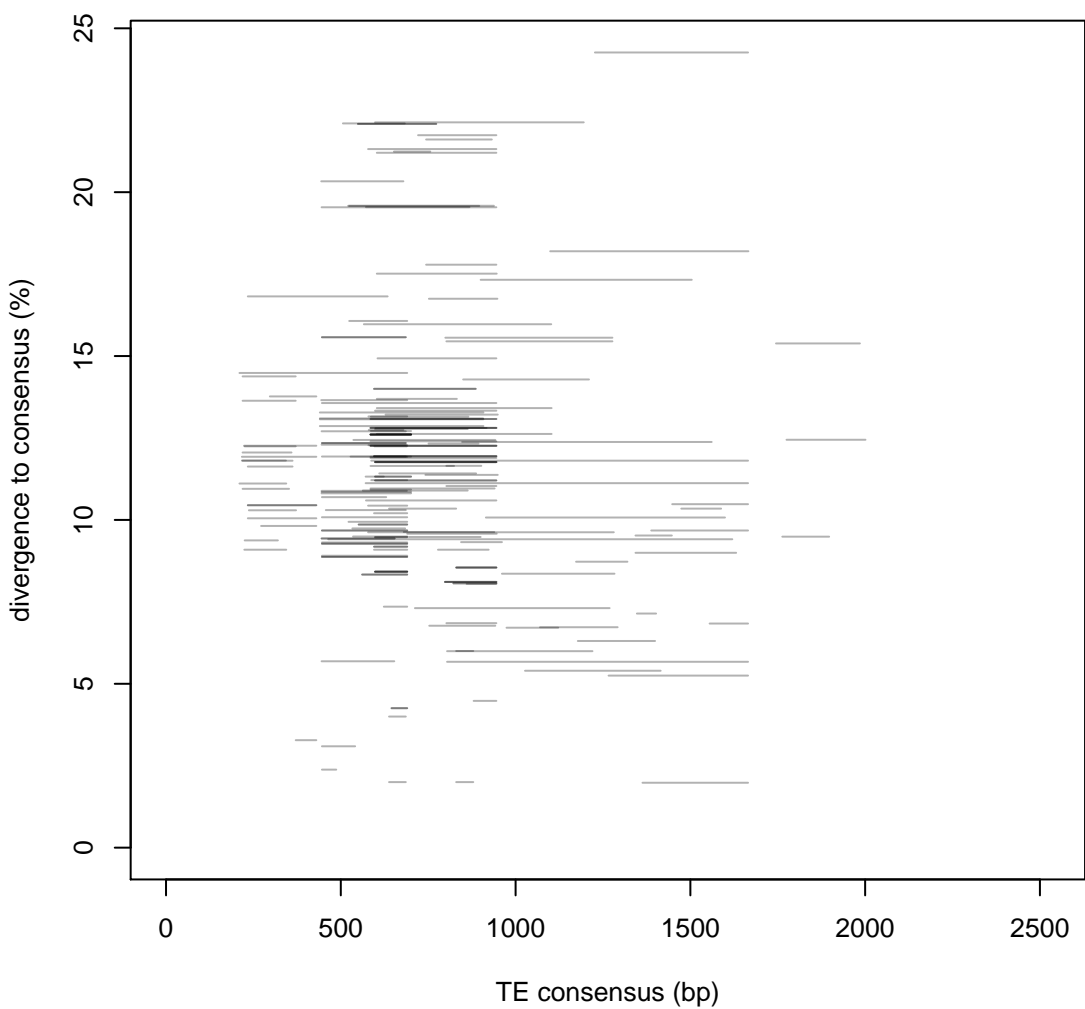
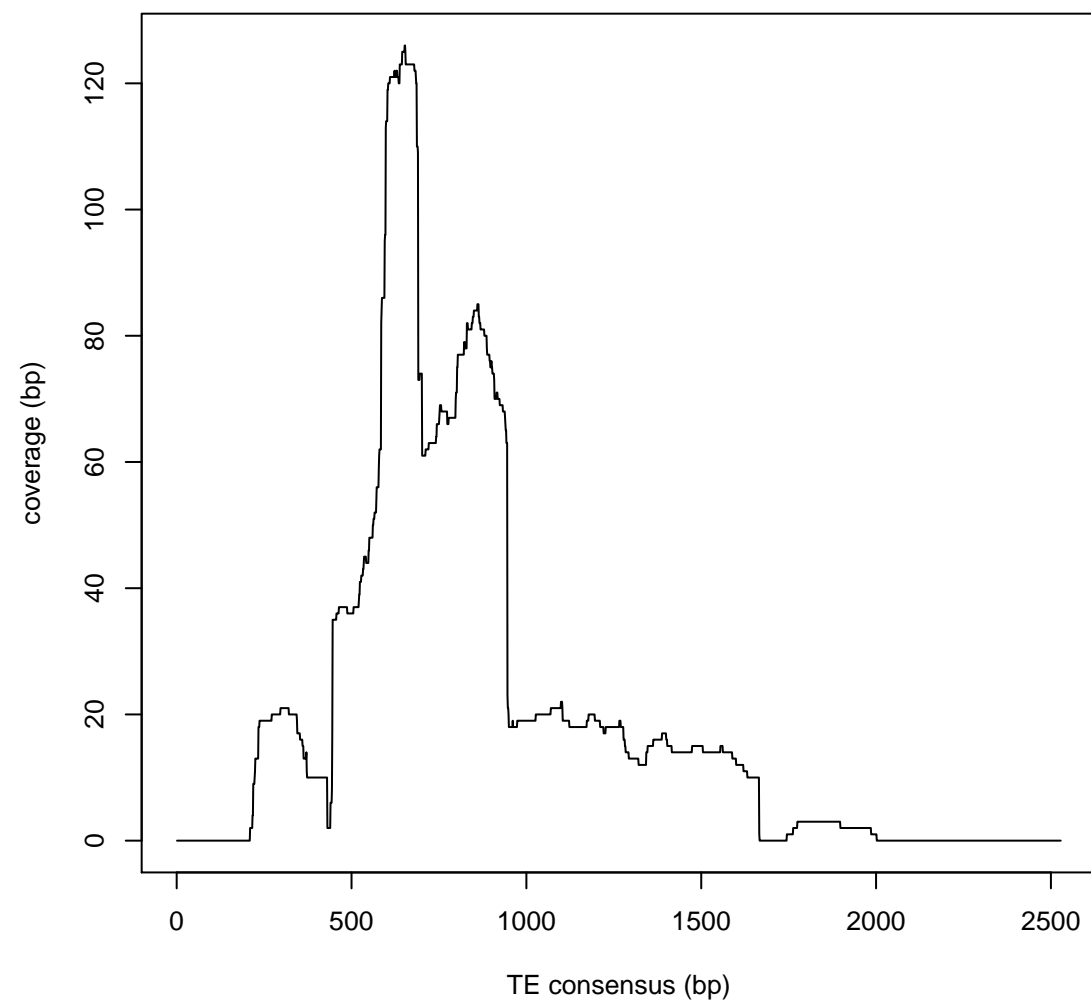


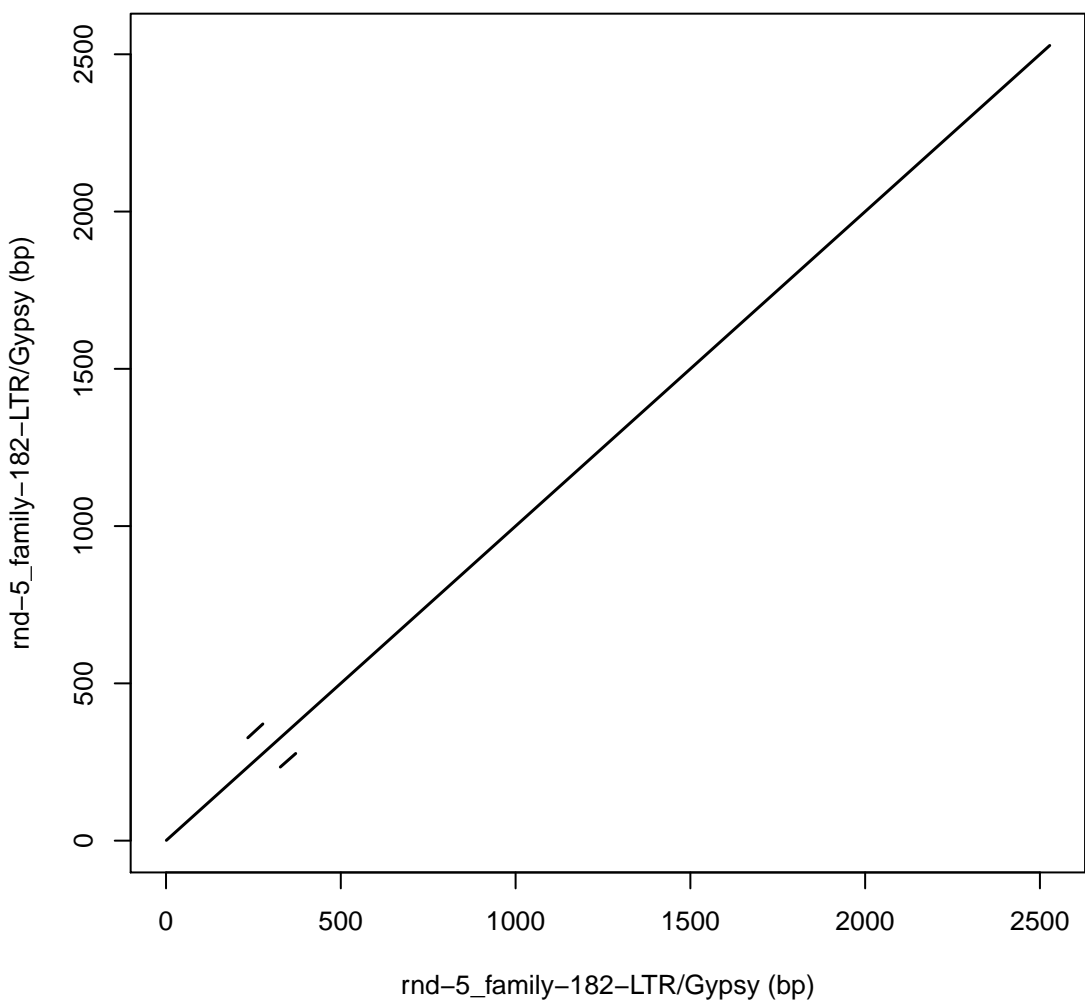
TE: rnd-5_family-182-LTR/Gypsy
consensus size: 2528bp; fragments: 207; full length: 0 (≥ 2275.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

