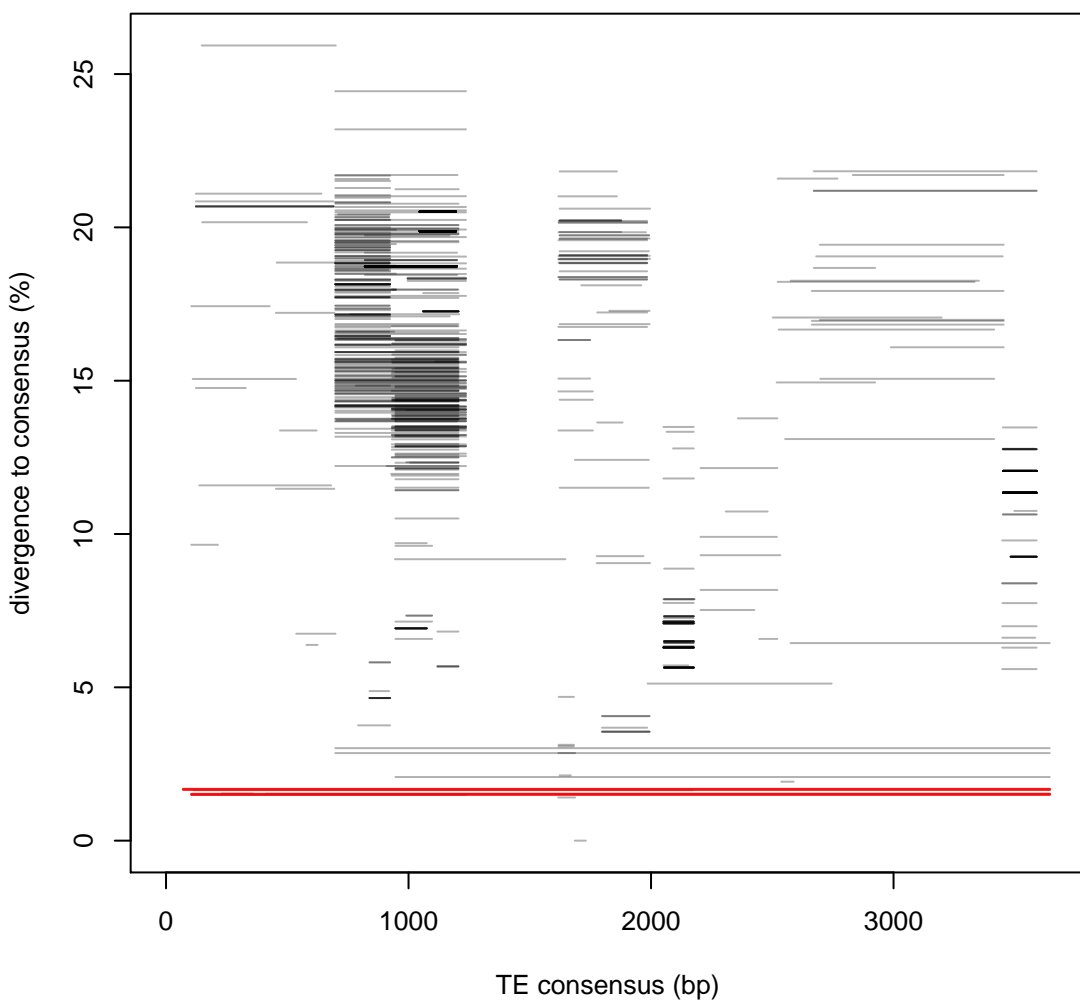
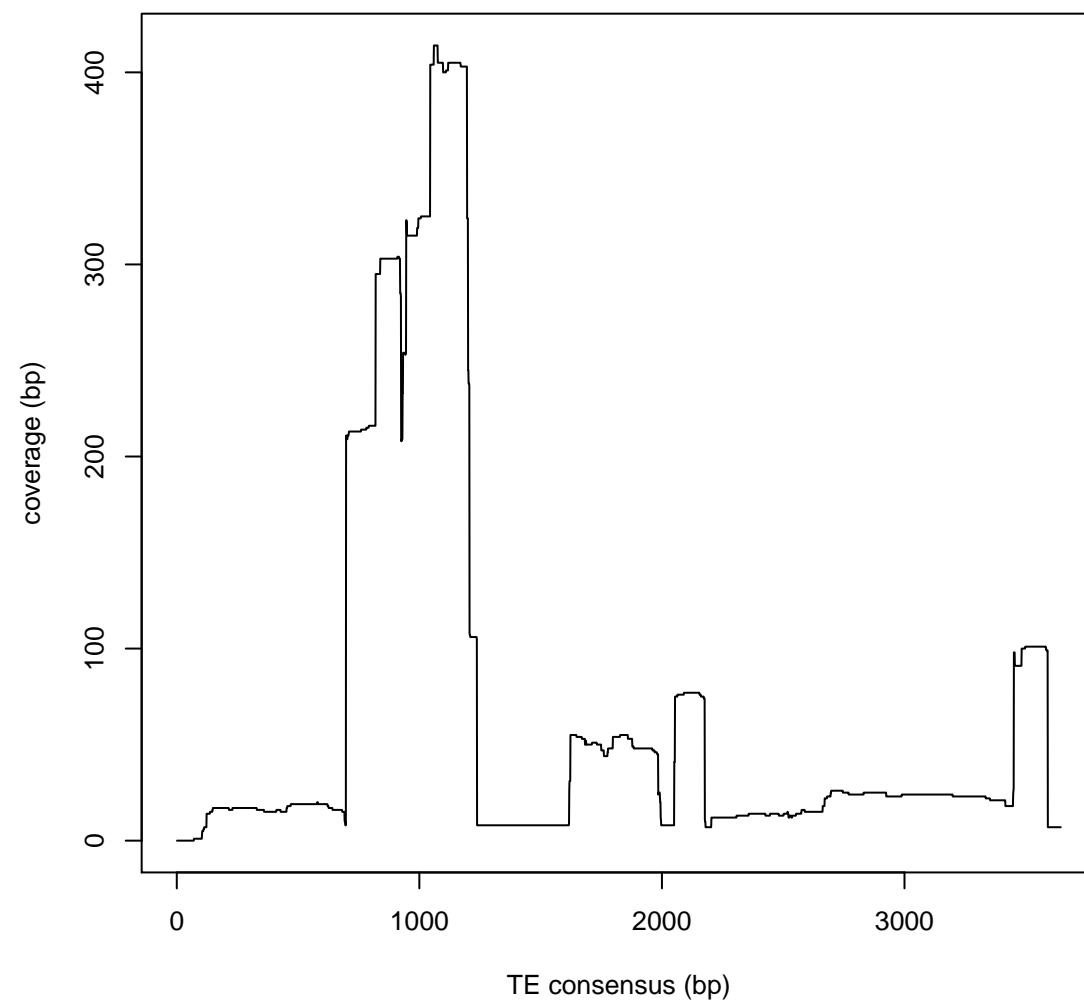


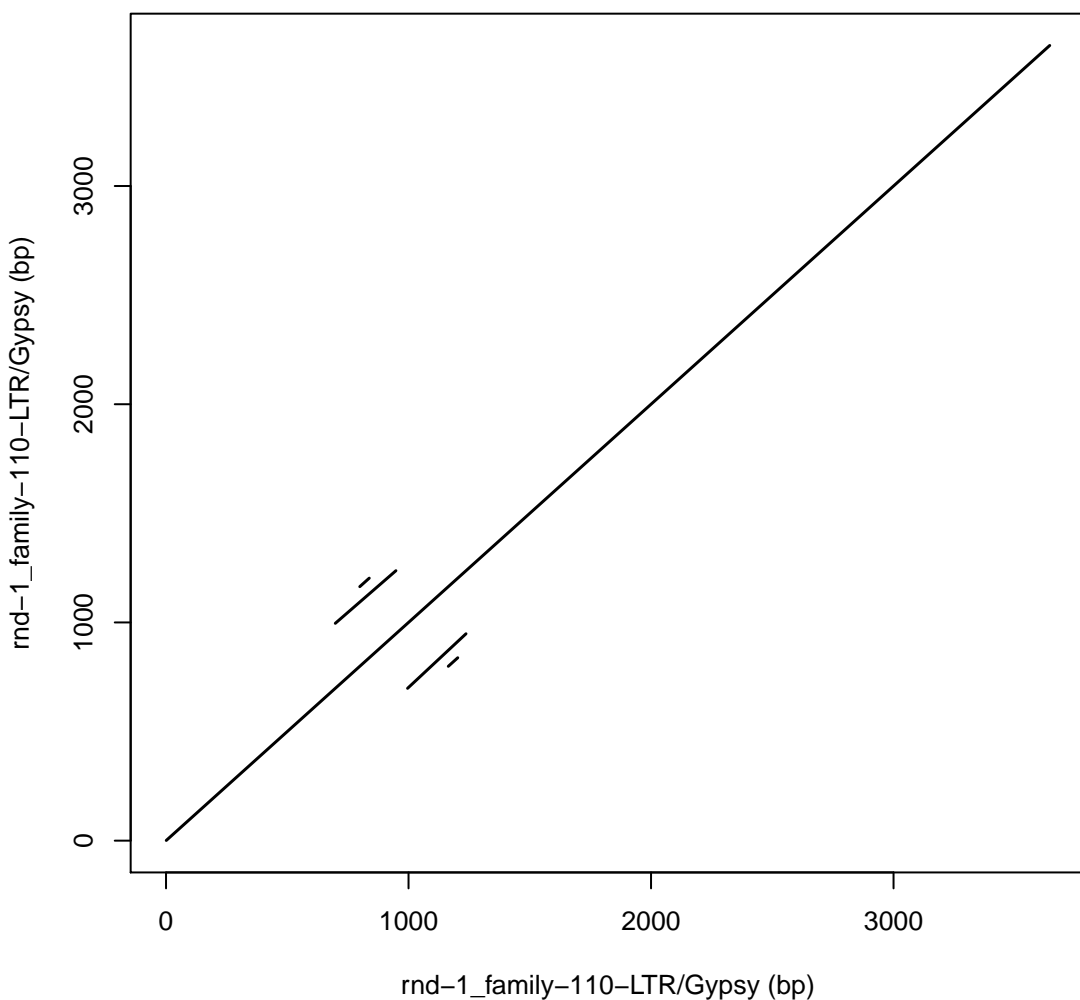
TE: rnd-1_family-110-LTR/Gypsy
consensus size: 3644bp; fragments: 796; full length: 2 (≥ 3279.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

