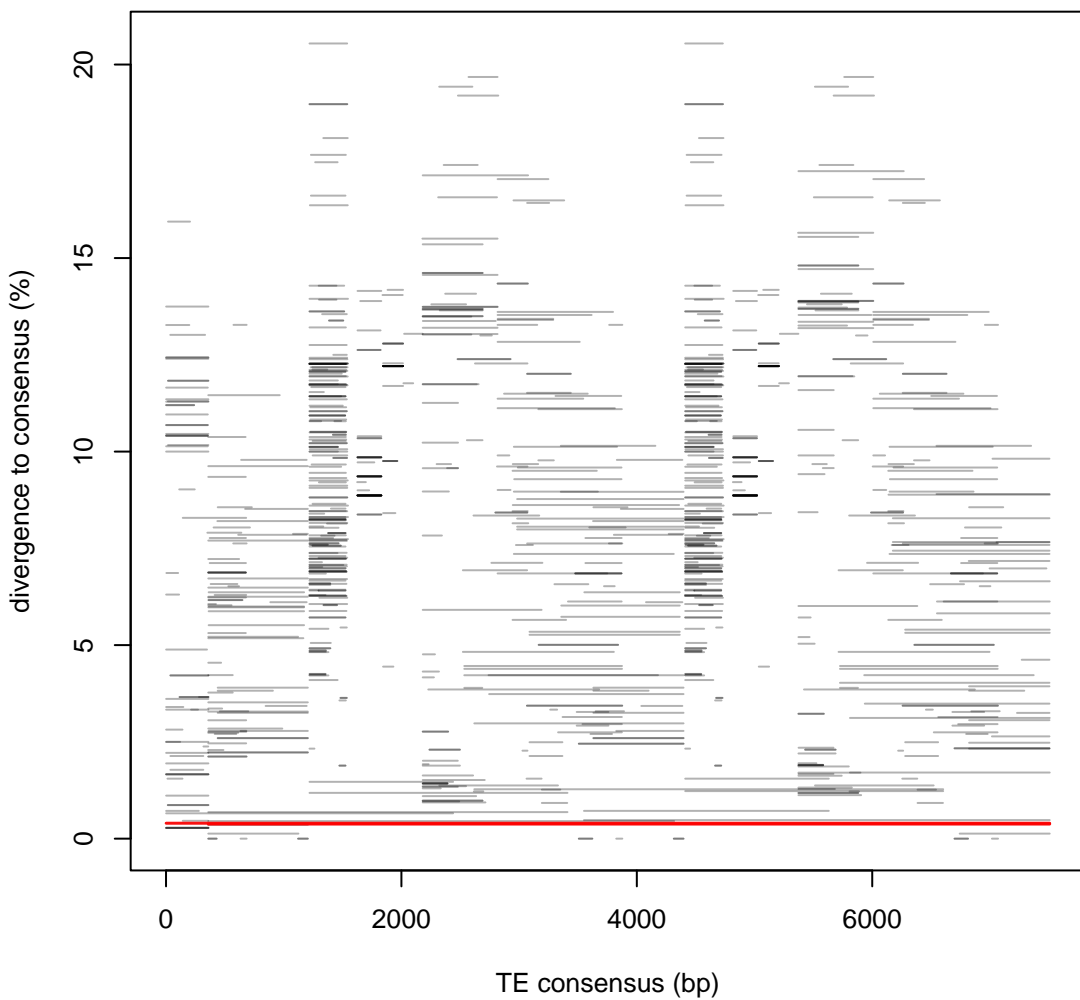
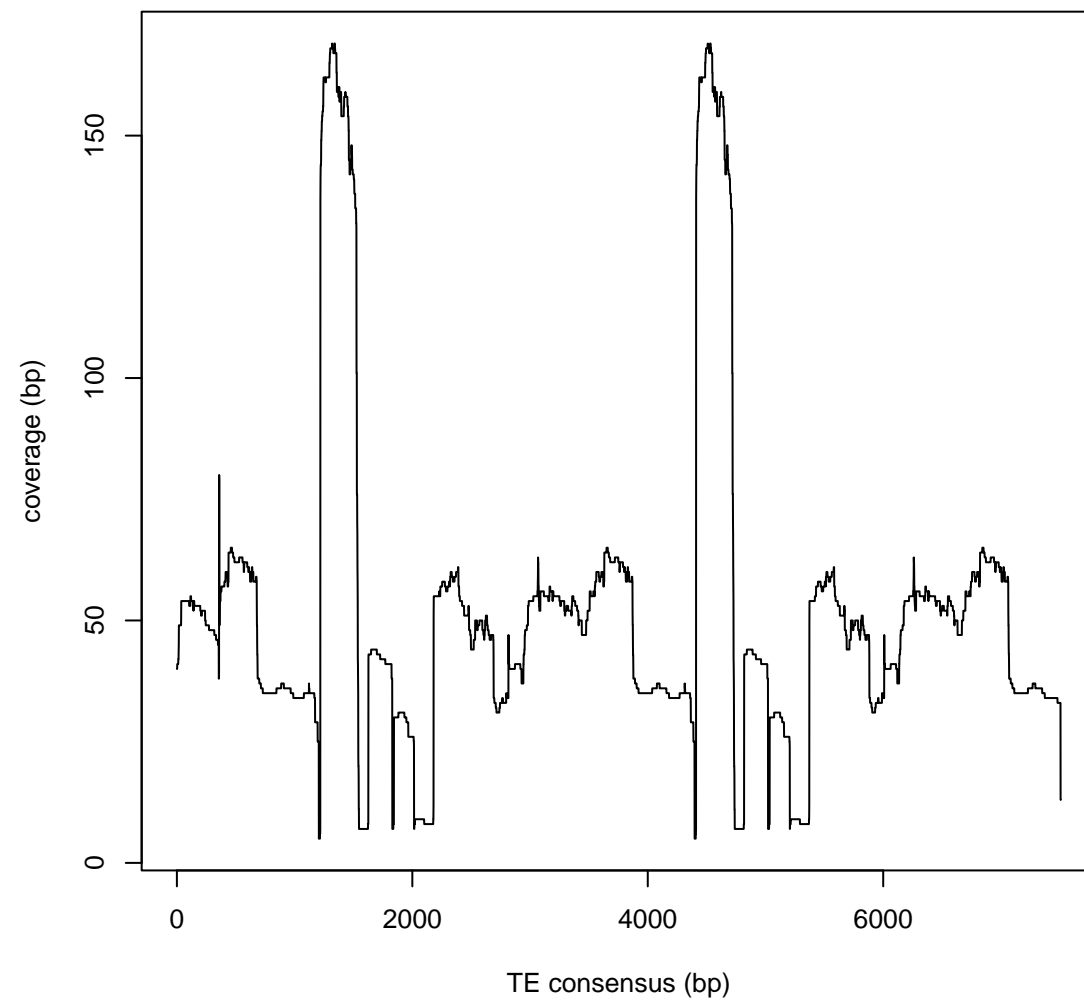


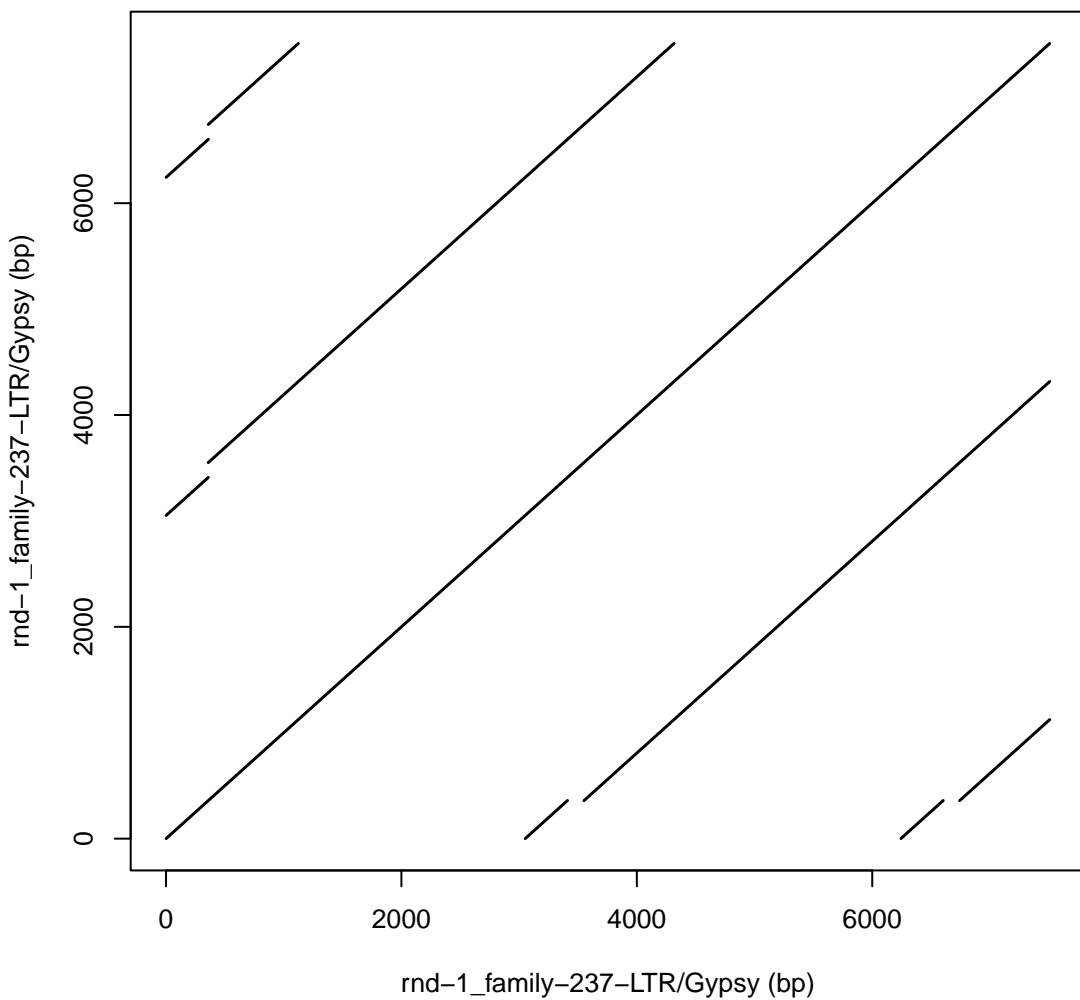
TE: rnd-1_family-237-LTR/Gypsy
consensus size: 7508bp; fragments: 1004; full length: 2 (≥ 6757.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

