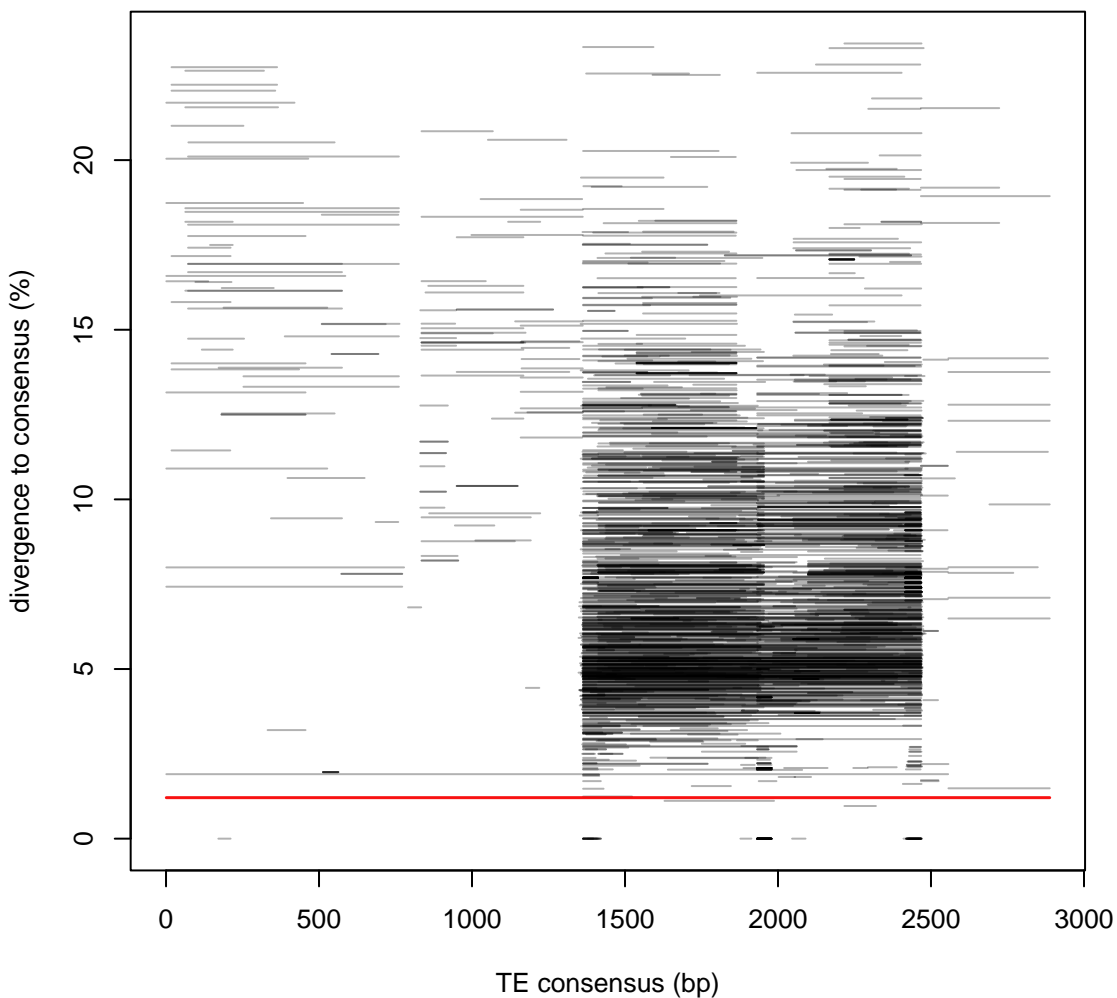
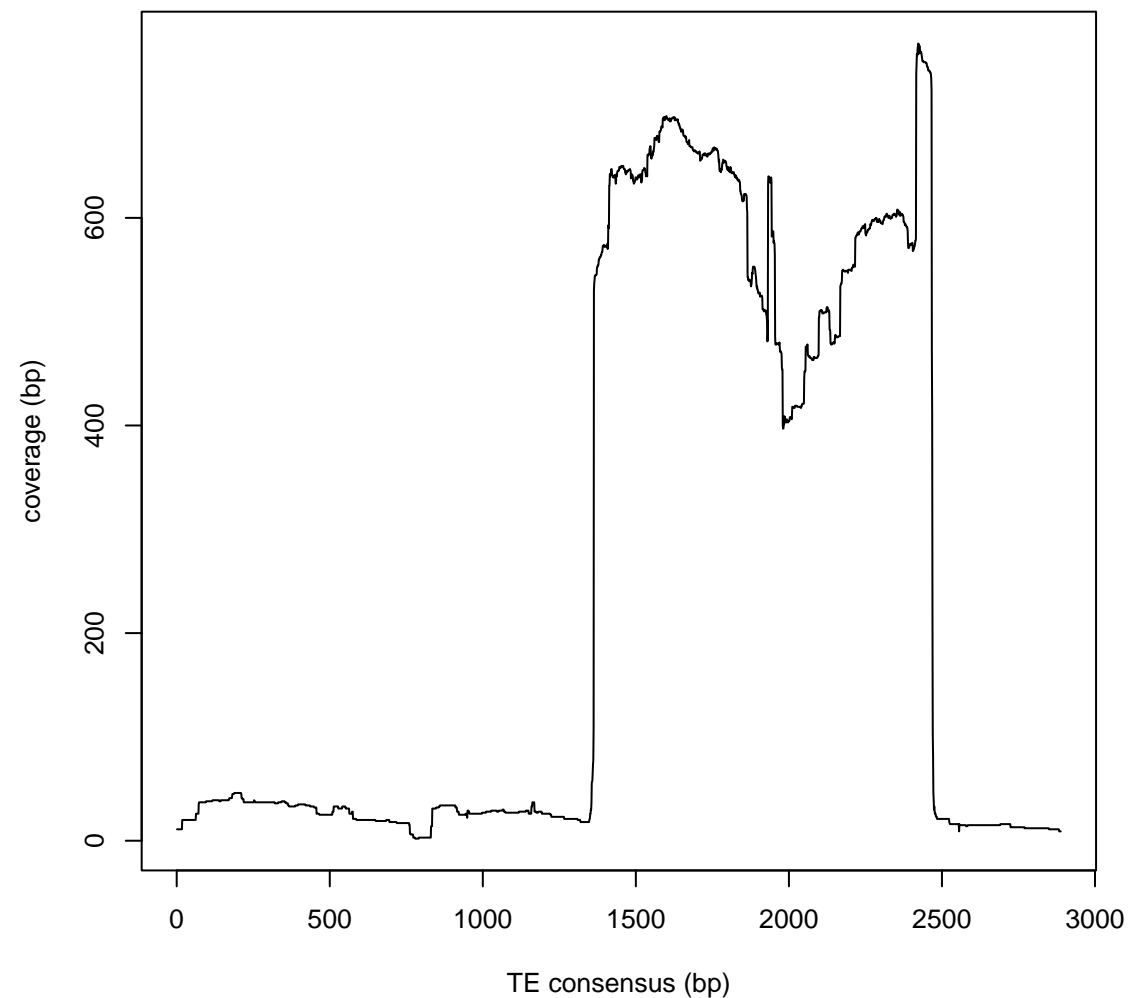


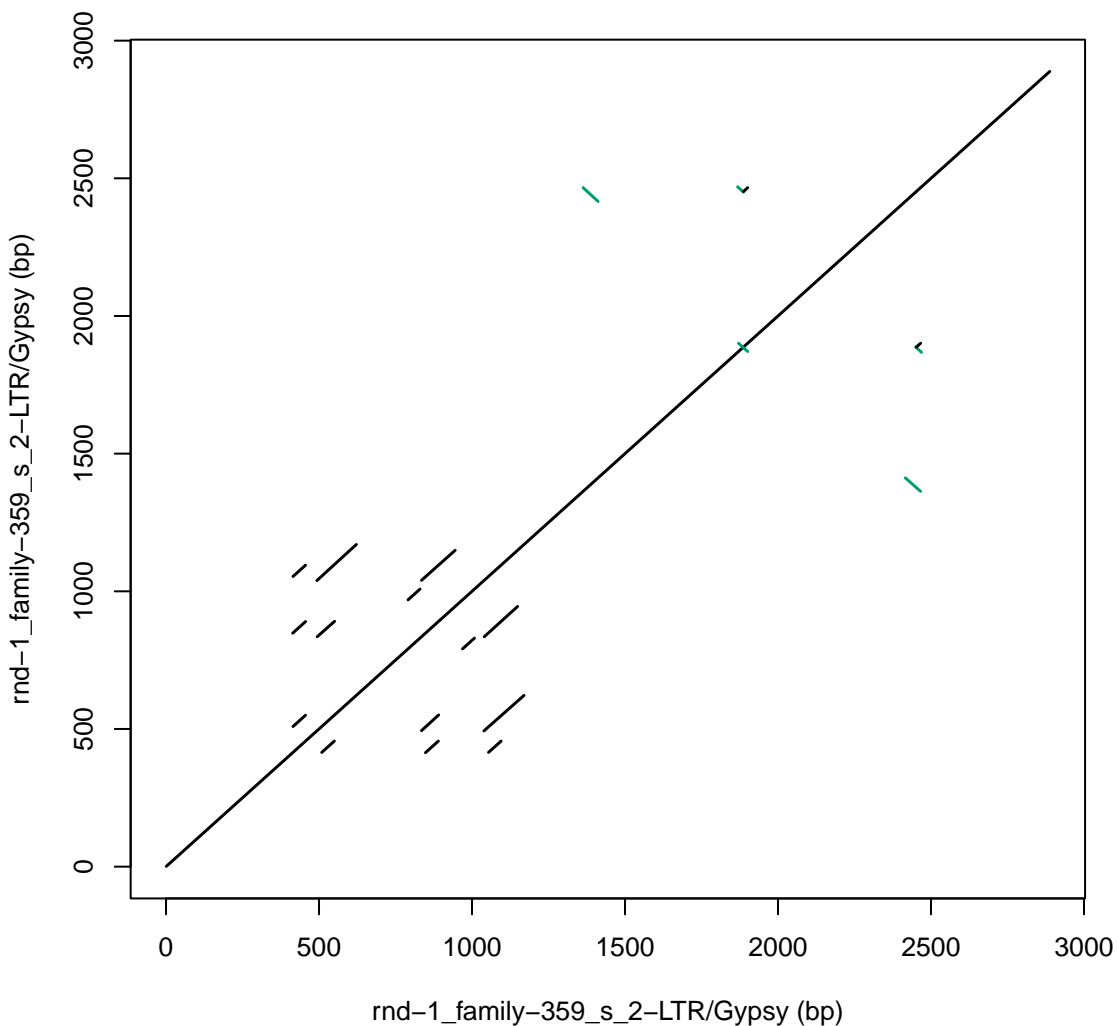
TE: rnd-1_family-359_s_2-LTR/Gypsy
consensus size: 2888bp; fragments: 2024; full length: 1 (>=2599.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

