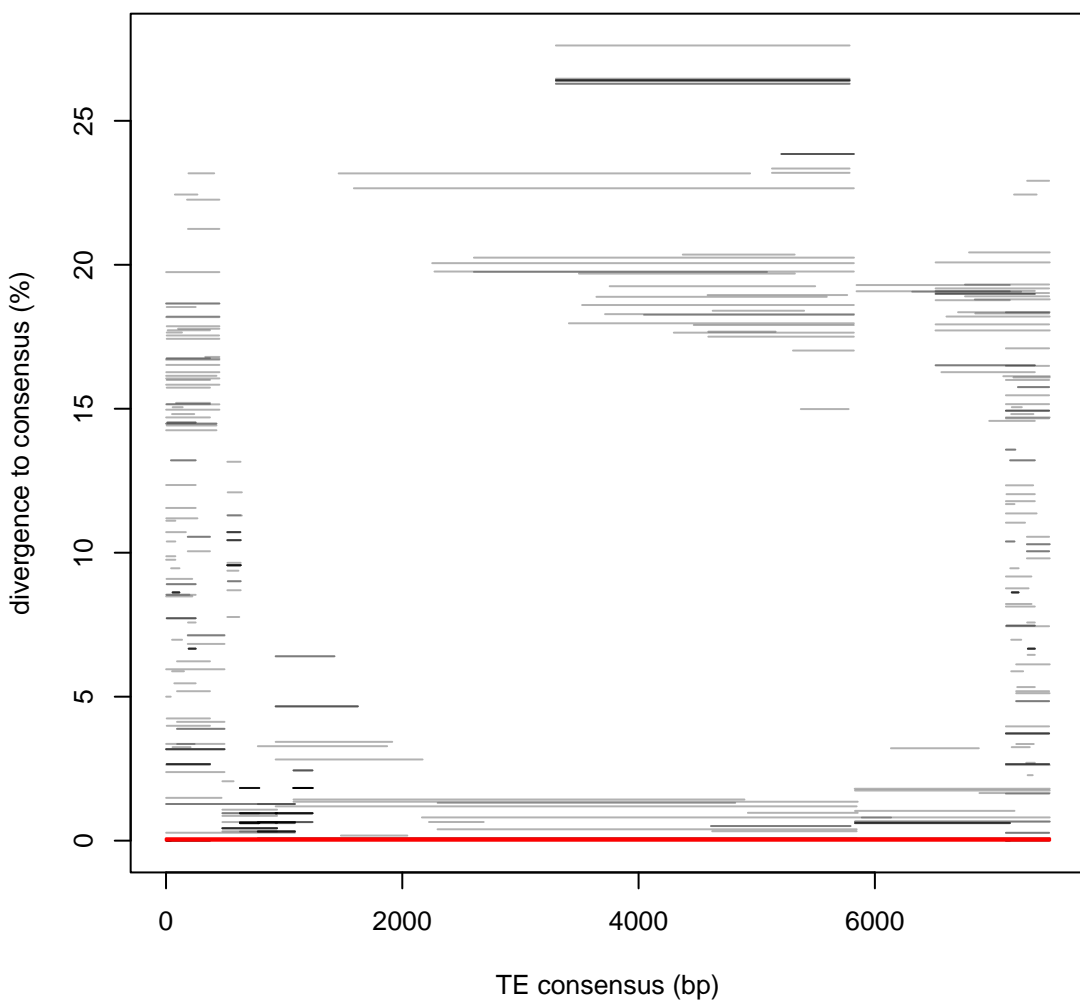
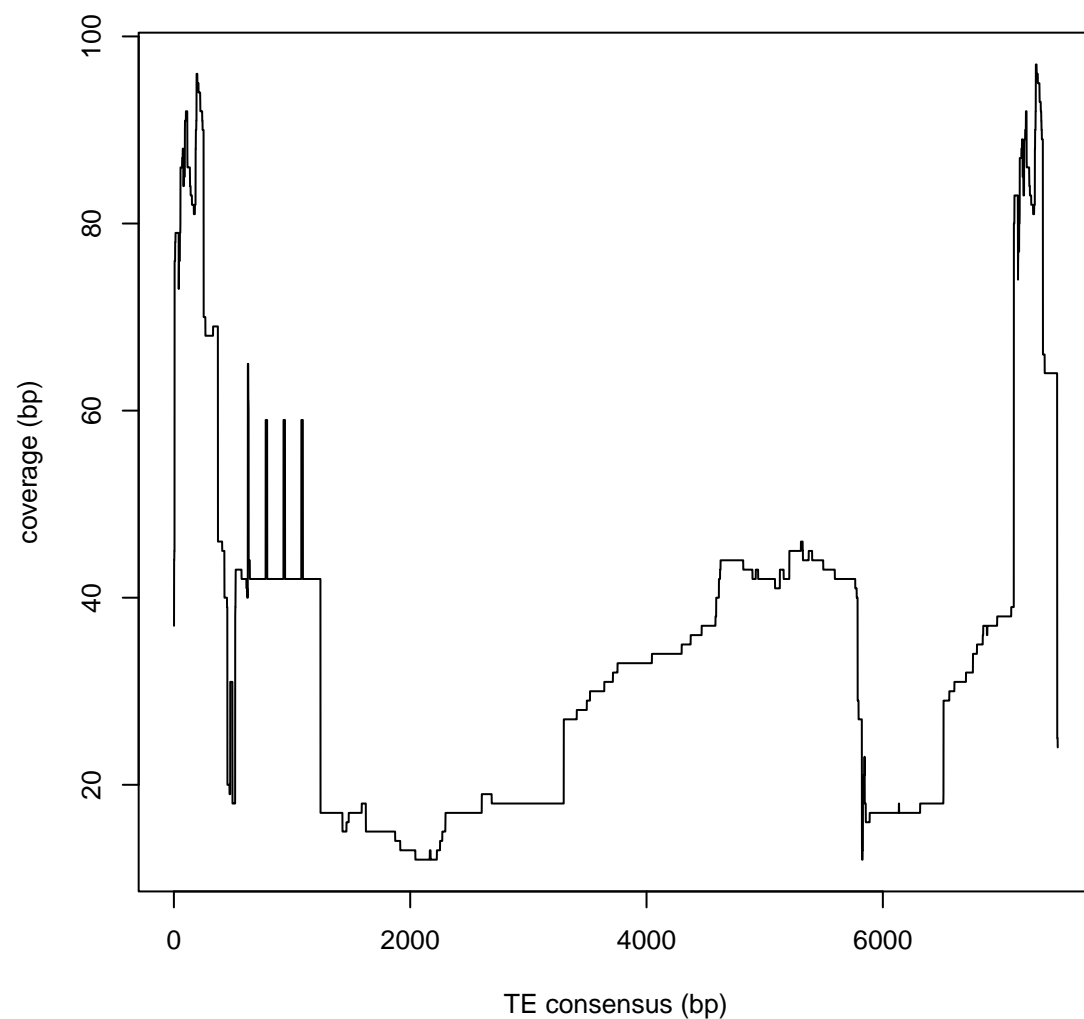


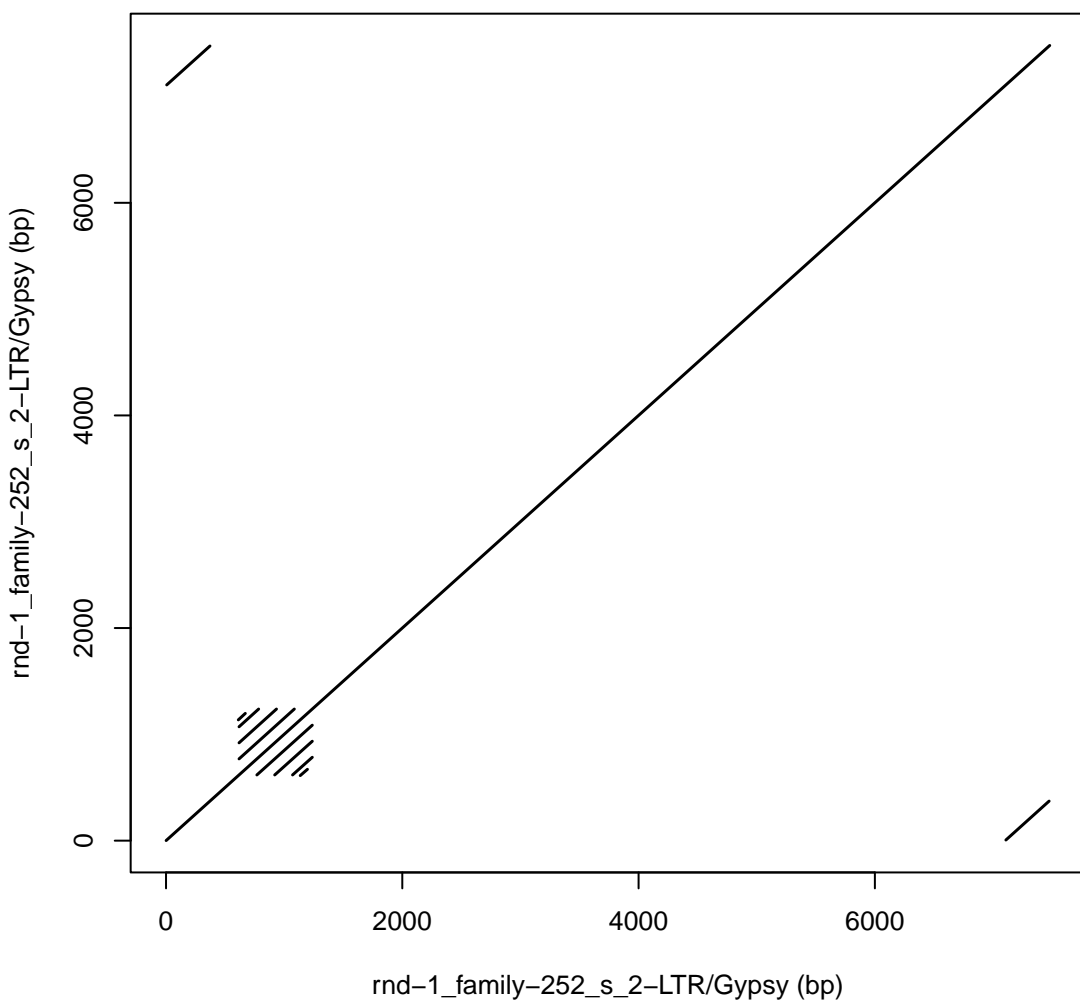
TE: rnd-1_family-252_s_2-LTR/Gypsy
consensus size: 7480bp; fragments: 396; full length: 3 (>=6732bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

