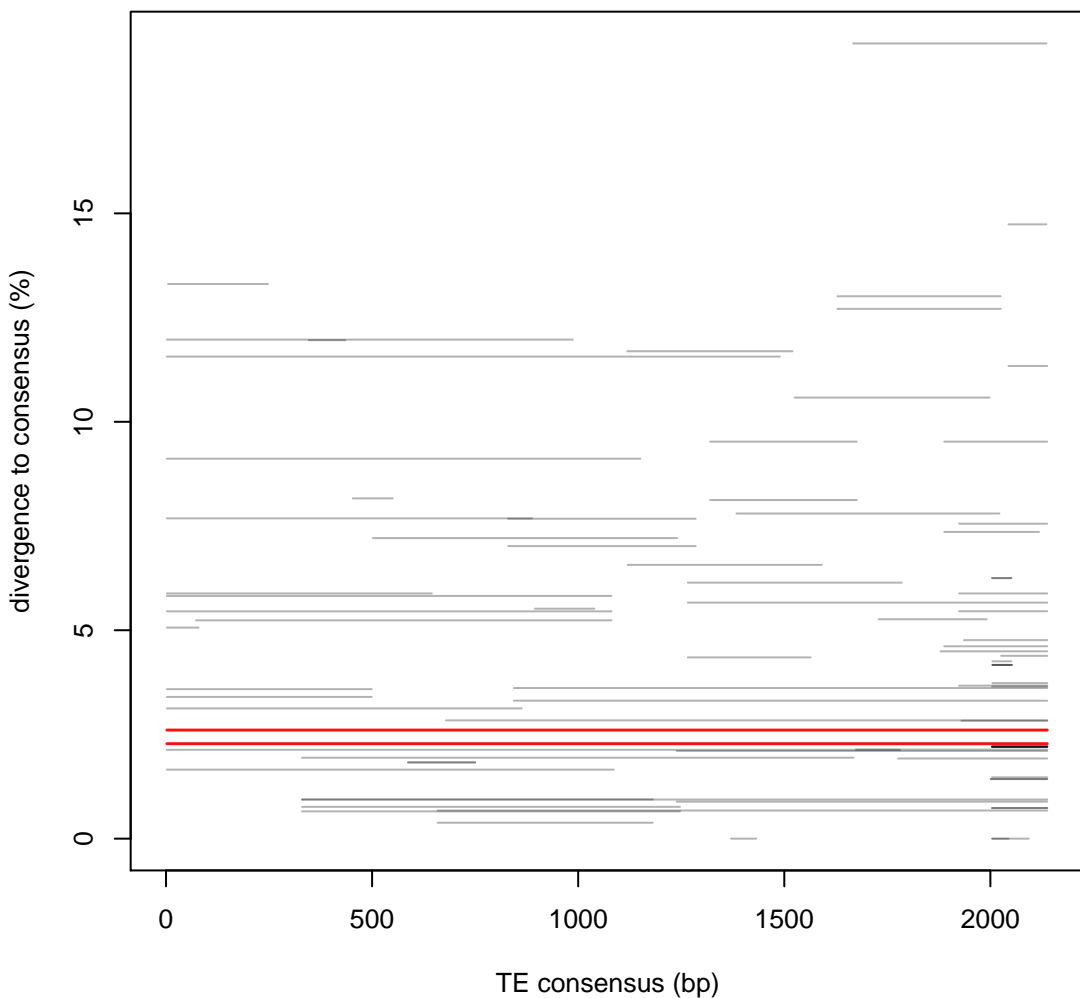
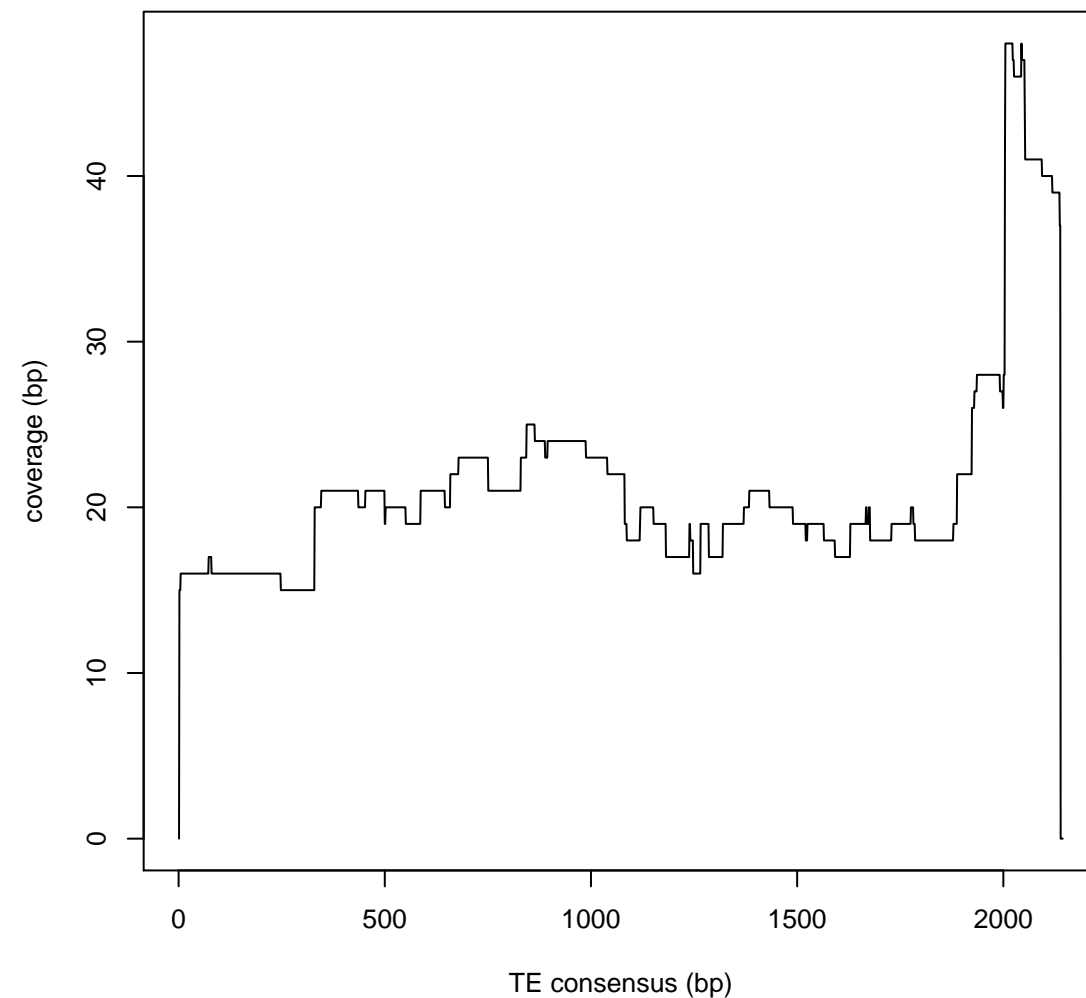


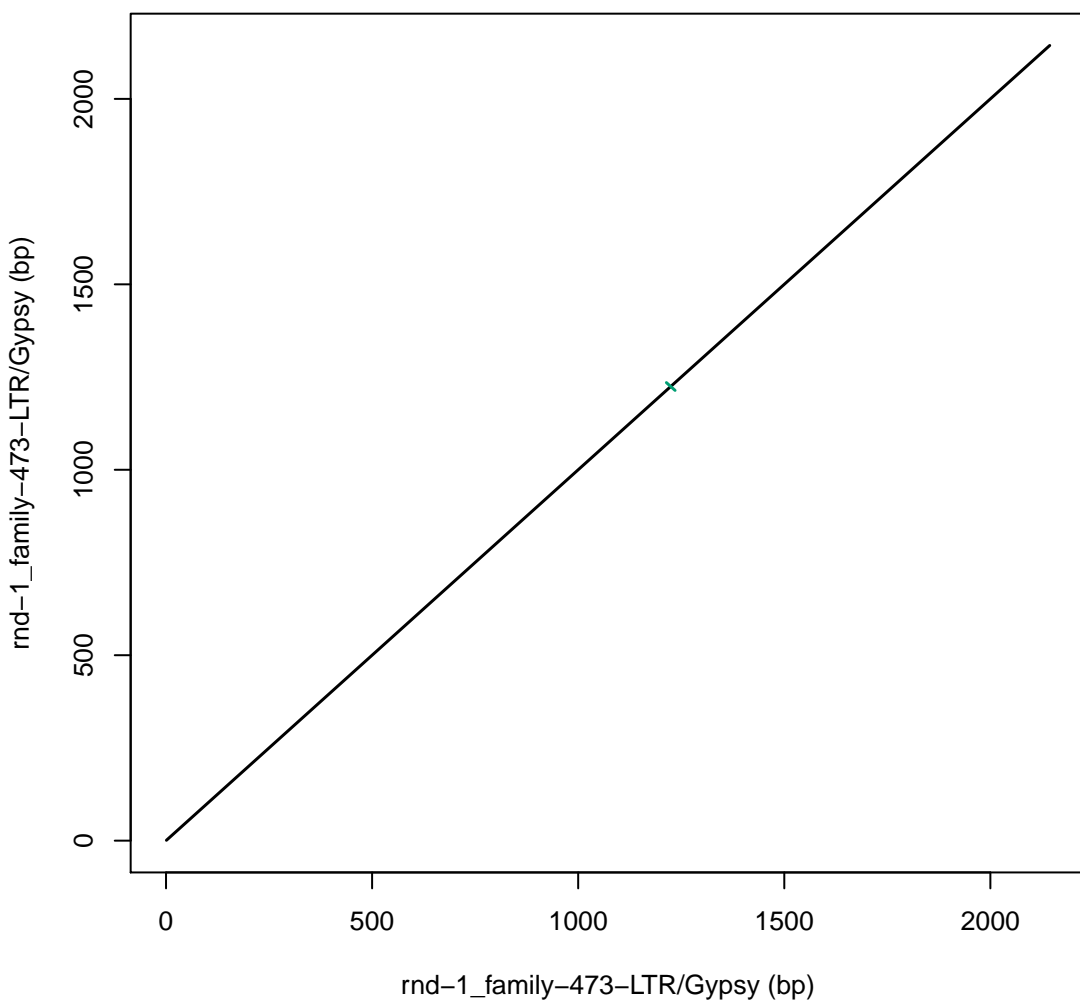
TE: rnd-1_family-473-LTR/Gypsy
consensus size: 2144bp; fragments: 88; full length: 2 (≥ 1929.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

