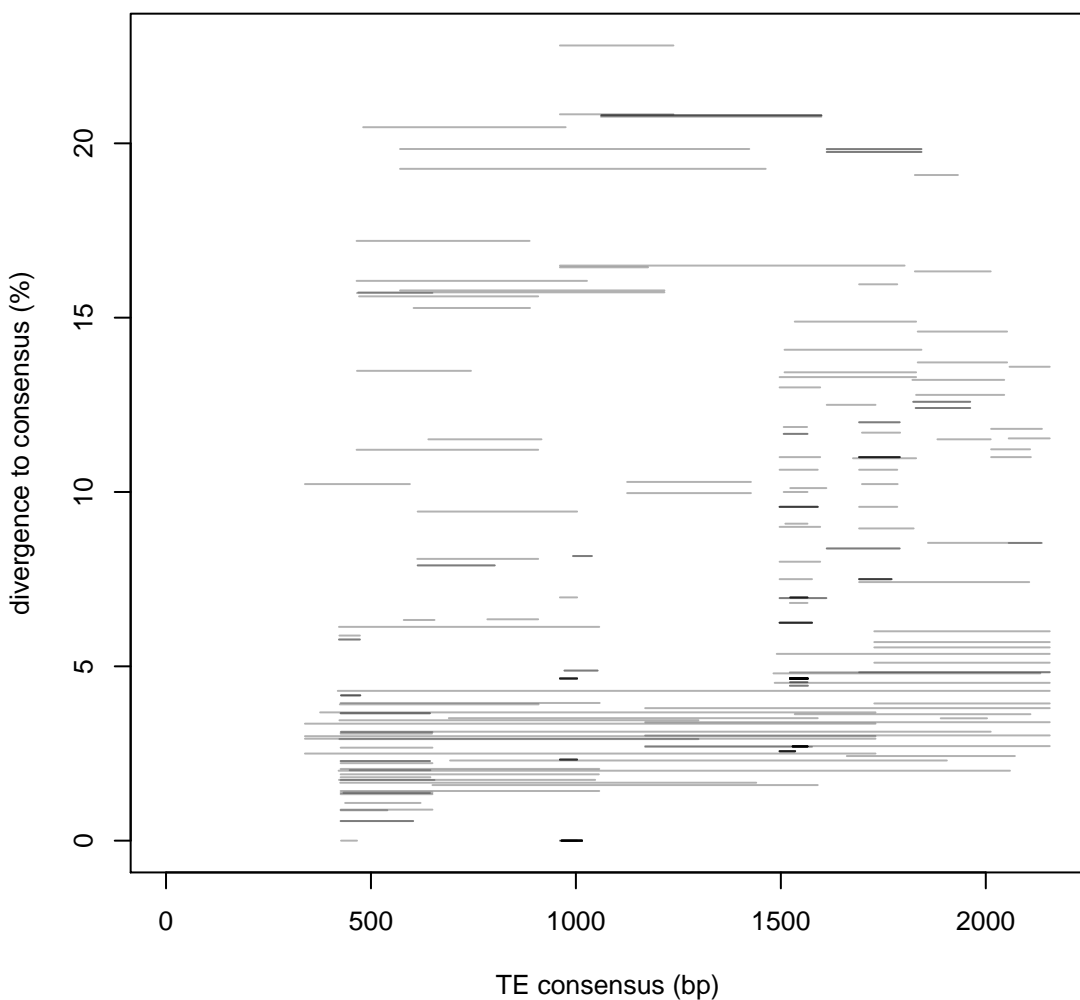
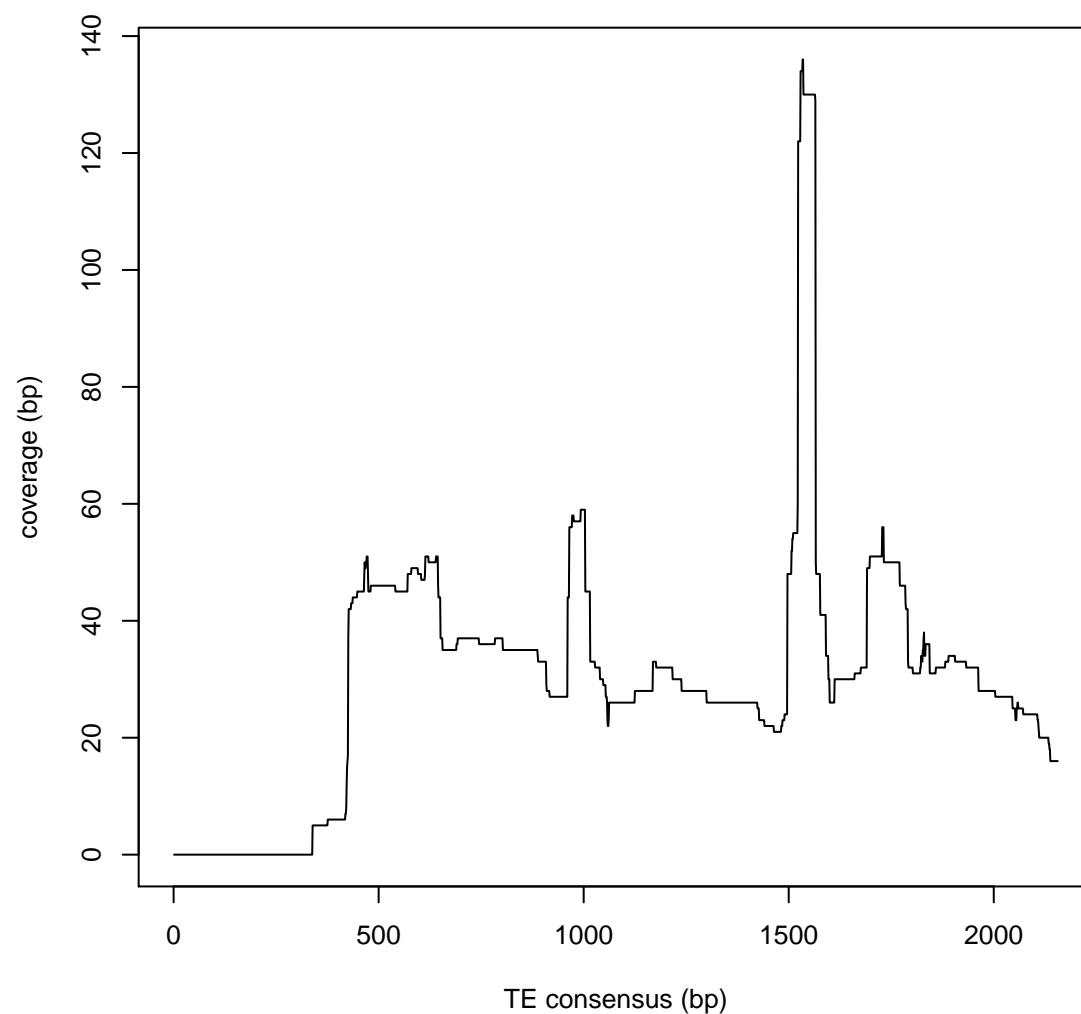


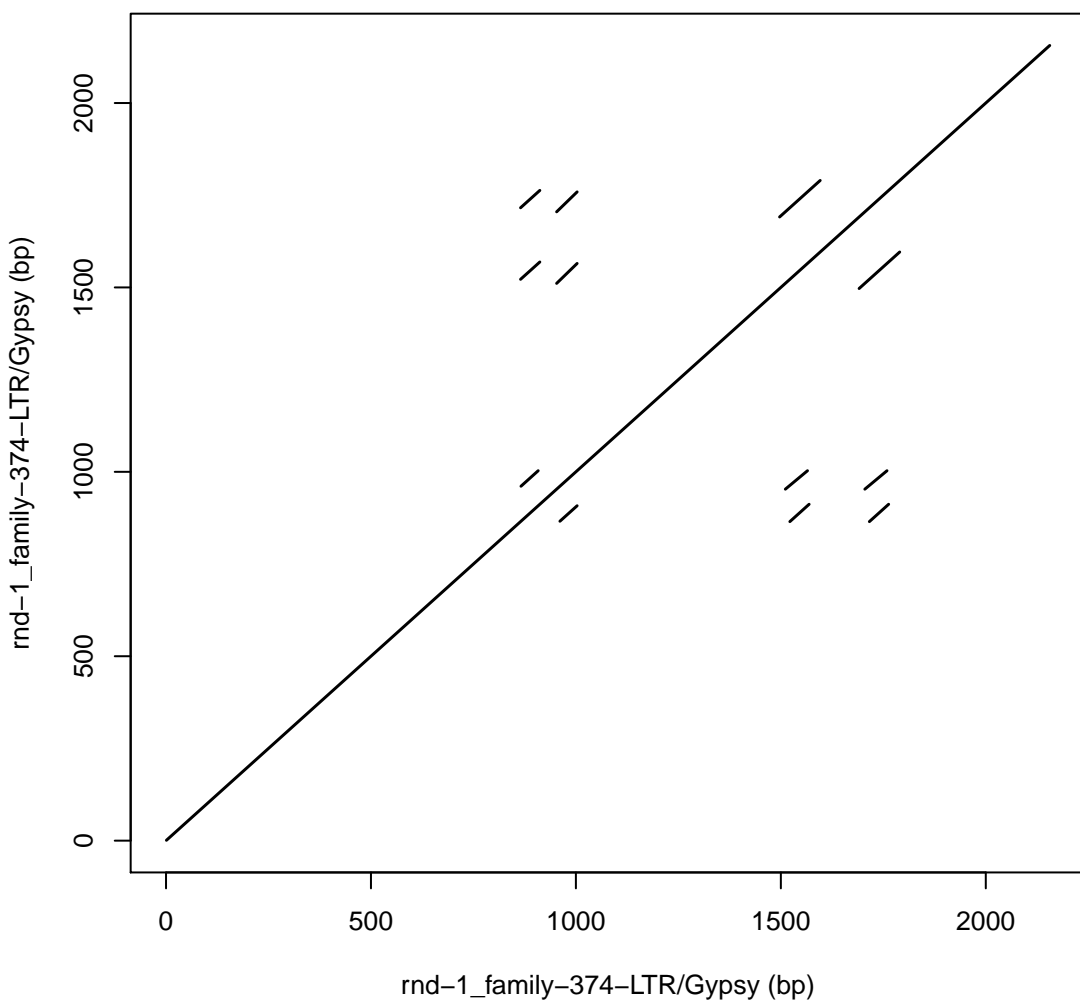
TE: rnd-1_family-374-LTR/Gypsy
consensus size: 2156bp; fragments: 278; full length: 0 (>=1940.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

