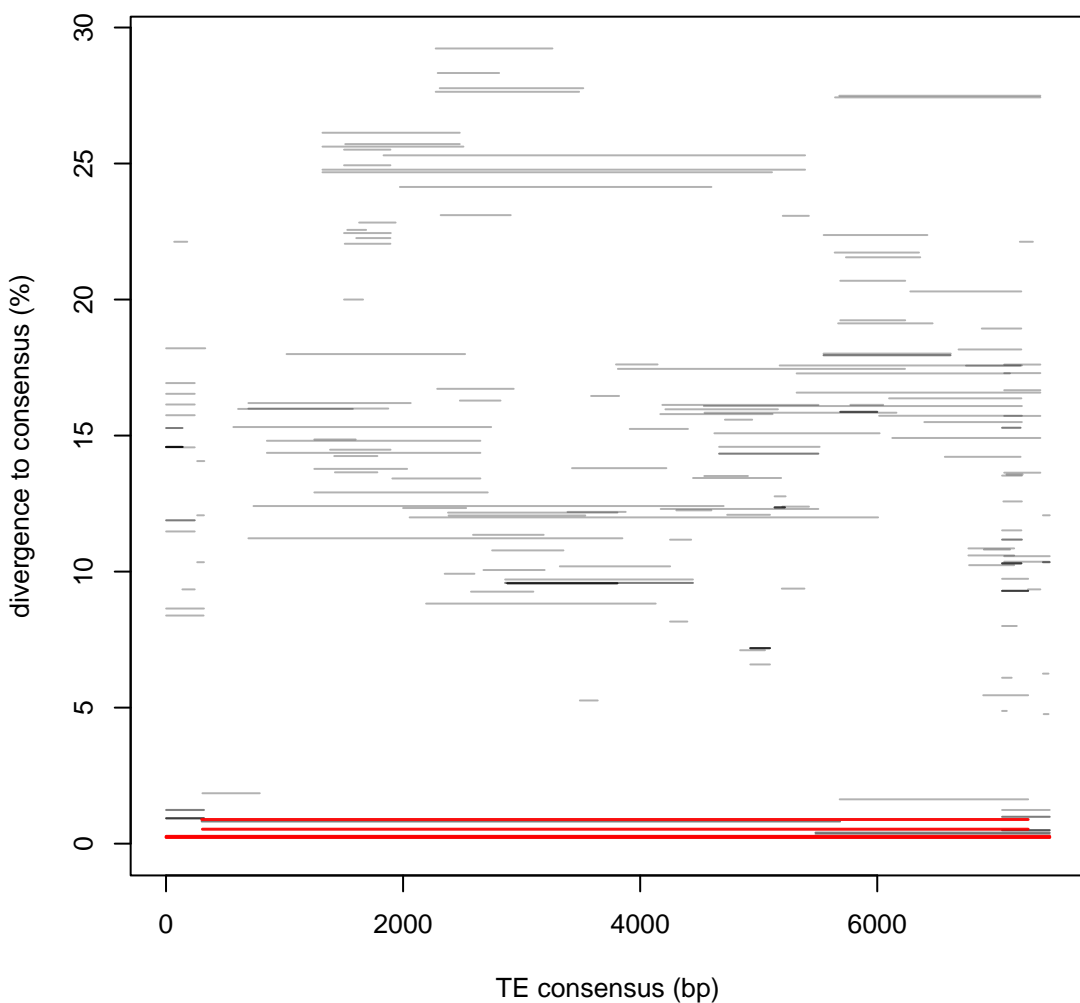
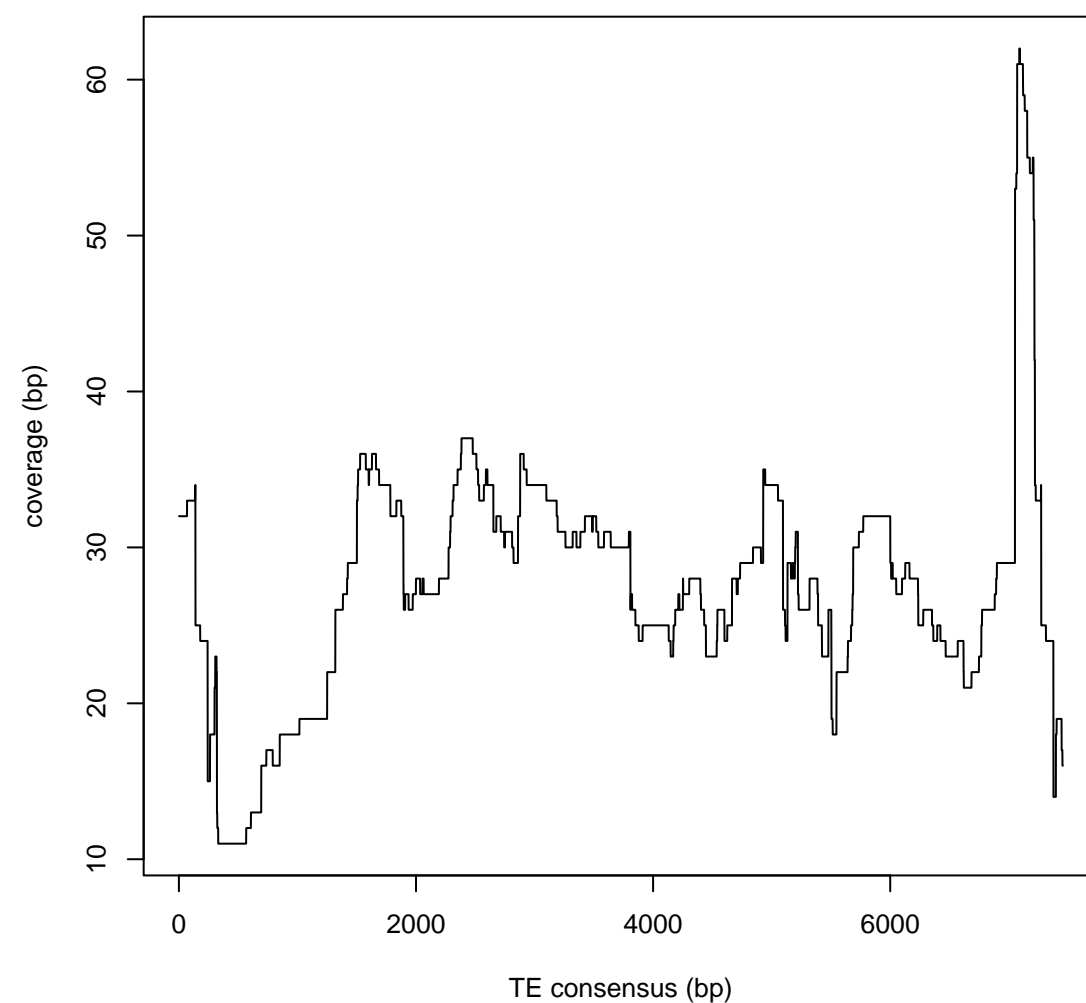


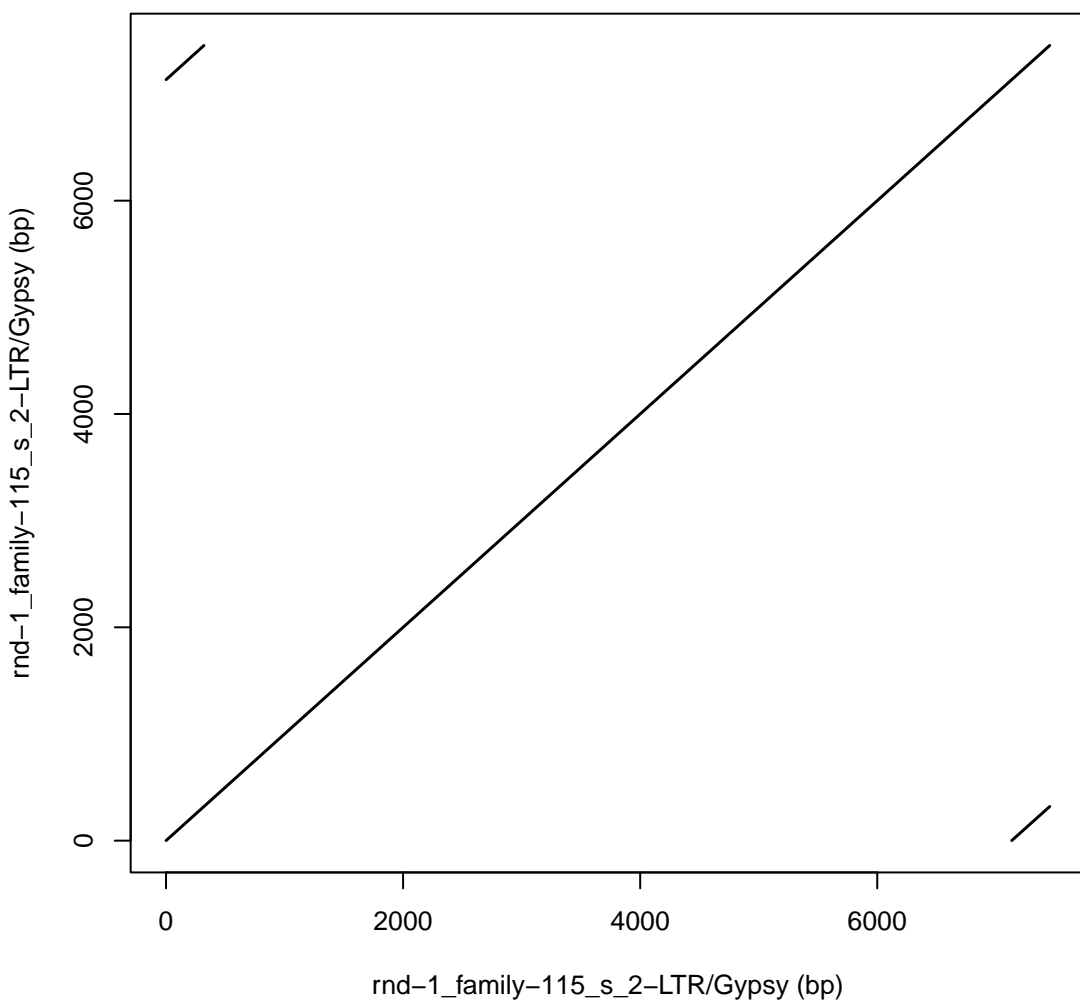
TE: rnd-1_family-115_s_2-LTR/Gypsy
consensus size: 7455bp; fragments: 213; full length: 5 (≥ 6709.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

