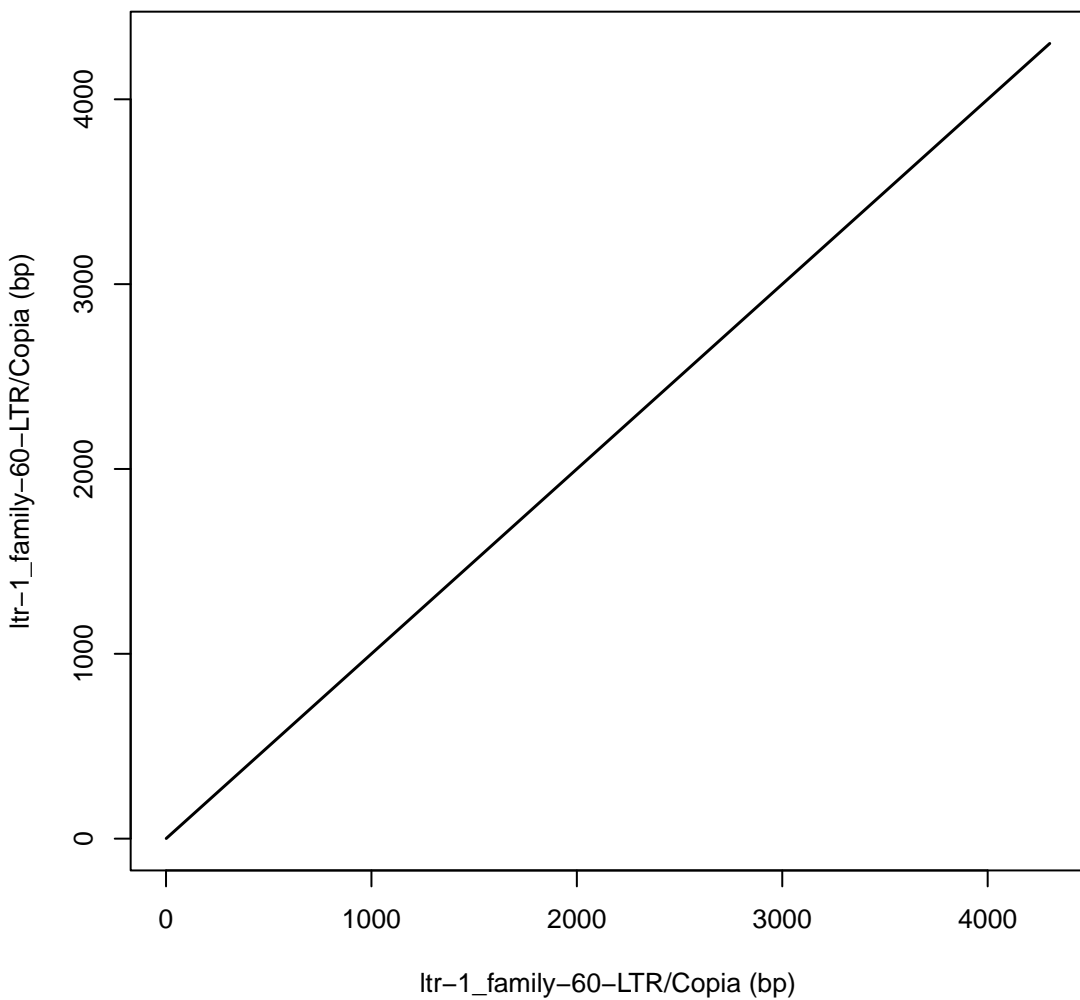
TE: ltr-1_family-60-LTR/Copia
consensus size: 4302bp; fragments: 946; full length: 1 (>=3871.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

