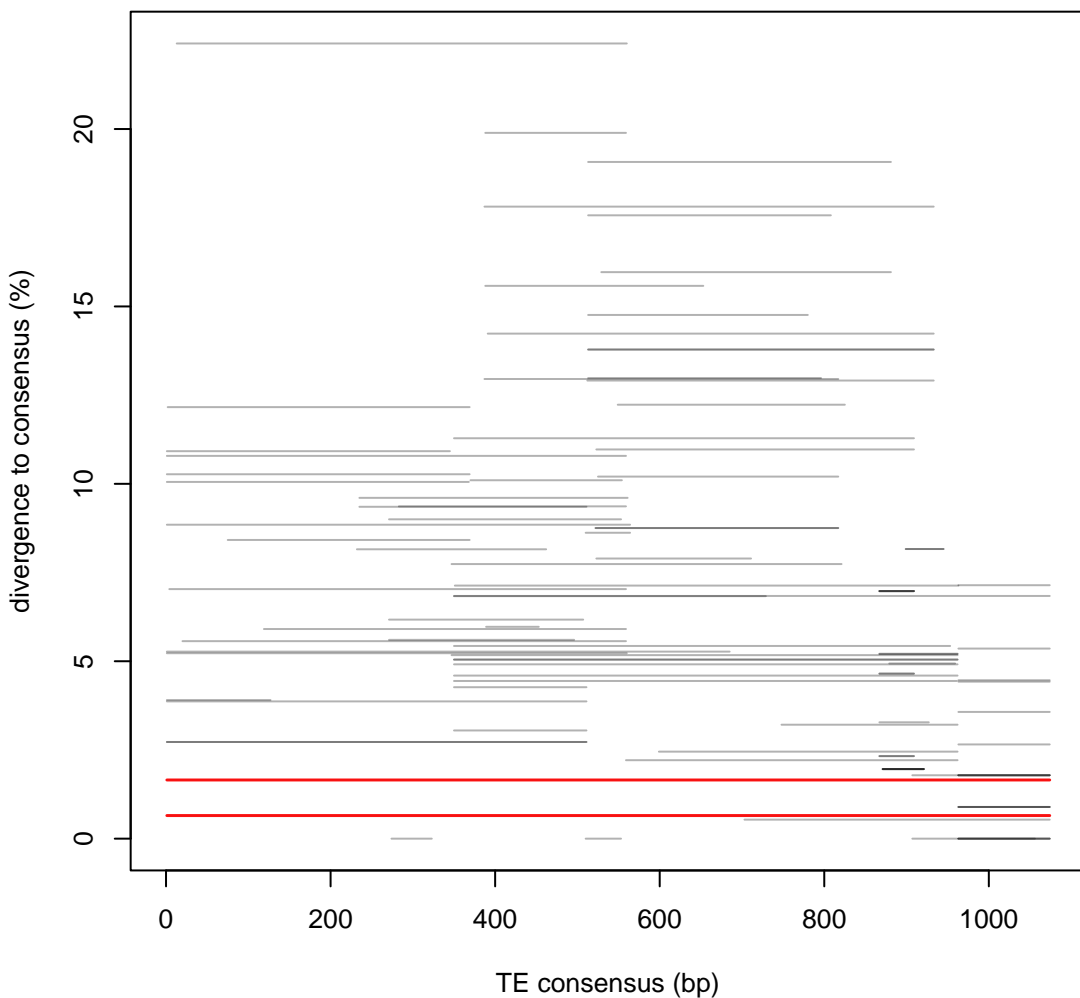
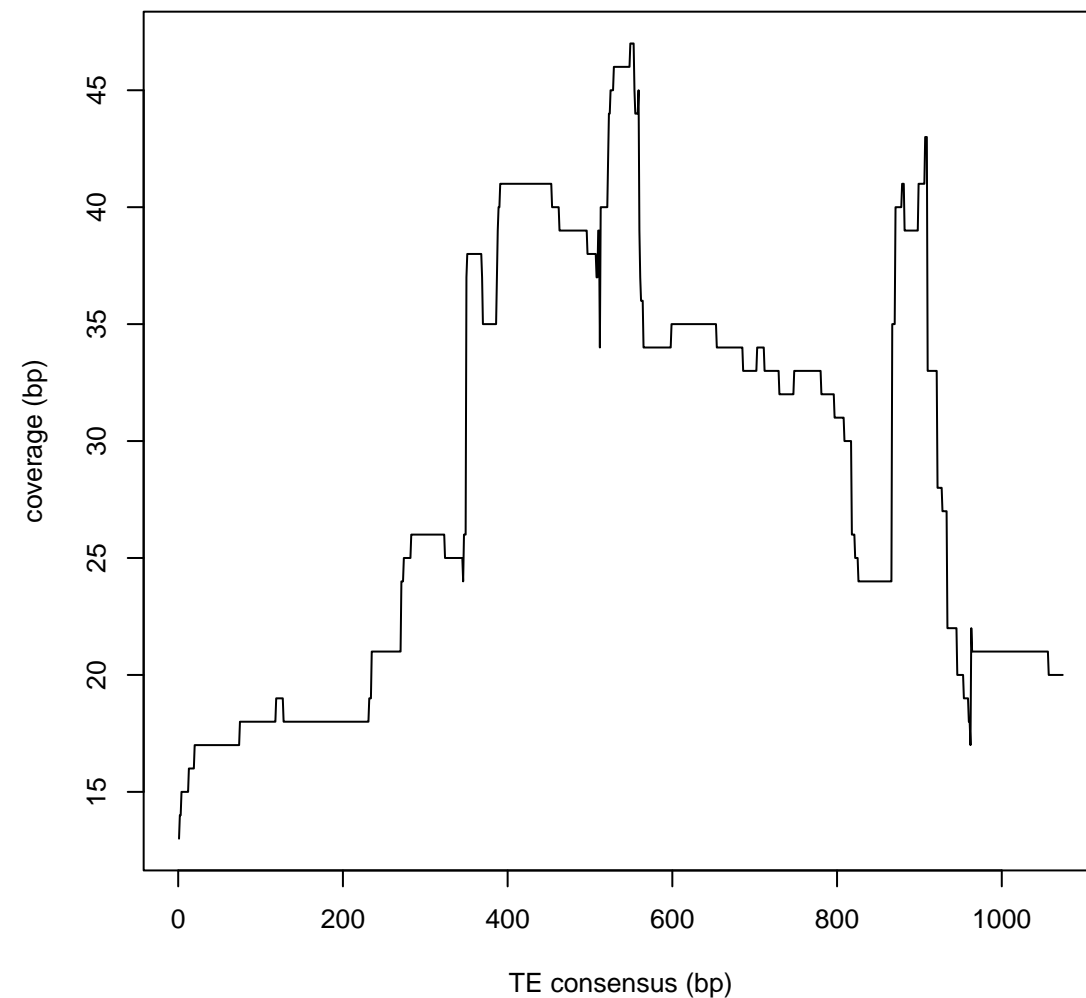


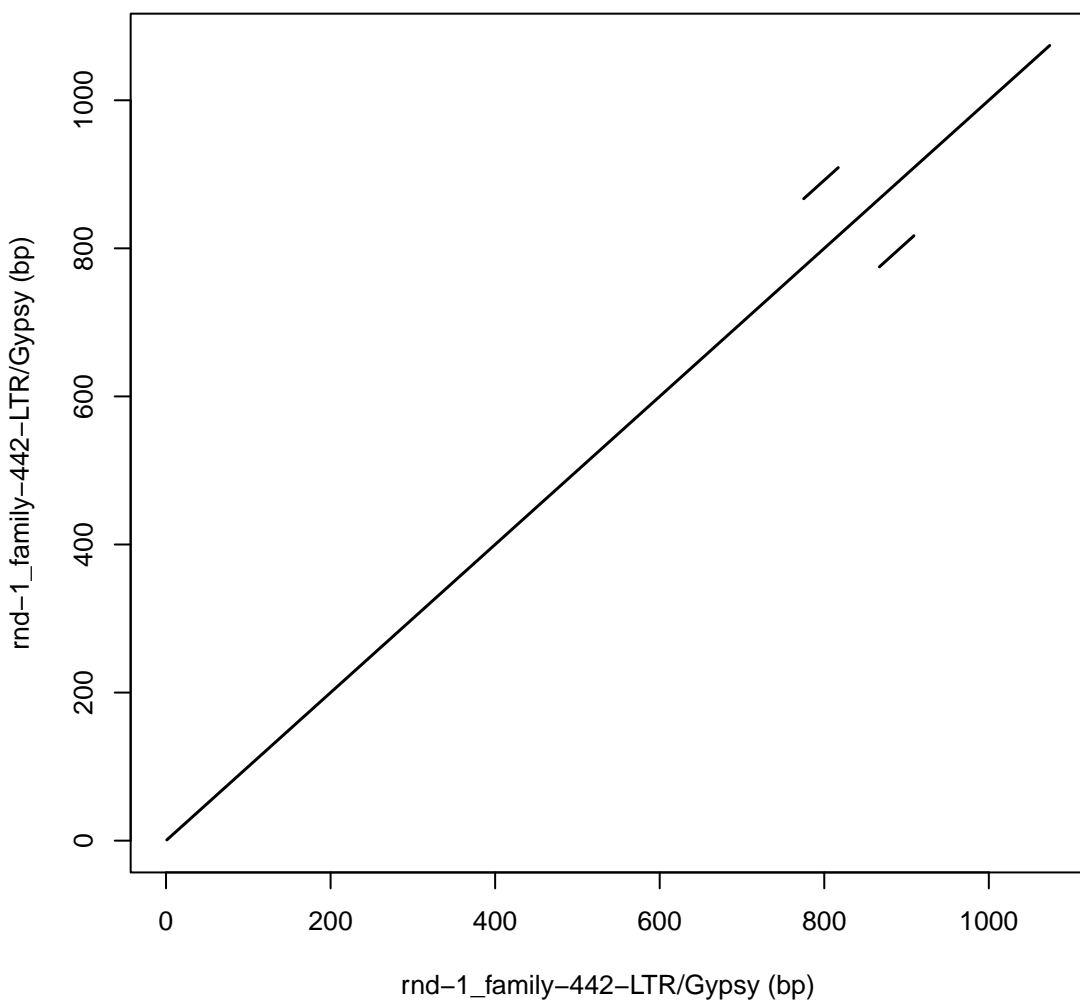
TE: rnd-1_family-442-LTR/Gypsy
consensus size: 1074bp; fragments: 106; full length: 2 (>=966.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

