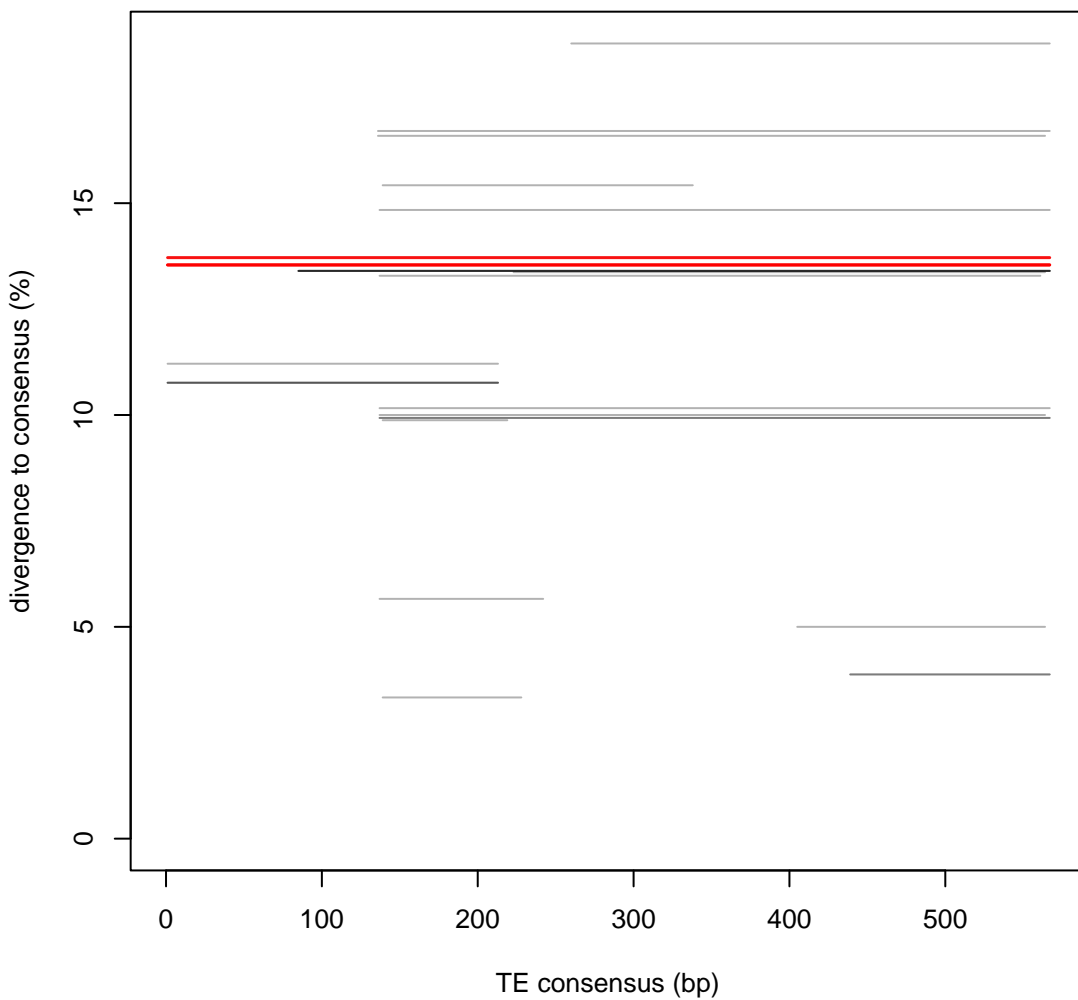
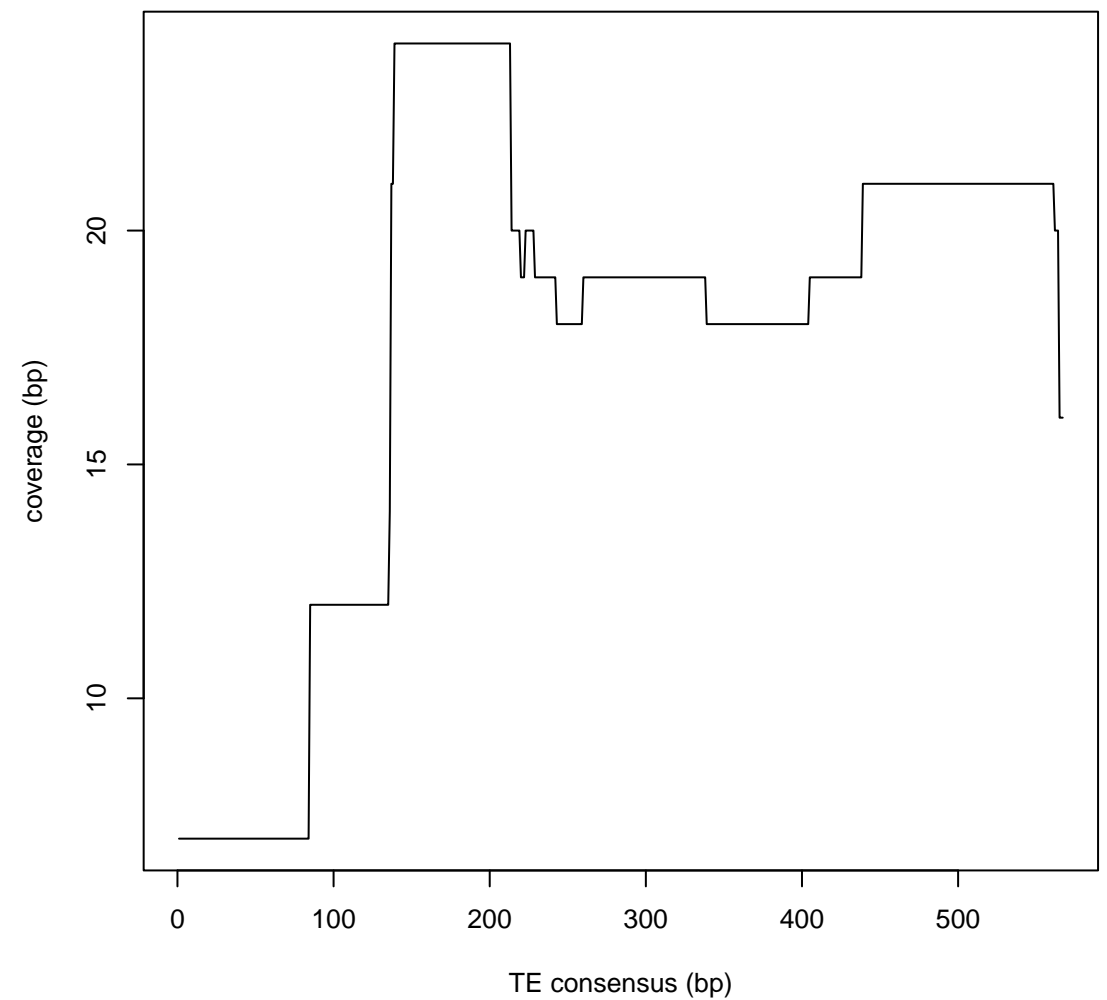


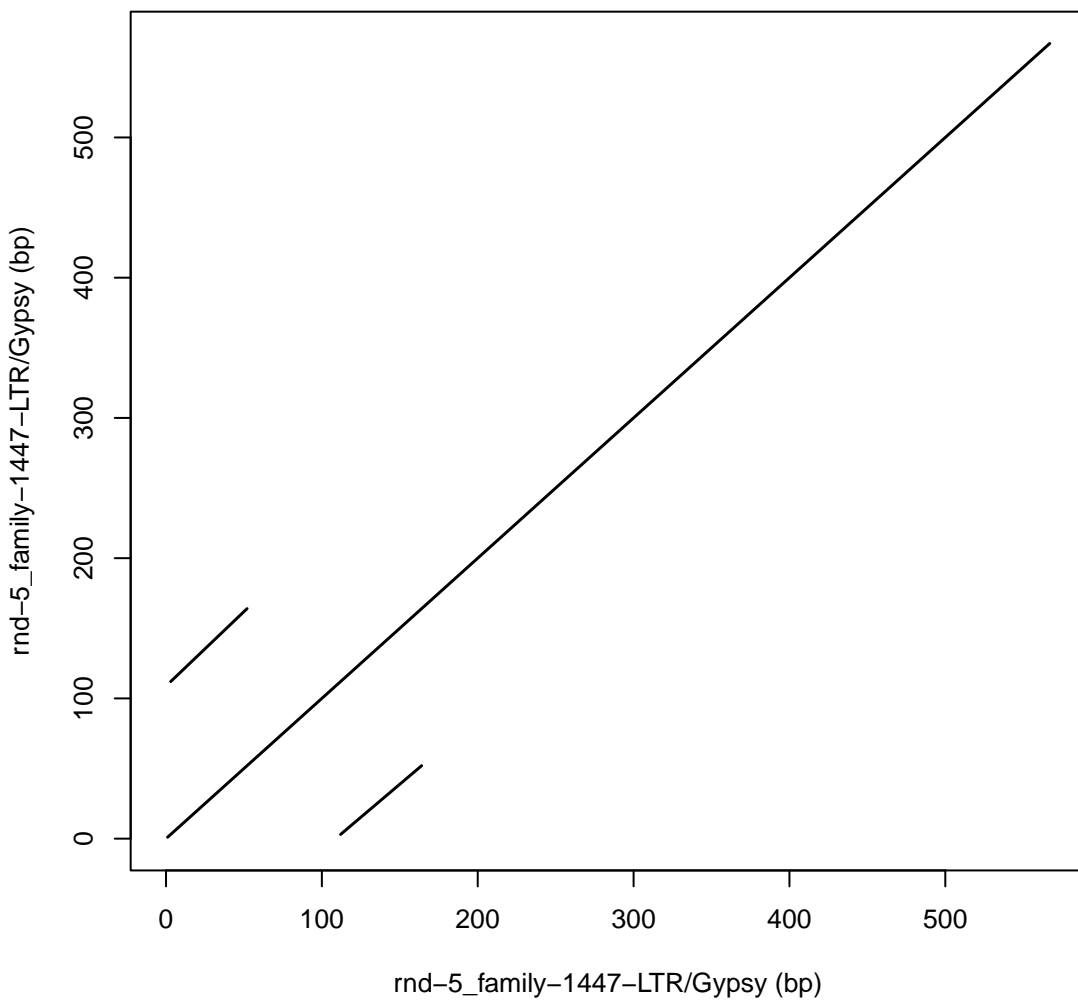
TE: rnd-5_family-1447-LTR/Gypsy
consensus size: 567bp; fragments: 29; full length: 3 (≥ 510.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

