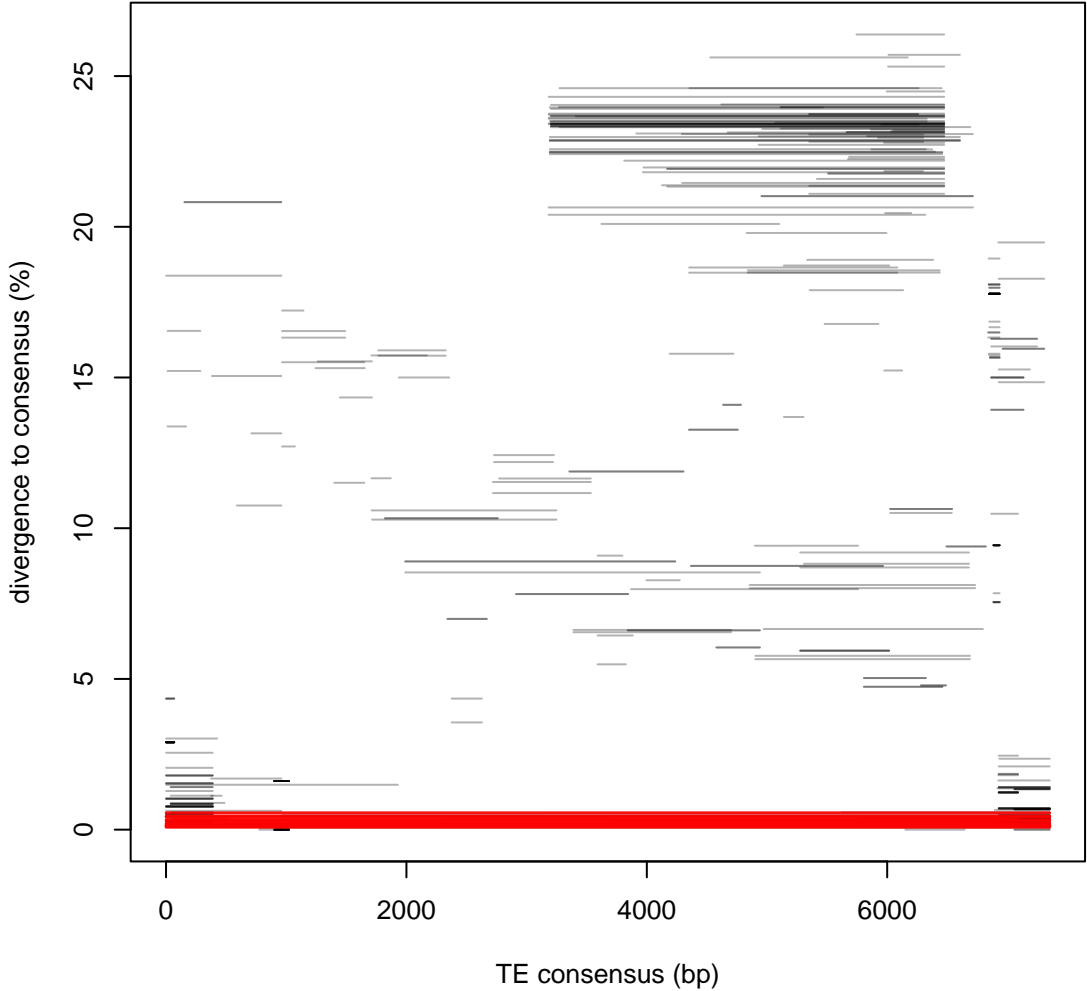
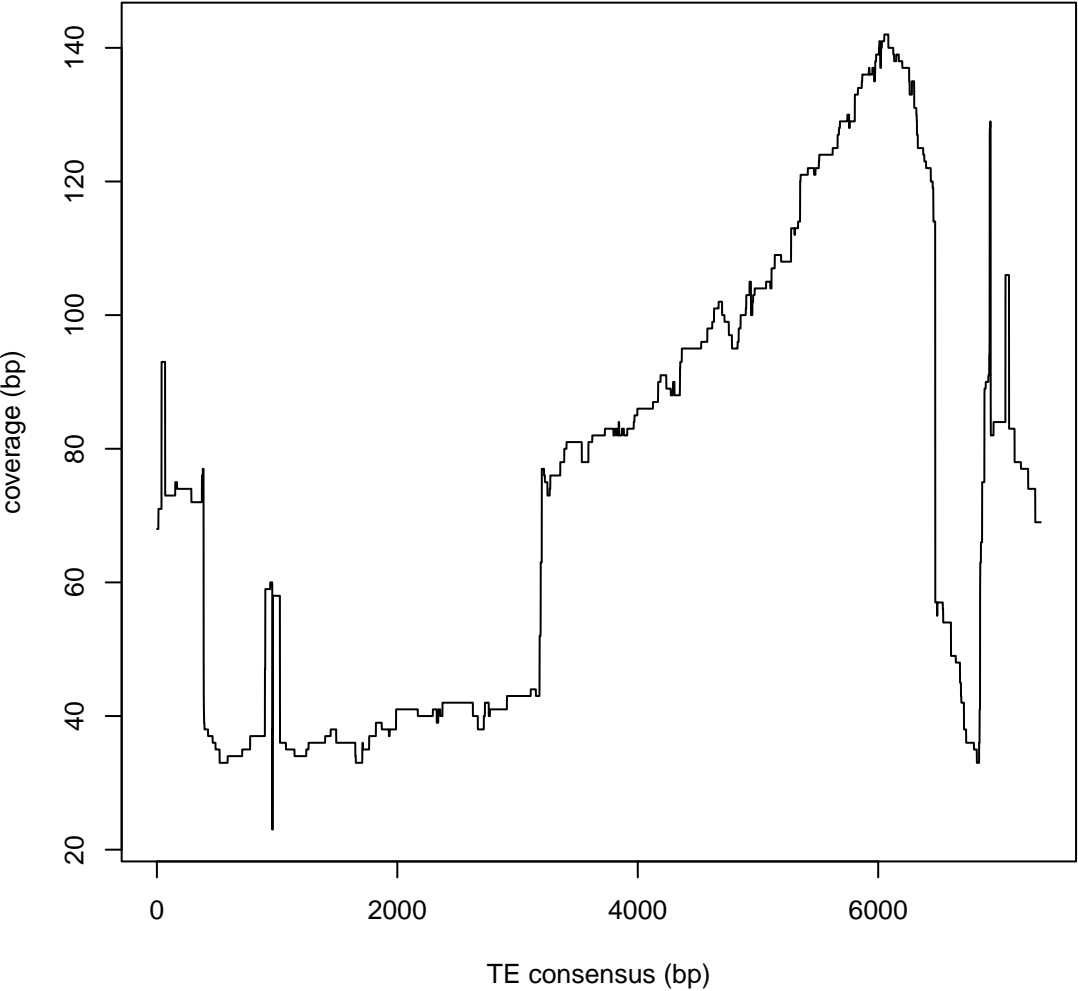


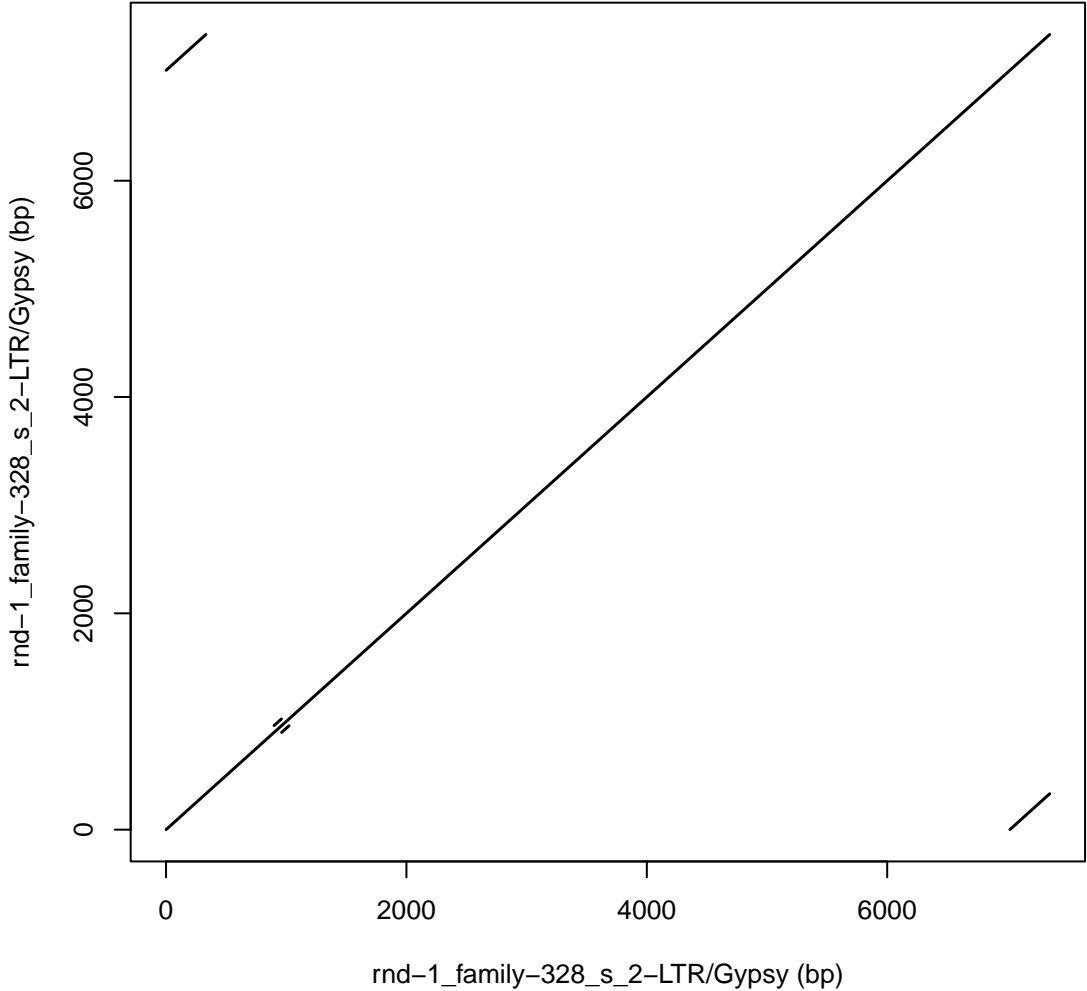
TE: rnd-1_family-328_s_2-LTR/Gypsy
consensus size: 7352bp; fragments: 458; full length: 20 (>=6616.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

