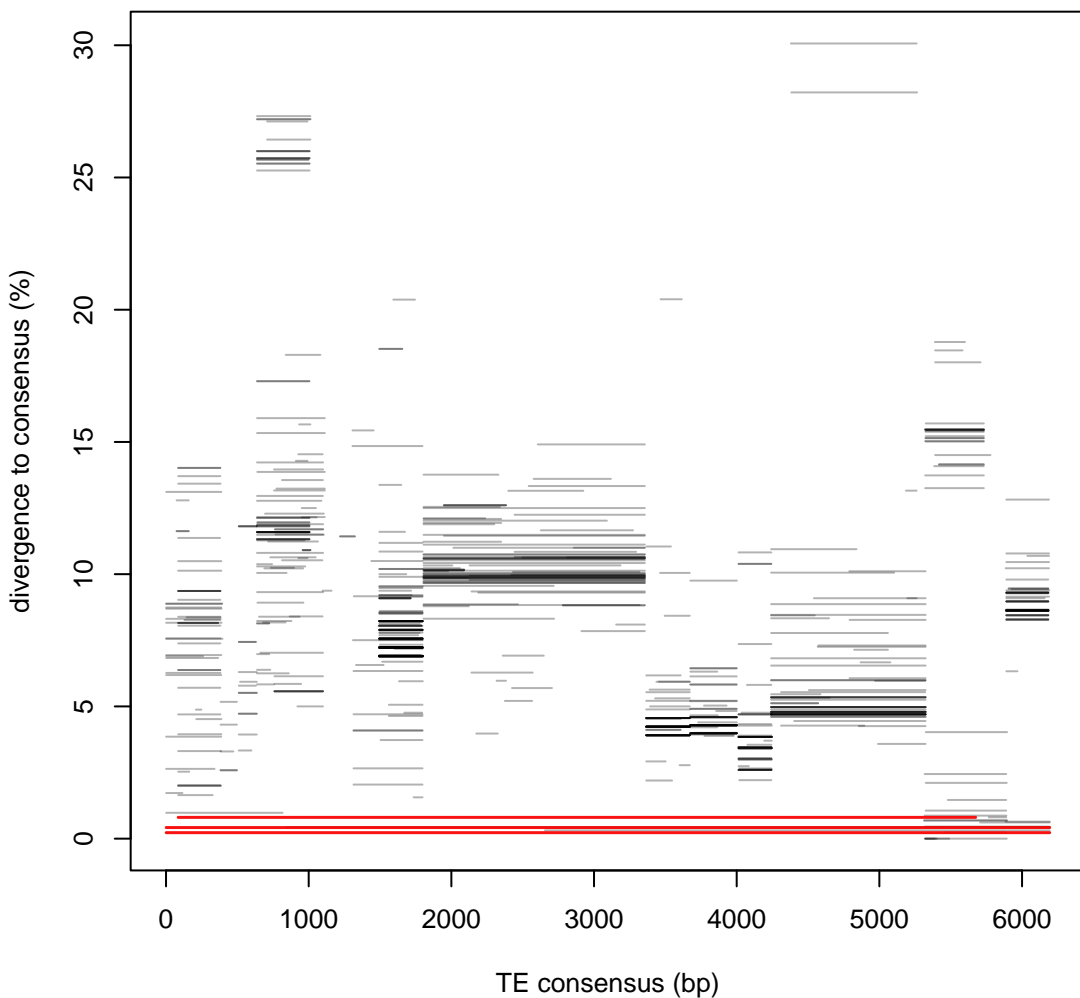
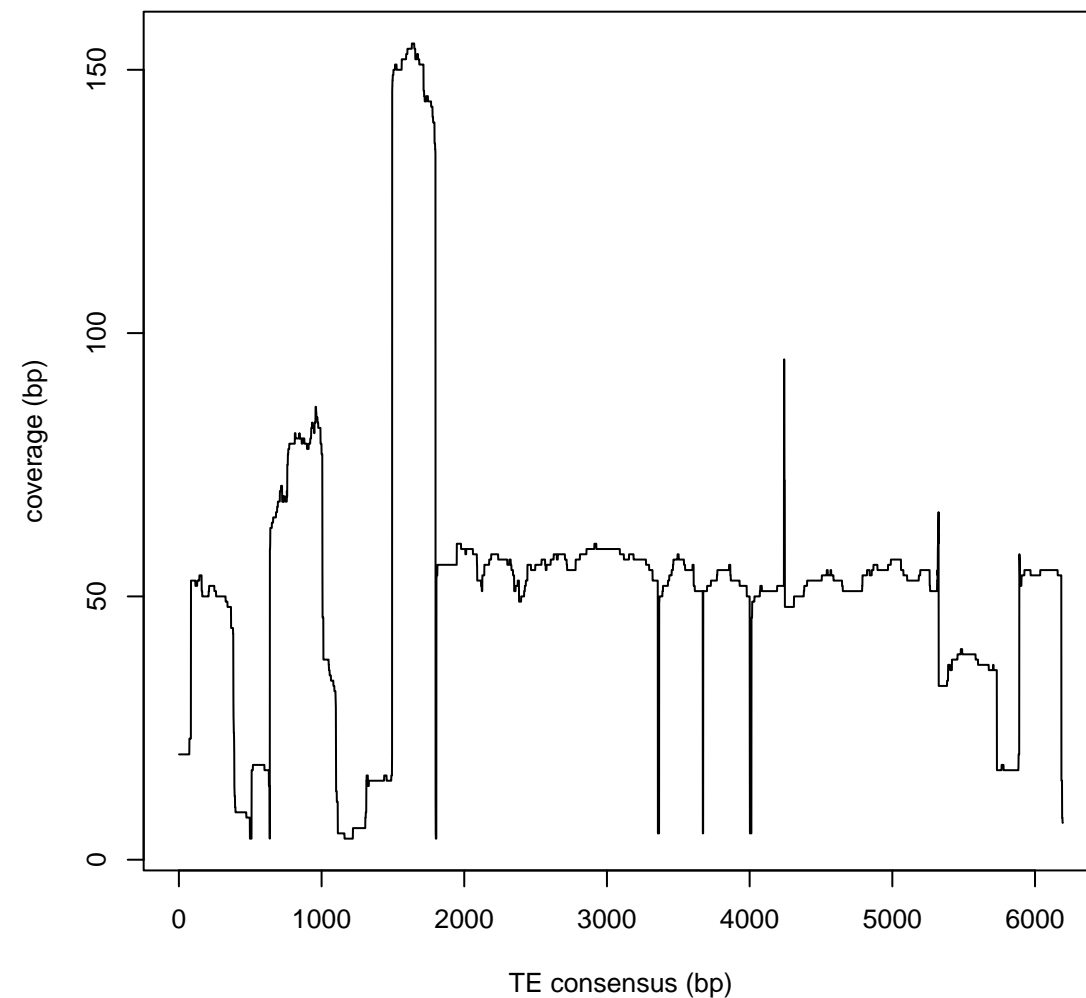


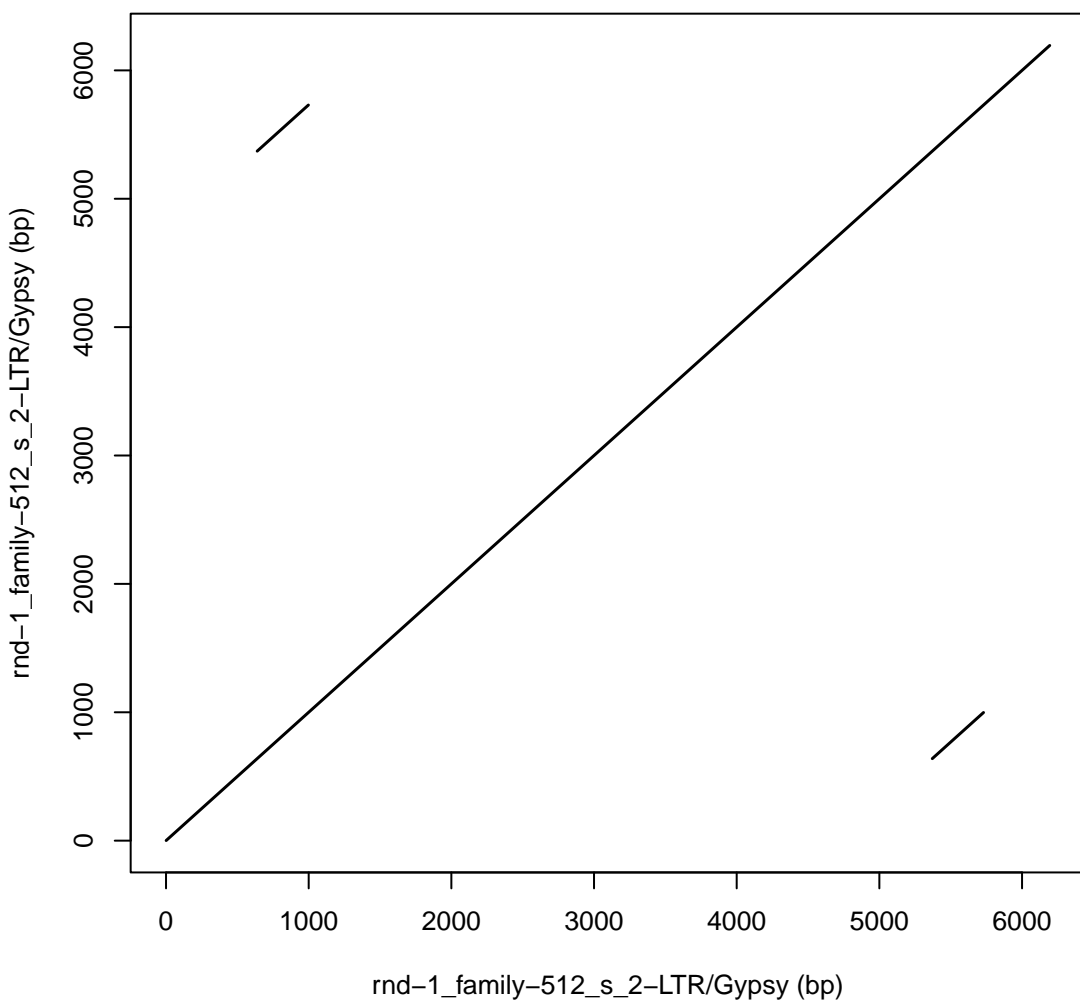
TE: rnd-1_family-512_s_2-LTR/Gypsy
consensus size: 6194bp; fragments: 720; full length: 3 (>=5574.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

