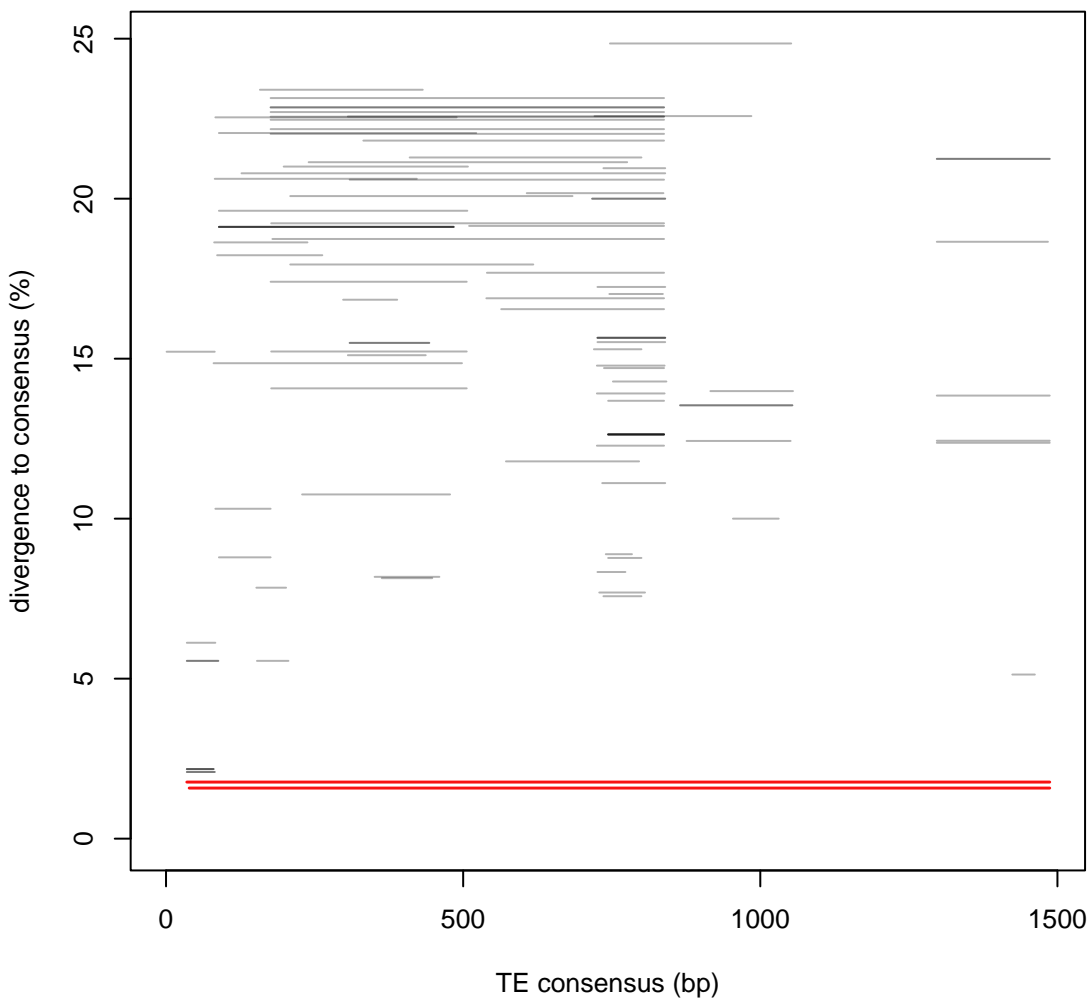
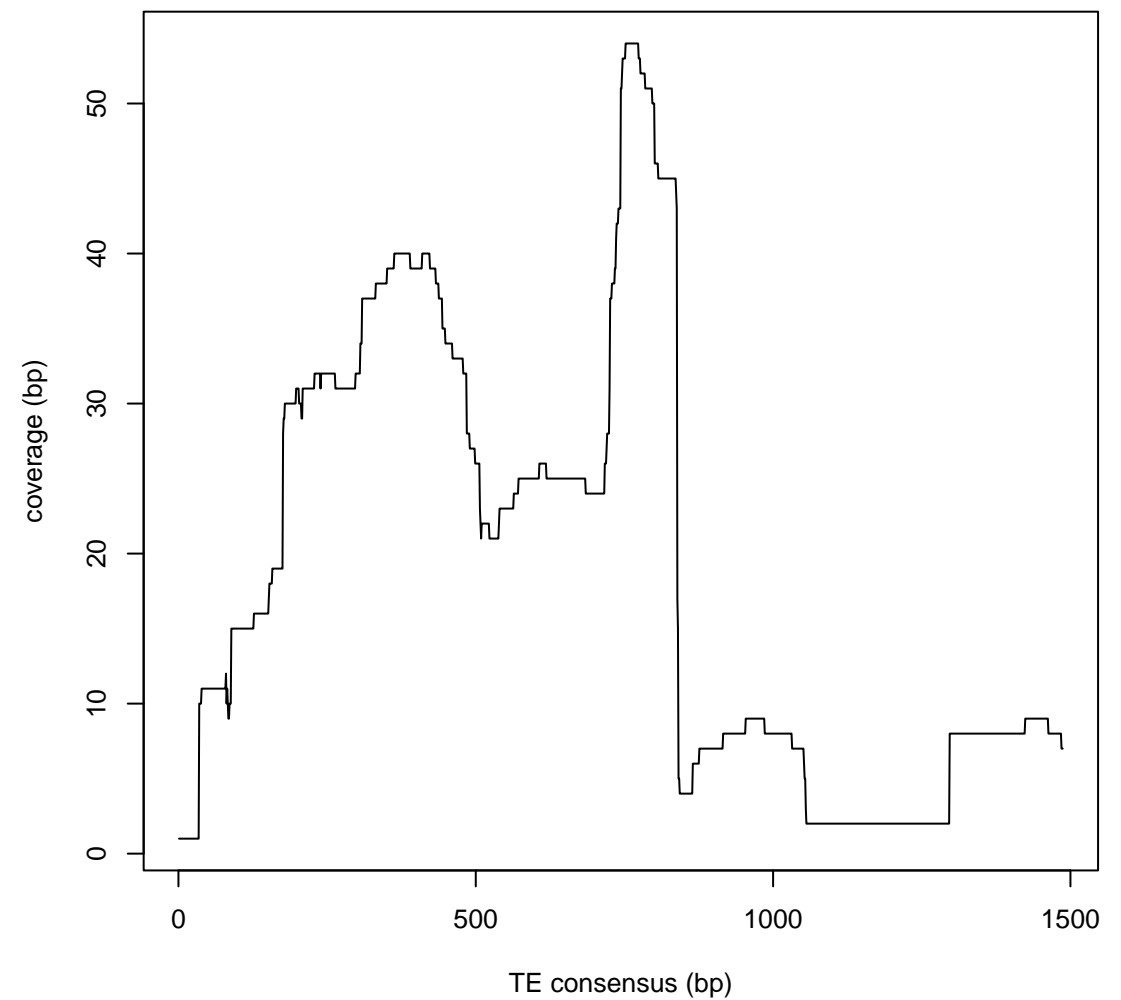


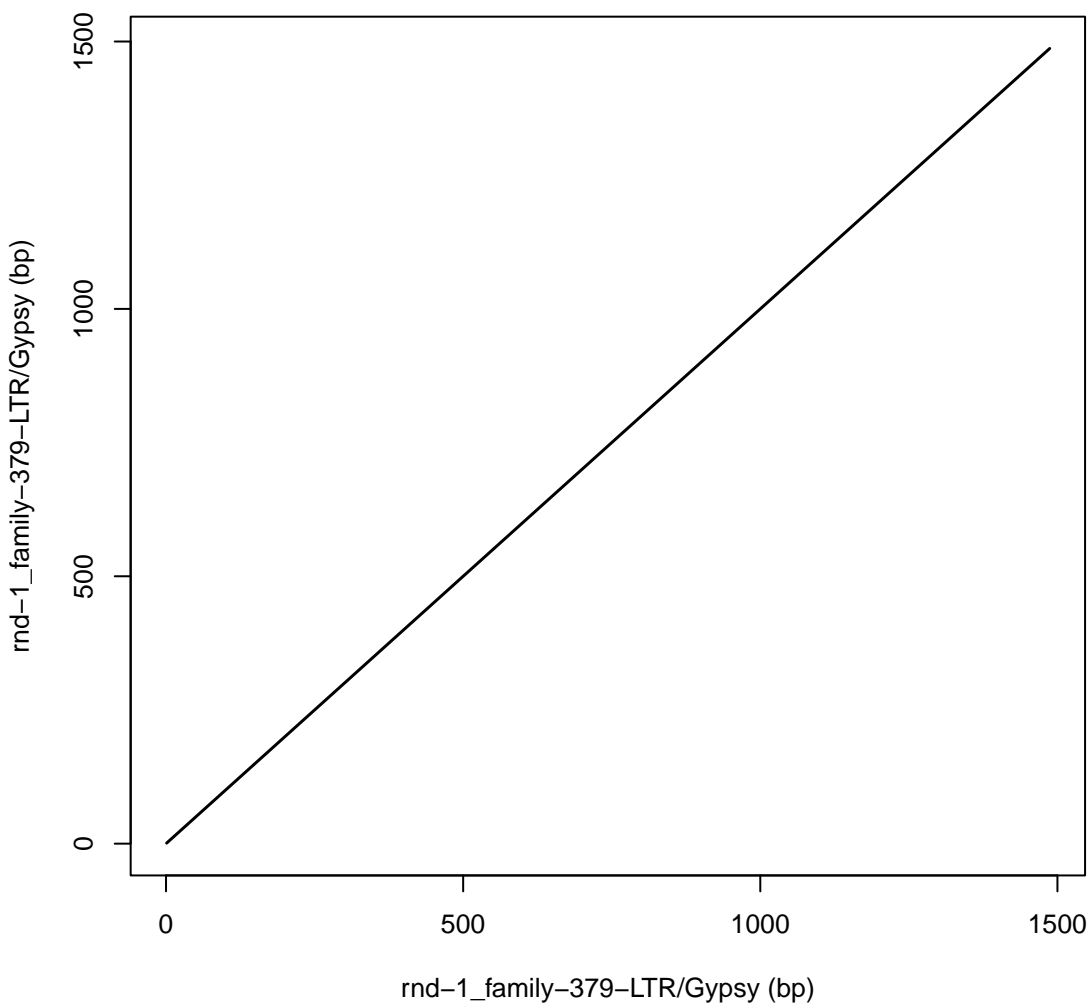
TE: rnd-1_family-379-LTR/Gypsy
consensus size: 1487bp; fragments: 104; full length: 2 (≥ 1338.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

