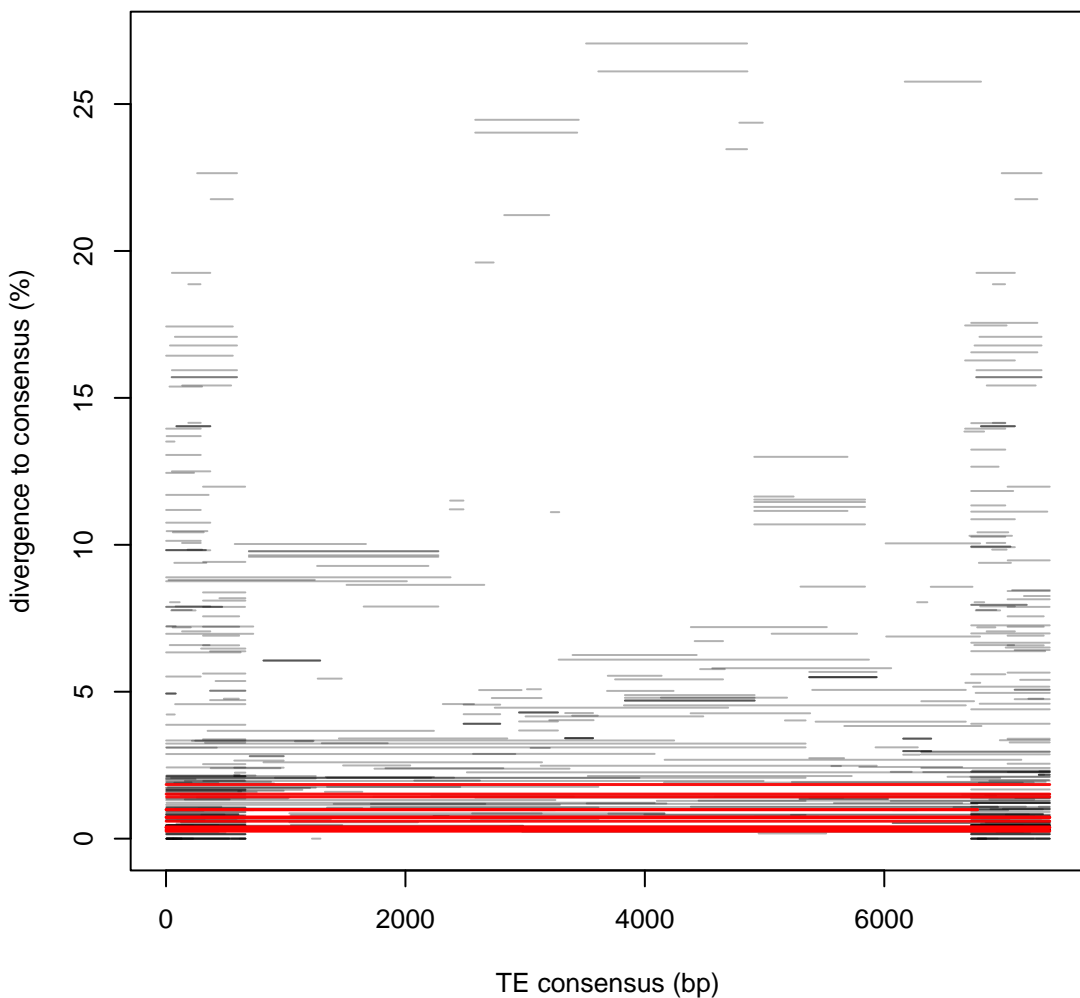
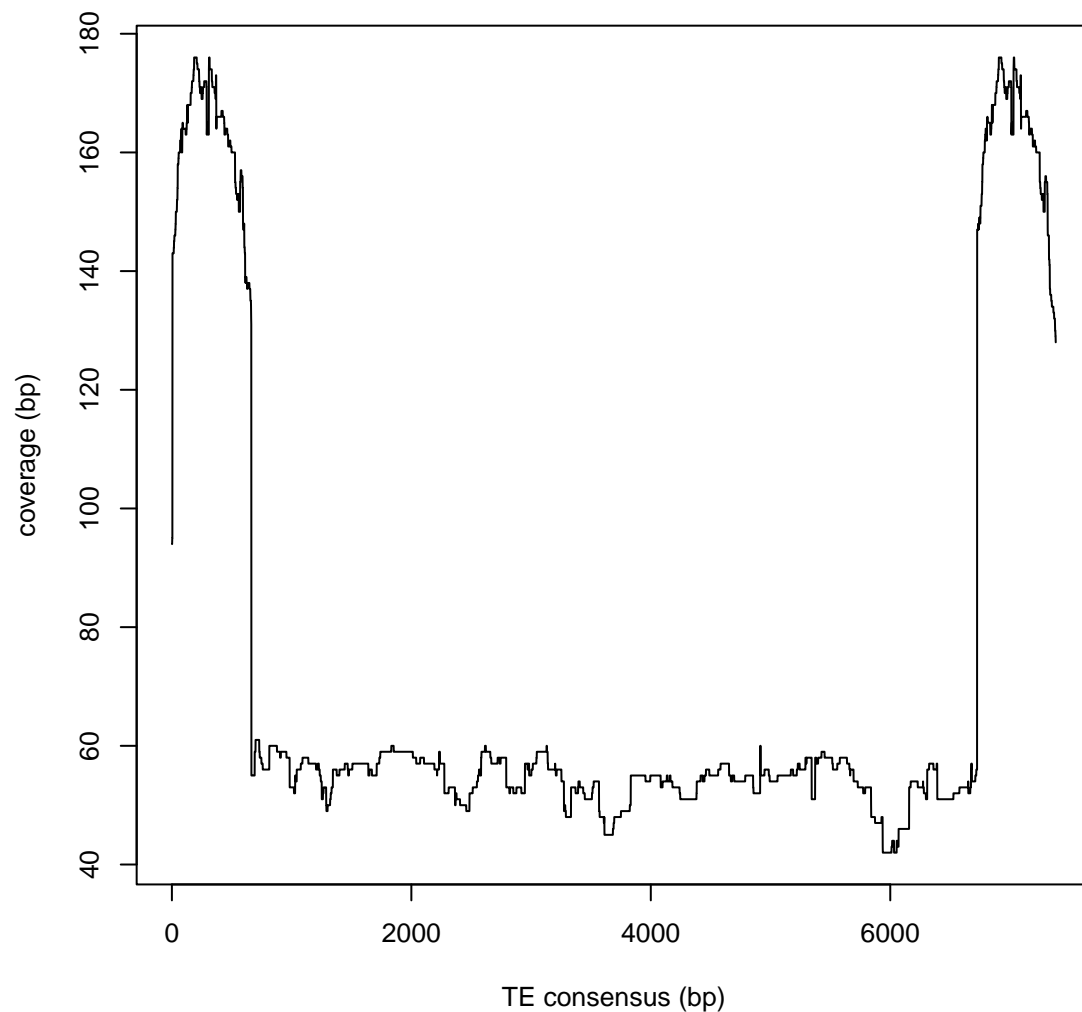


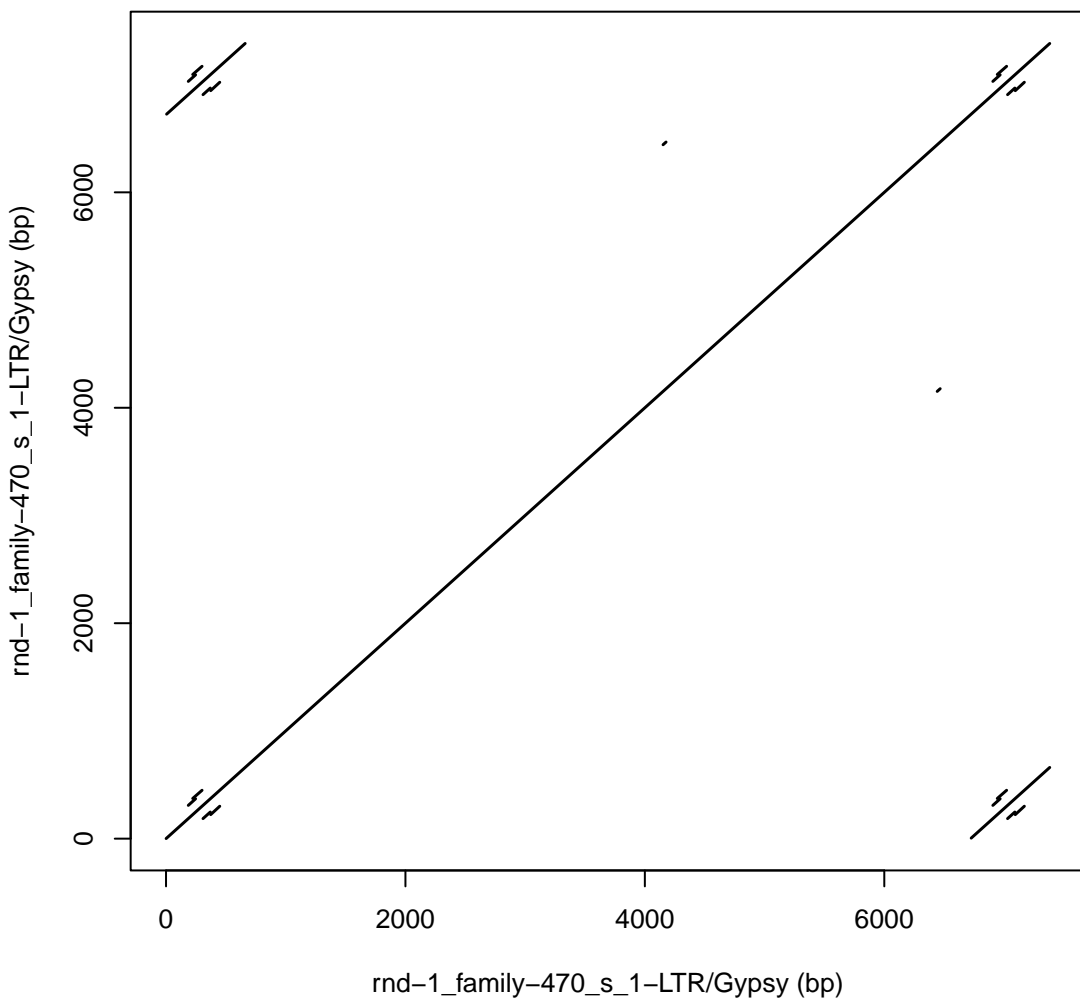
TE: rnd-1_family-470_s_1-LTR/Gypsy
consensus size: 7382bp; fragments: 583; full length: 16 (>=6643.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

