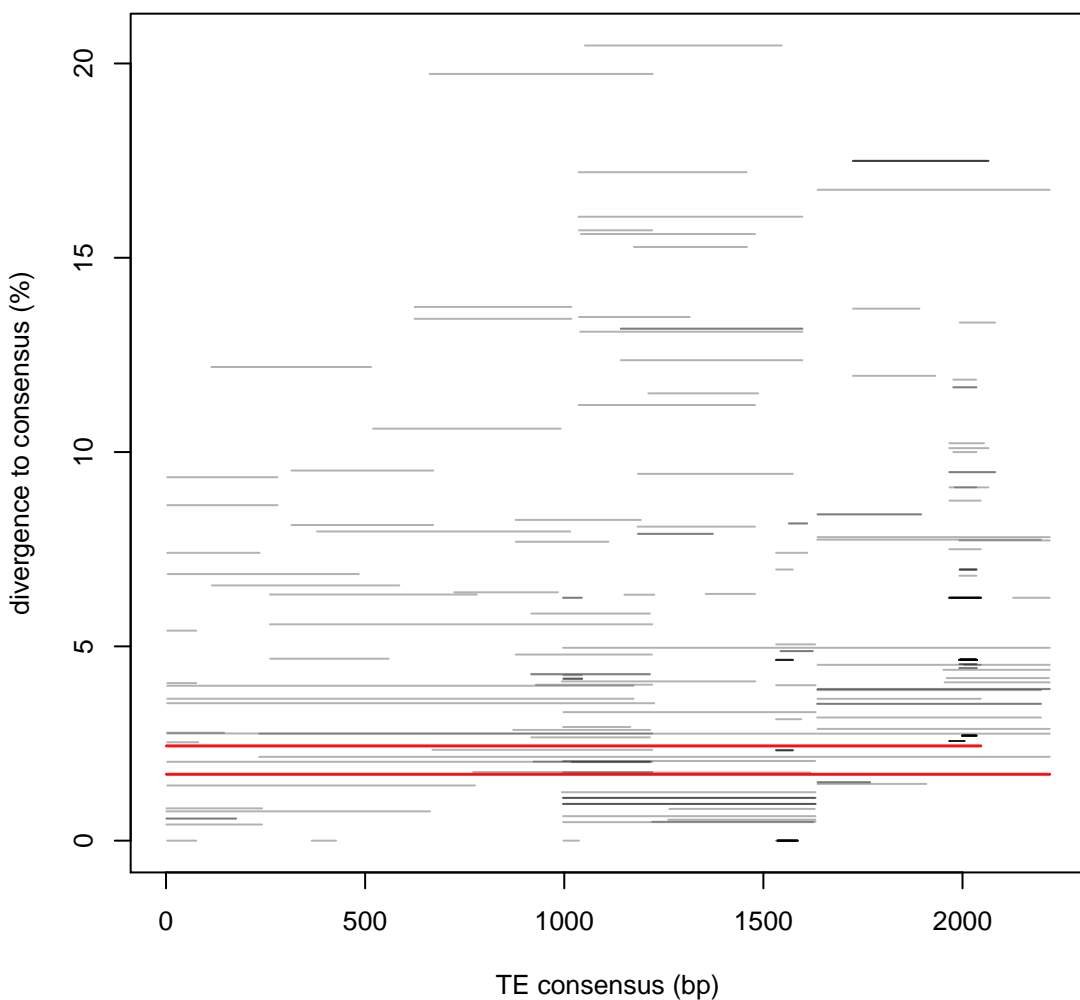
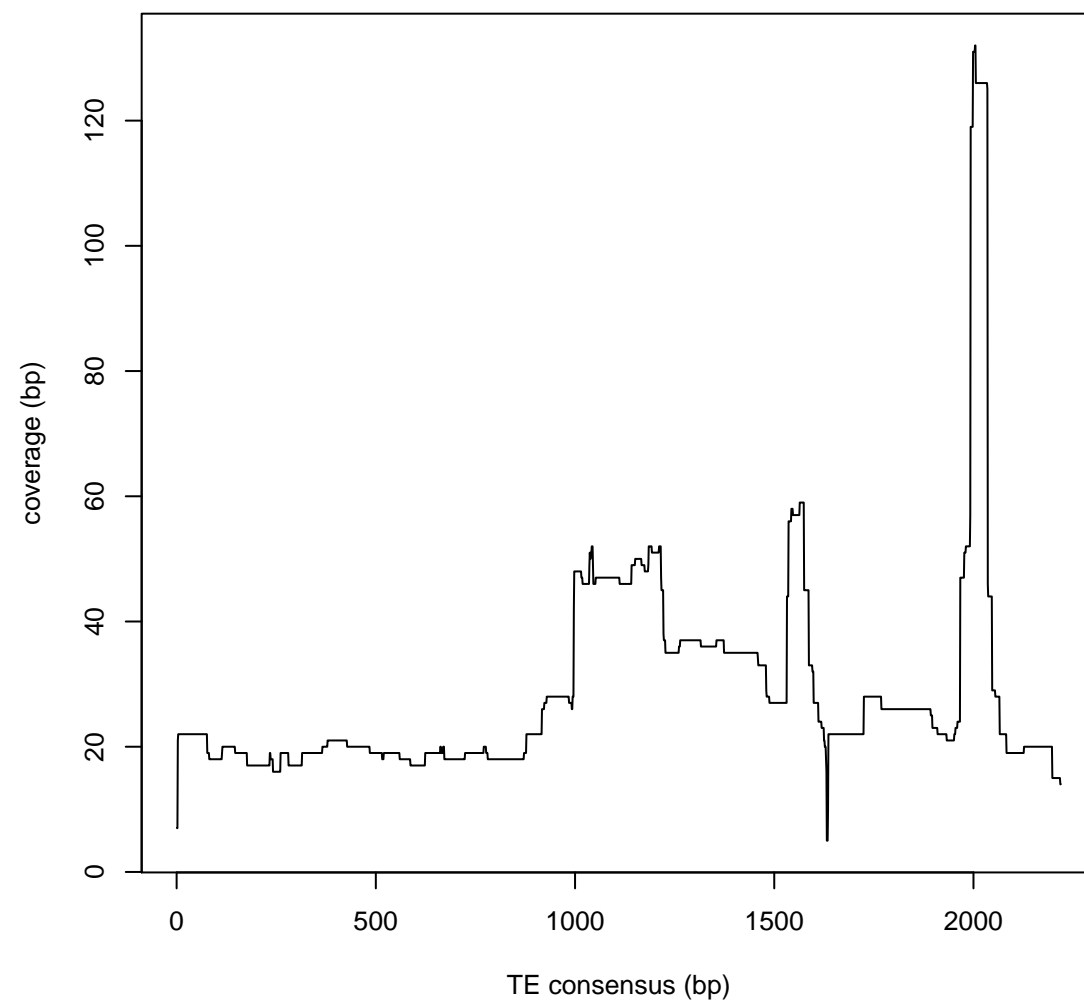


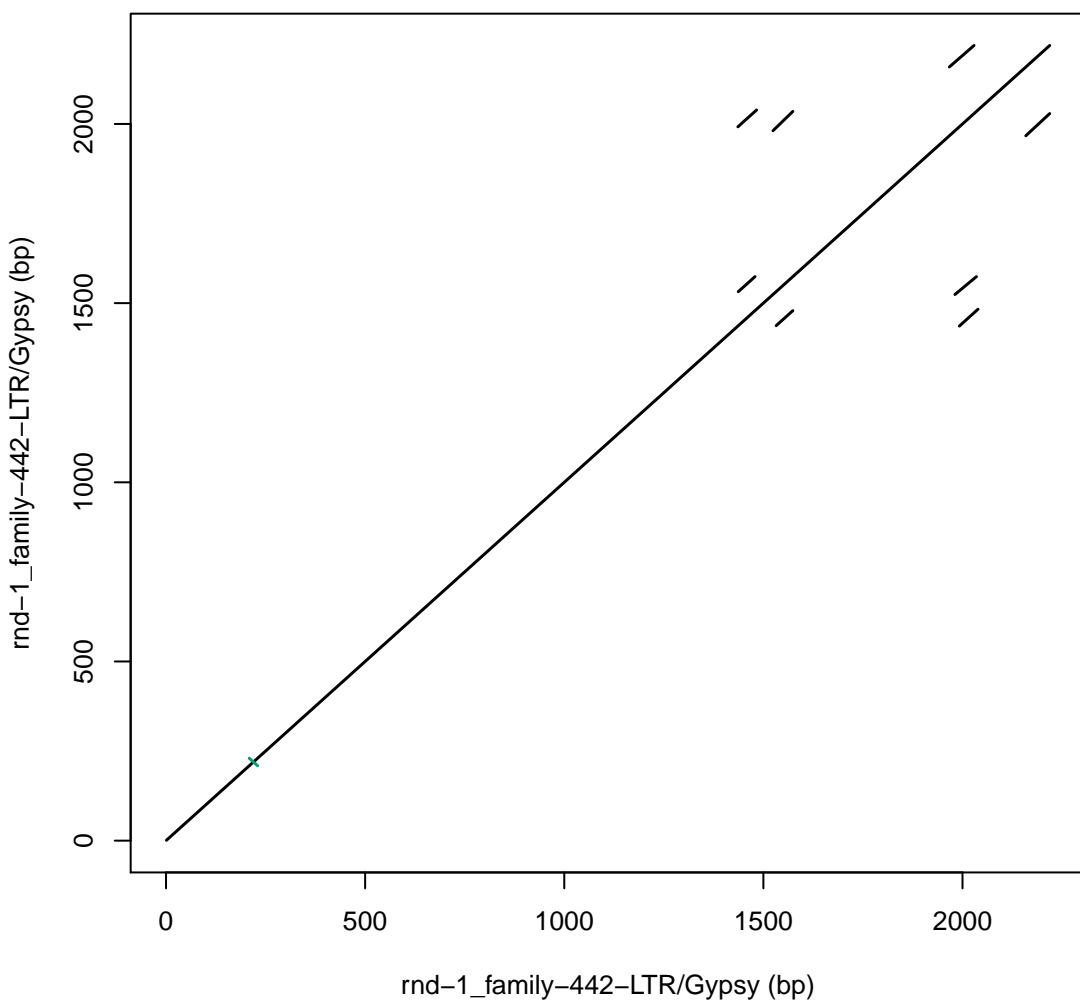
TE: rnd-1_family-442-LTR/Gypsy
consensus size: 2219bp; fragments: 263; full length: 2 (≥ 1997.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

