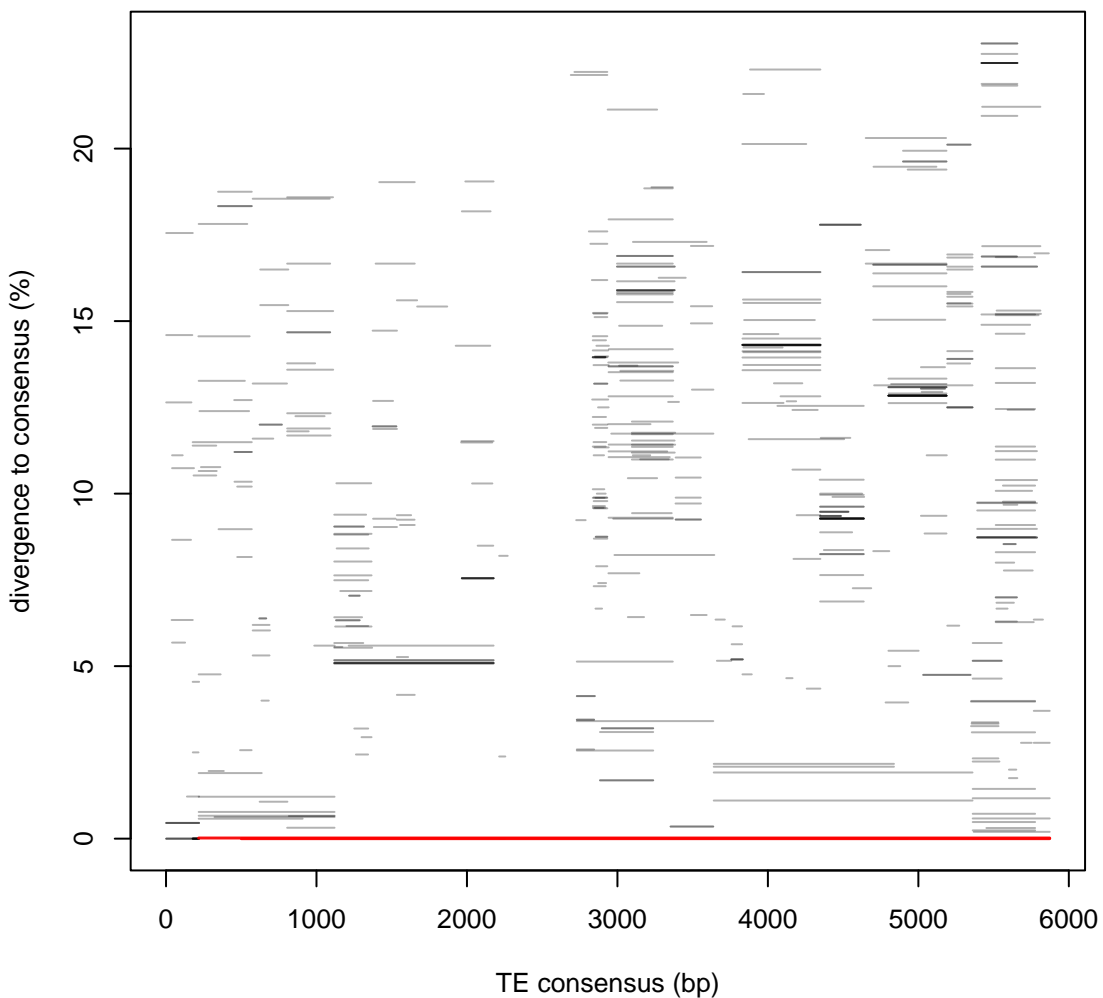
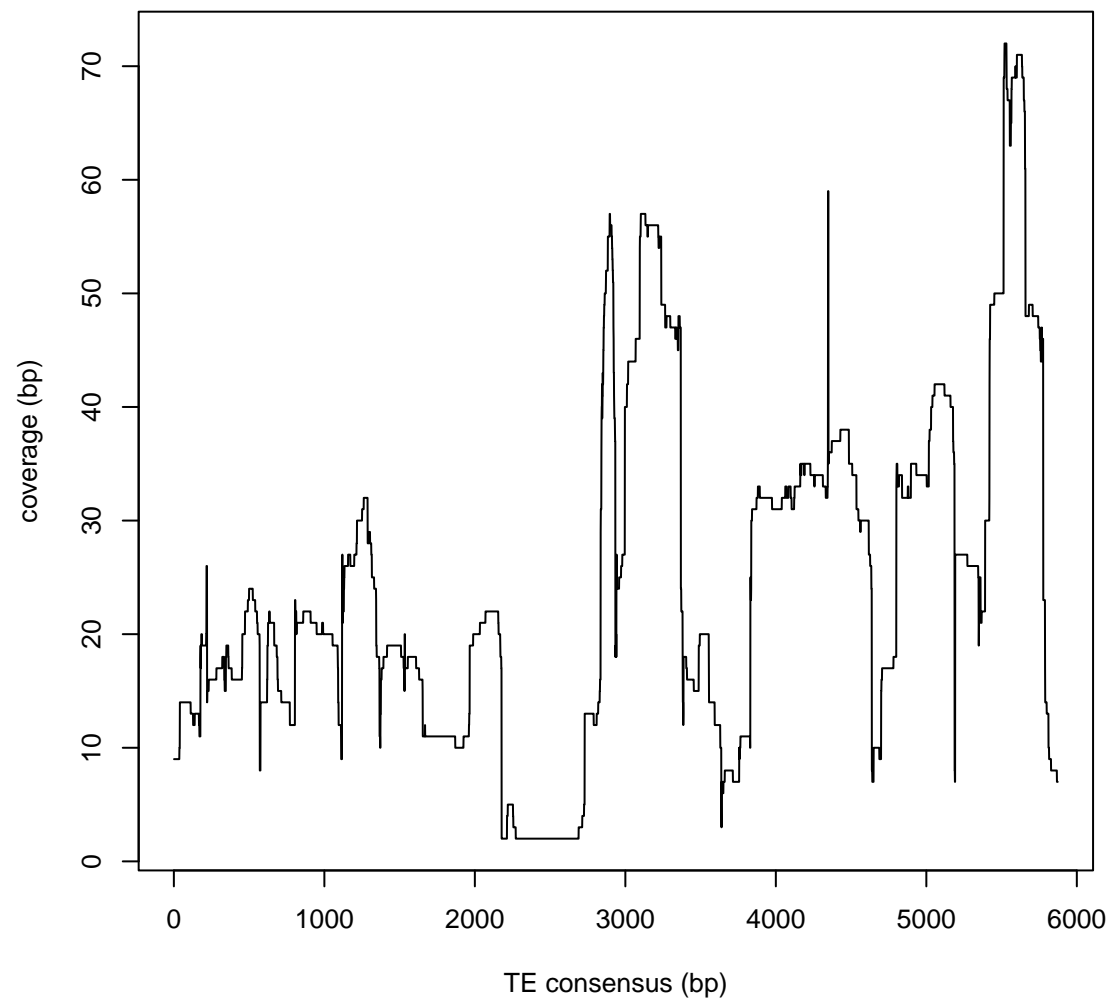


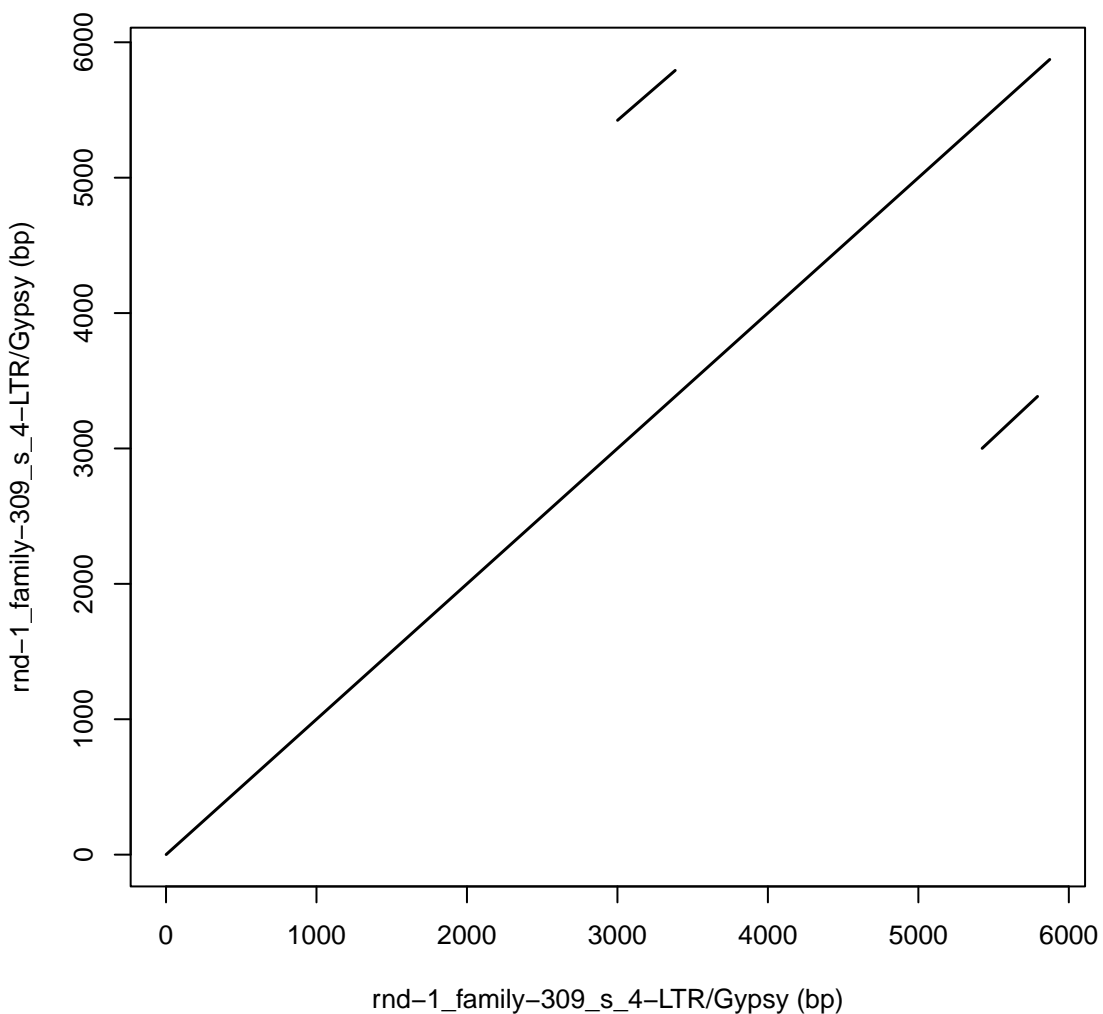
TE: rnd-1_family-309_s_4-LTR/Gypsy
consensus size: 5873bp; fragments: 500; full length: 2 (>=5285.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

