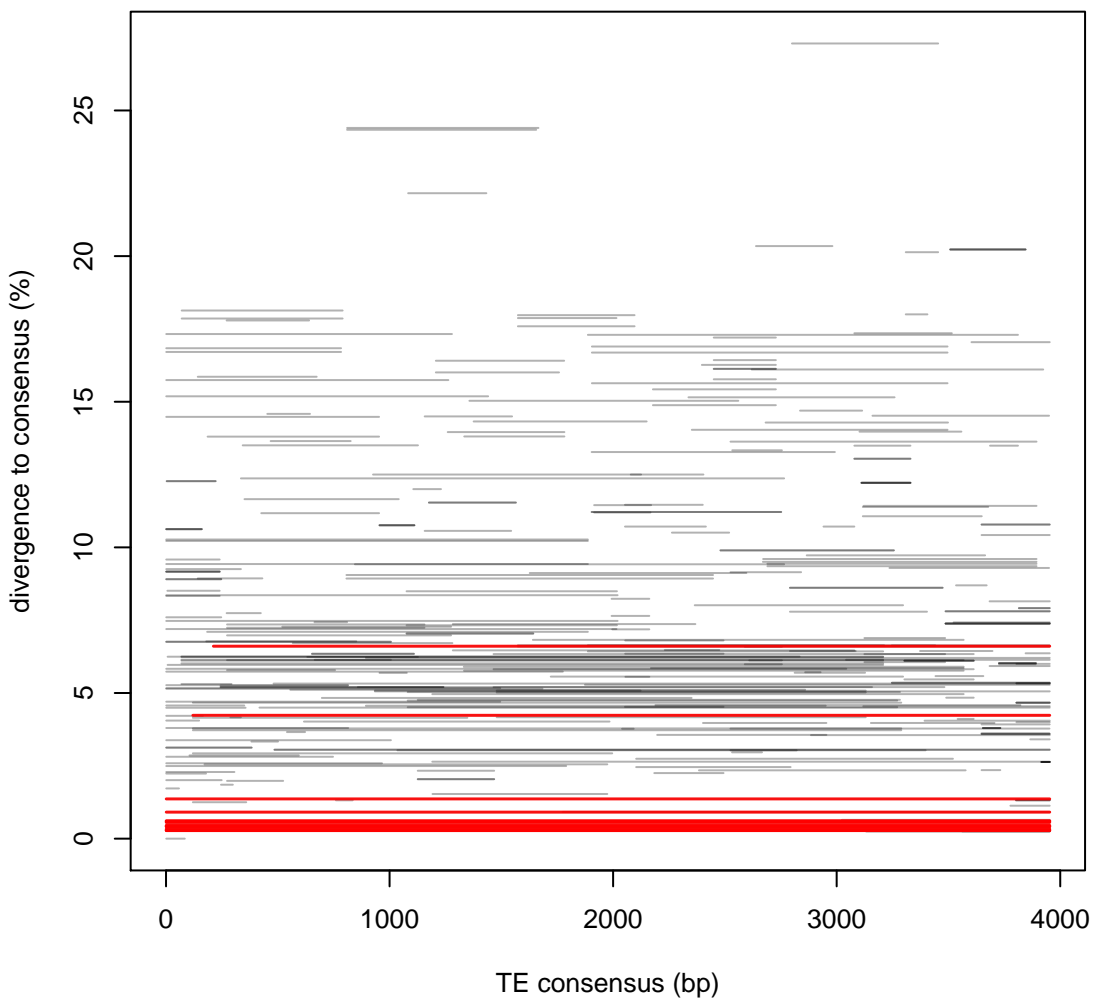
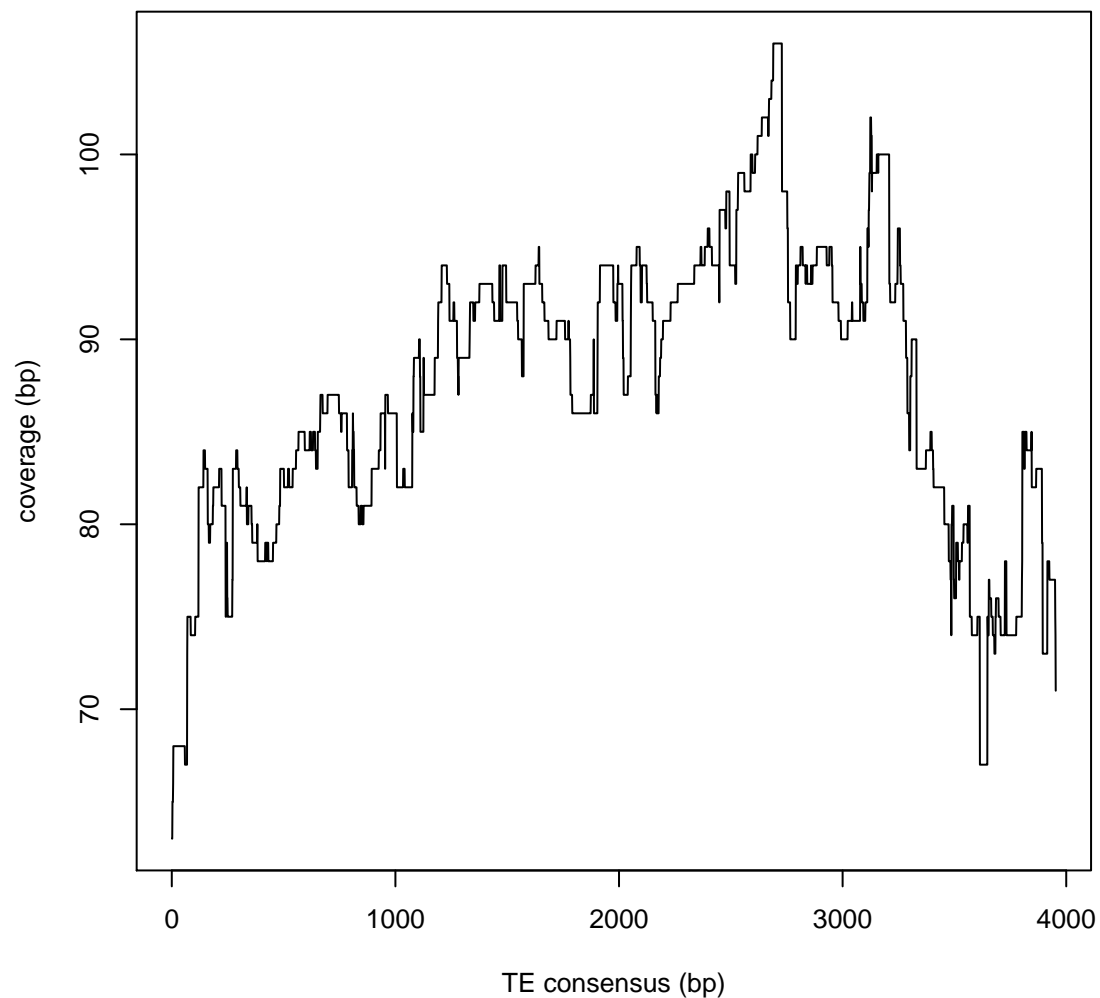


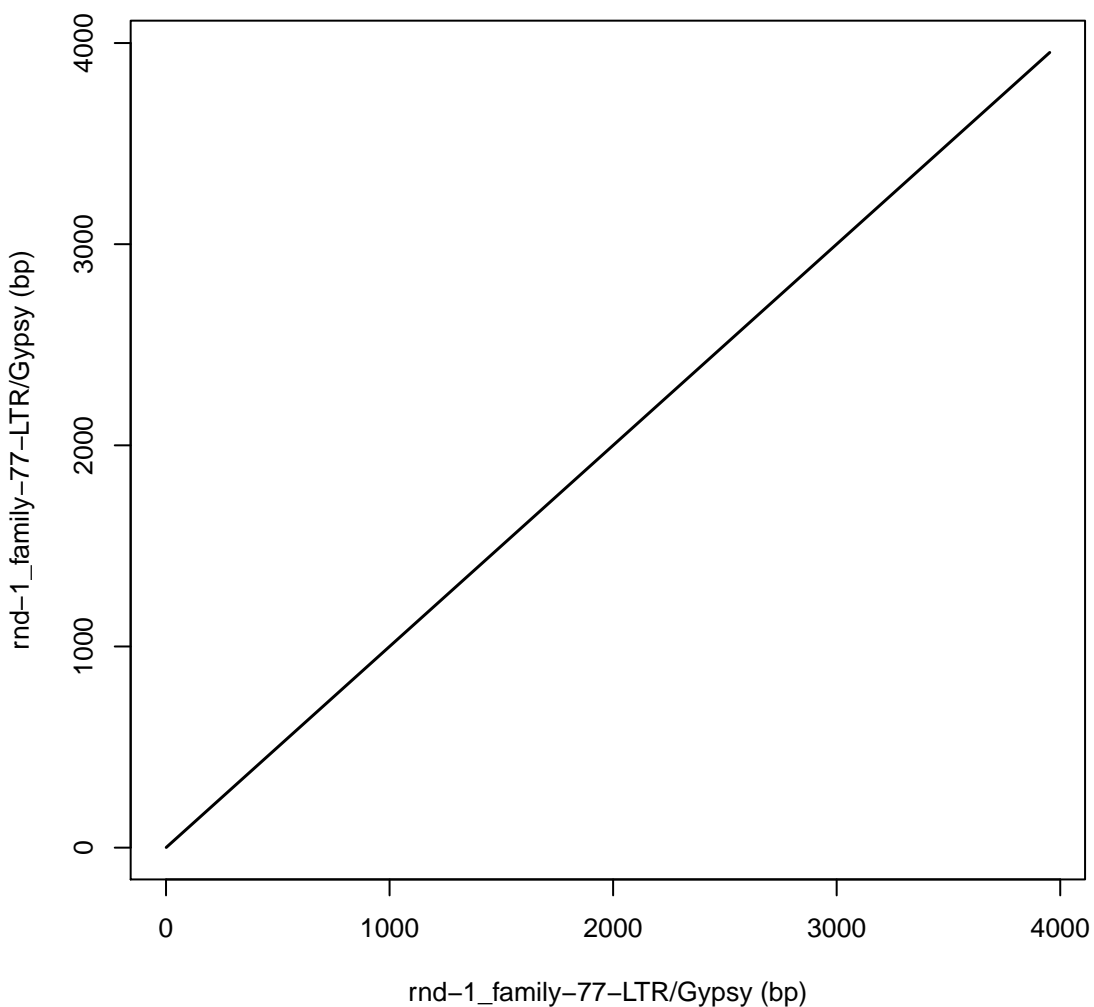
TE: rnd-1_family-77-LTR/Gypsy
consensus size: 3953bp; fragments: 349; full length: 16 (≥ 3557.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

