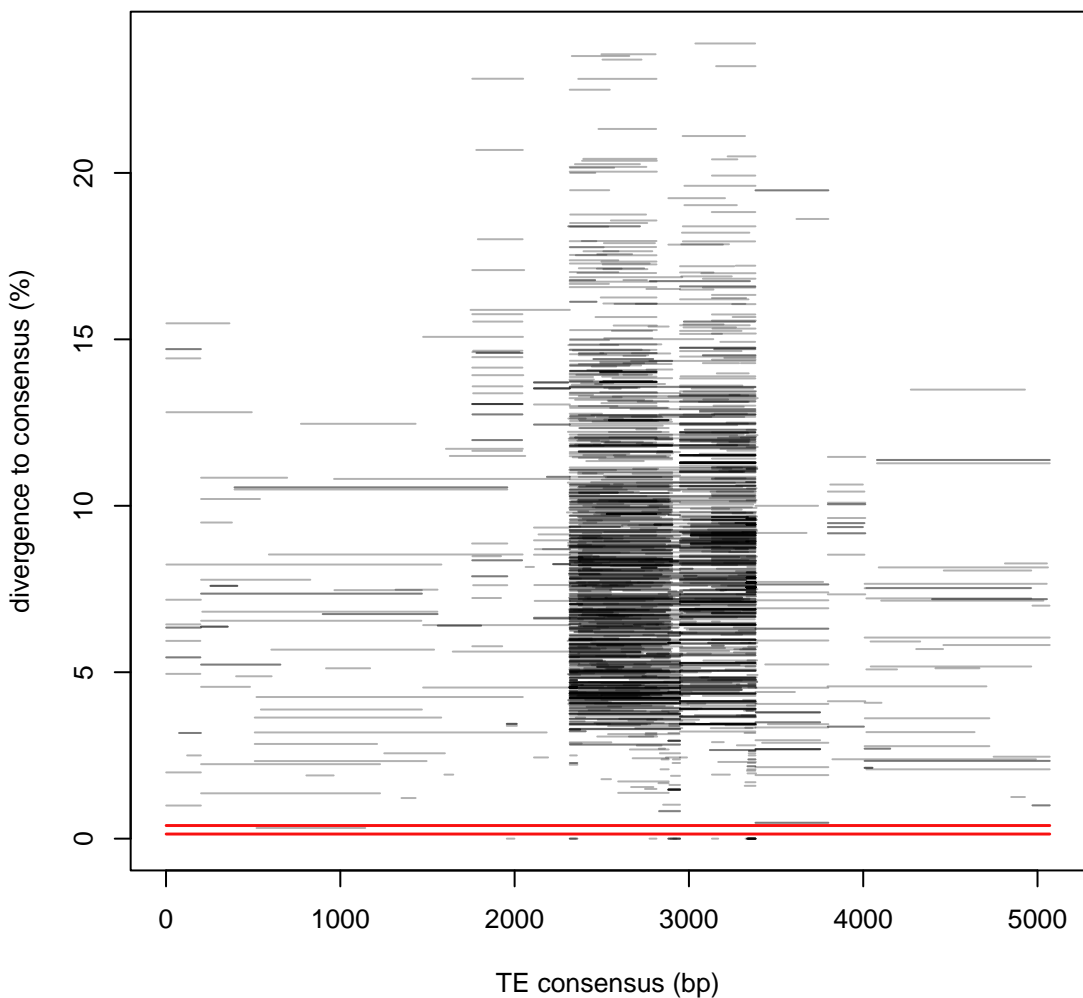
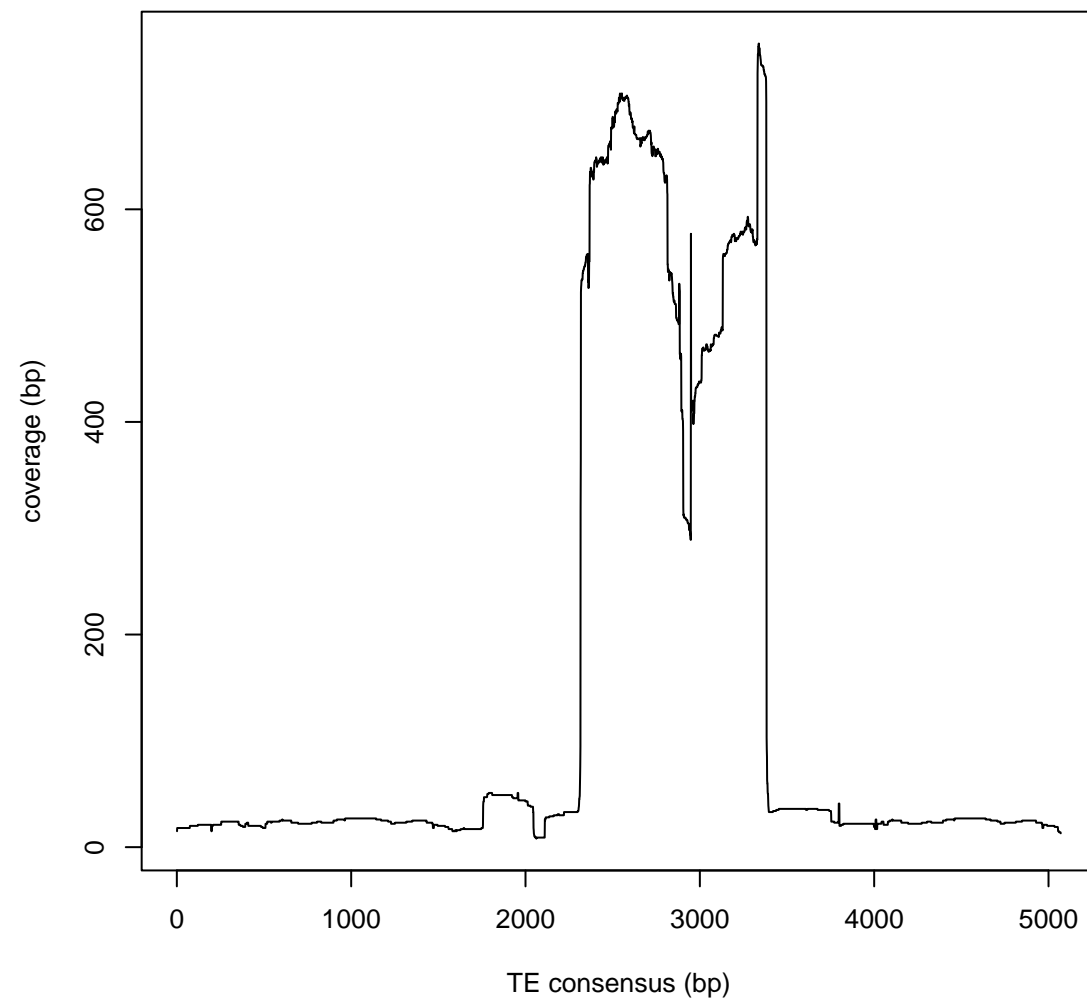


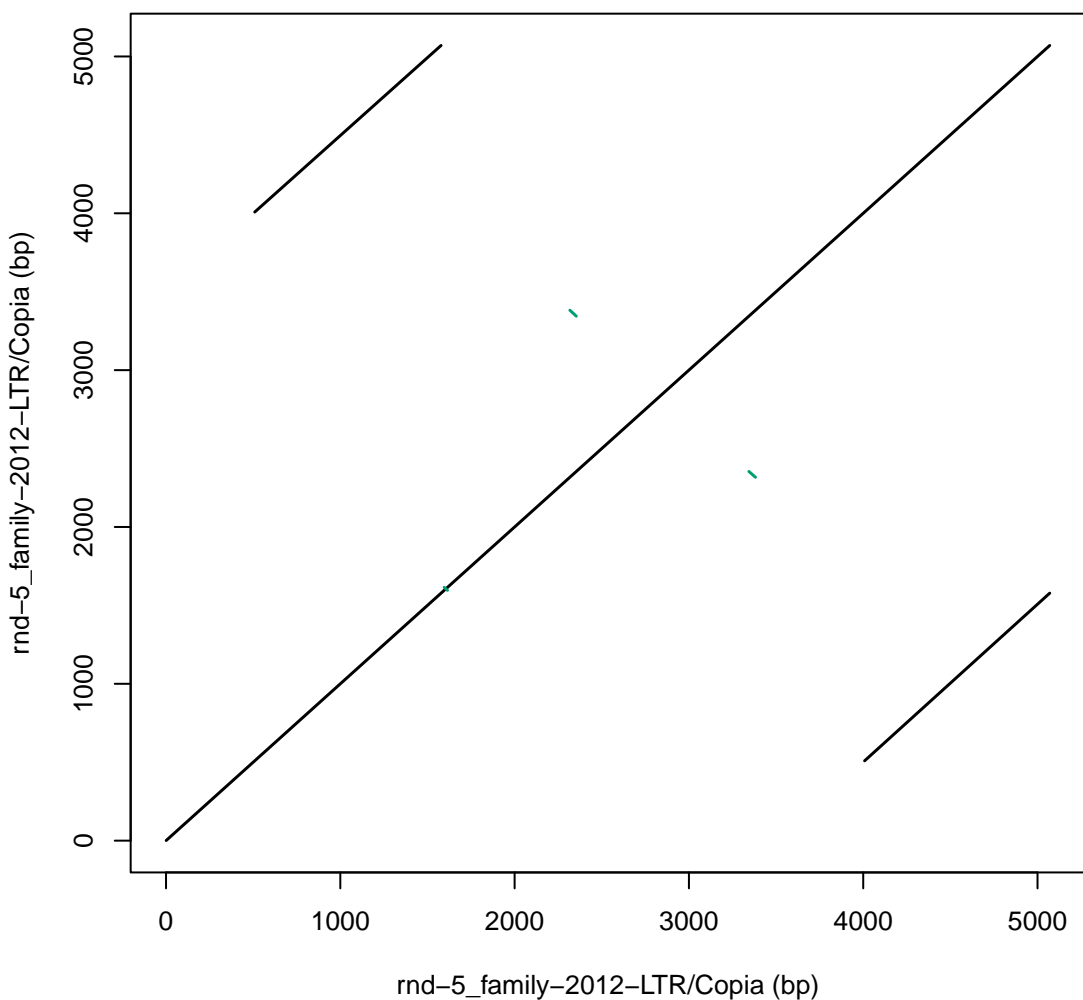
TE: rnd-5\_family-2012-LTR/Copia  
consensus size: 5070bp; fragments: 2085; full length: 2 (>=4563bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

