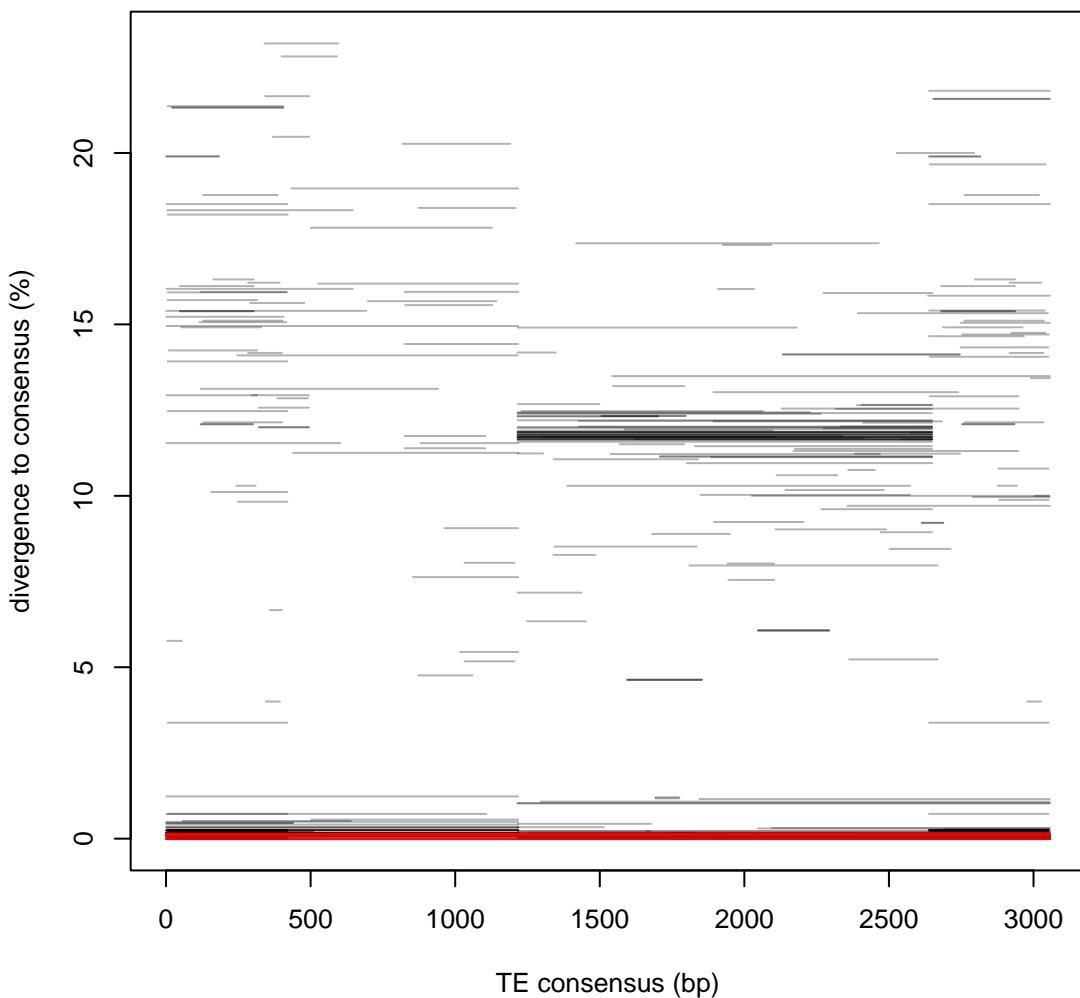
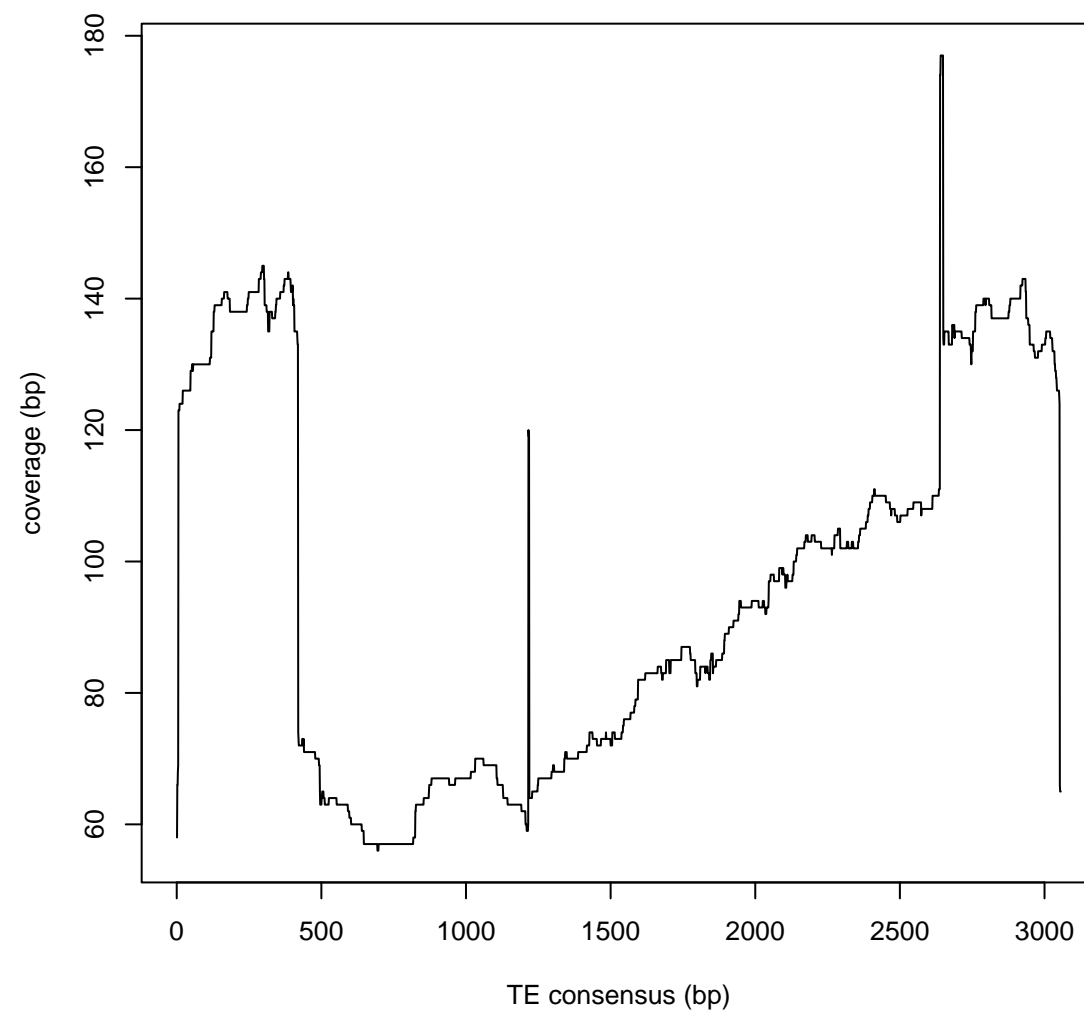


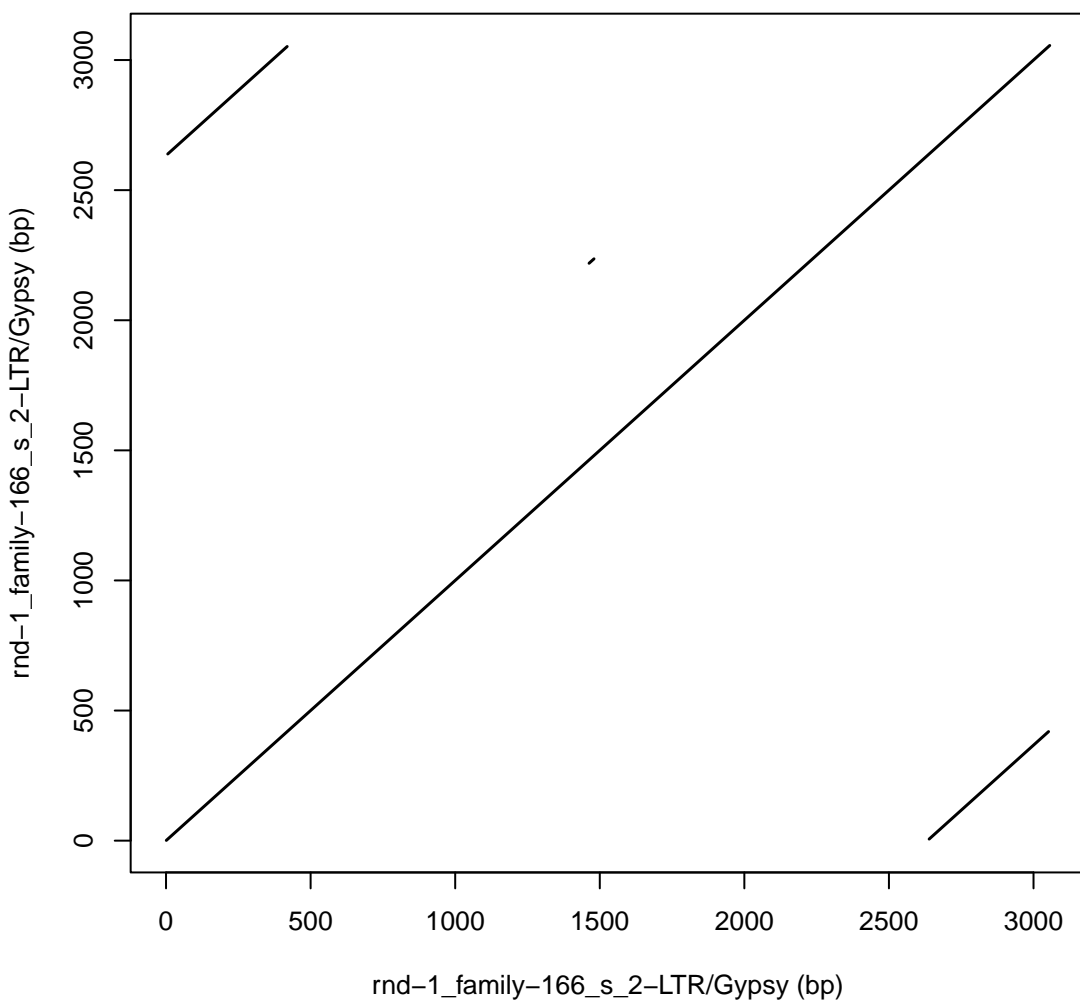
TE: rnd-1_family-166_s_2-LTR/Gypsy
consensus size: 3056bp; fragments: 433; full length: 3 (>=2750.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

