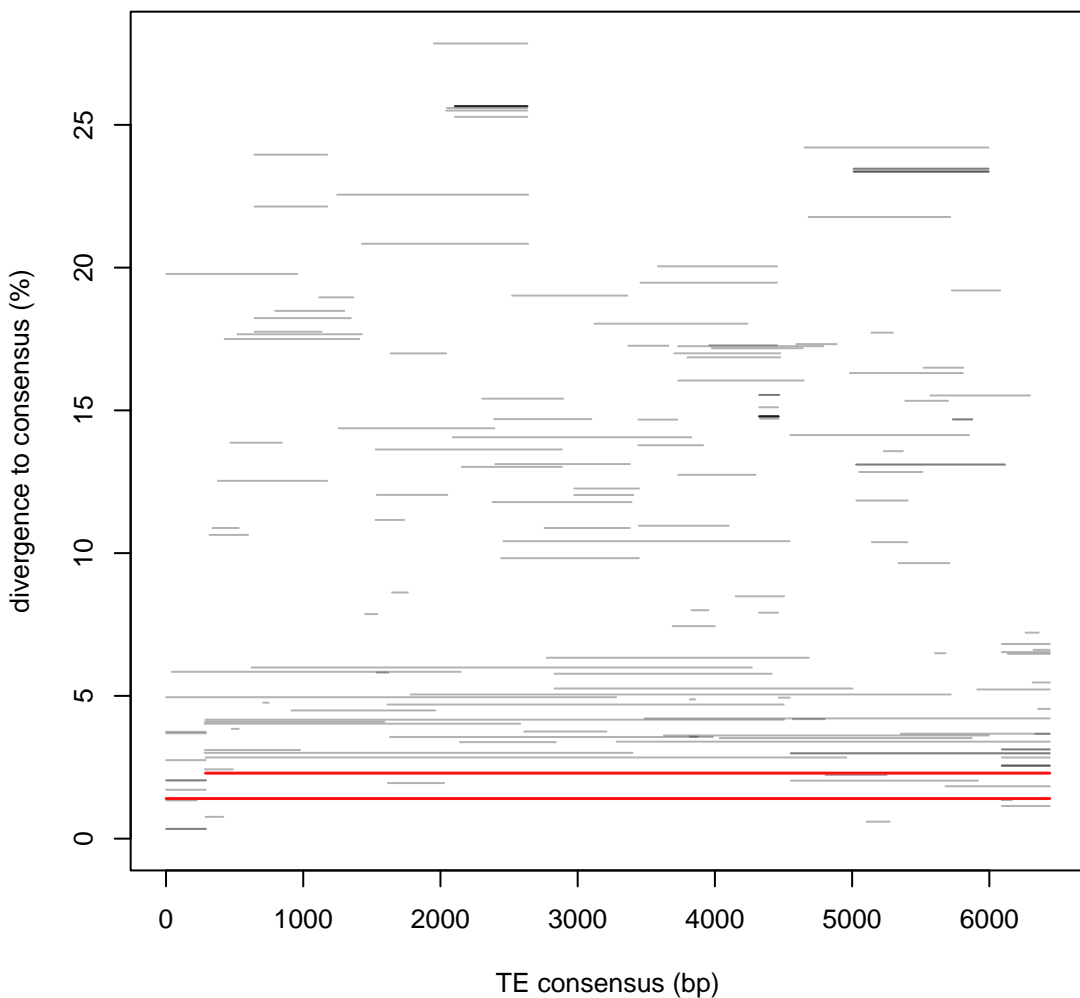
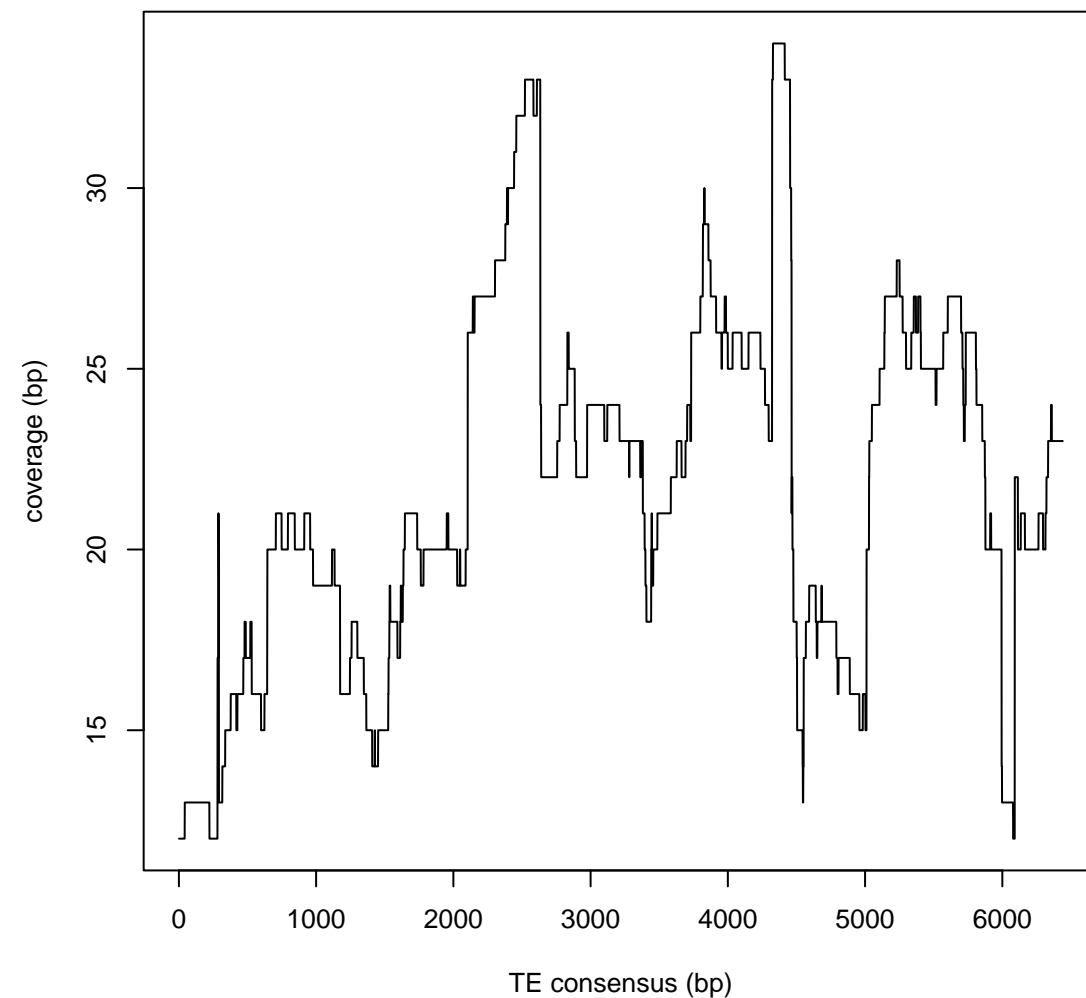


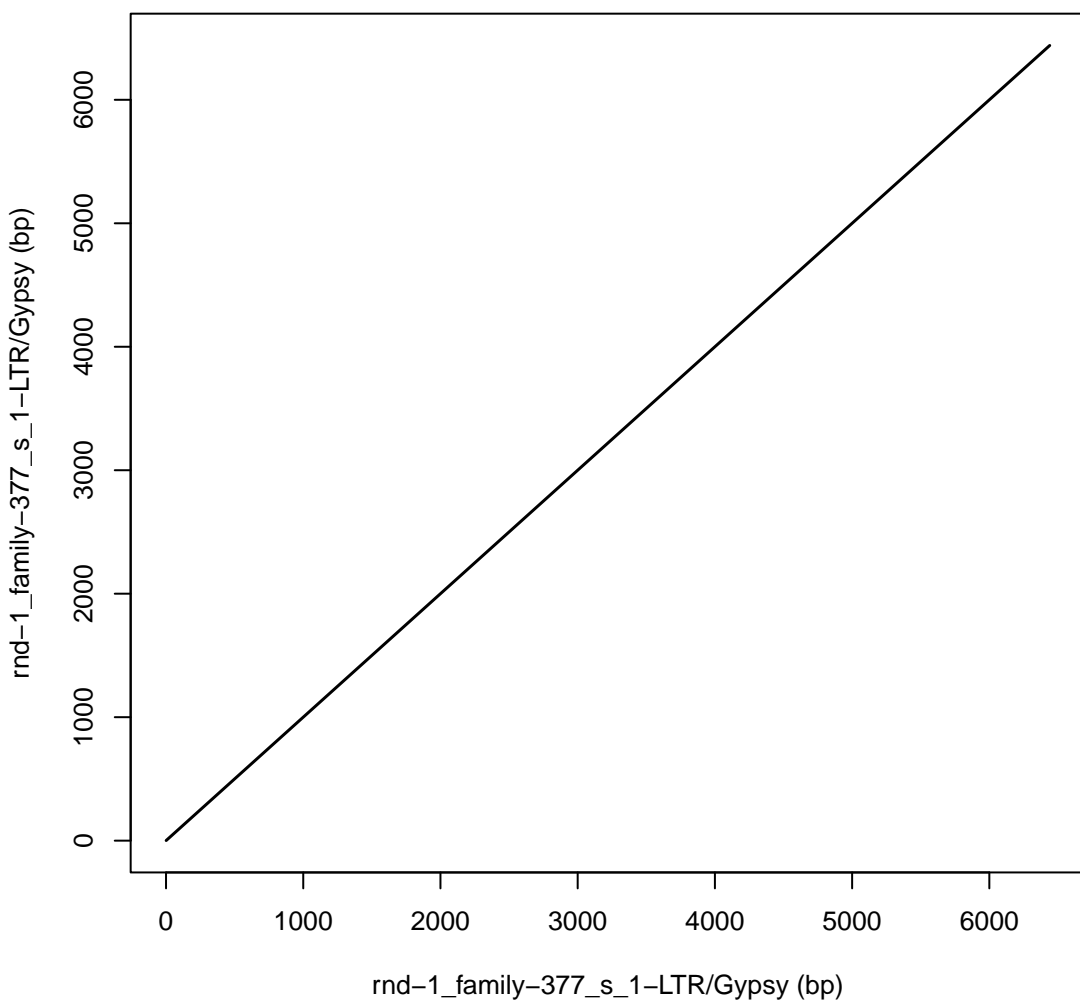
TE: rnd-1_family-377_s_1-LTR/Gypsy
consensus size: 6440bp; fragments: 163; full length: 2 (≥ 5796 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

