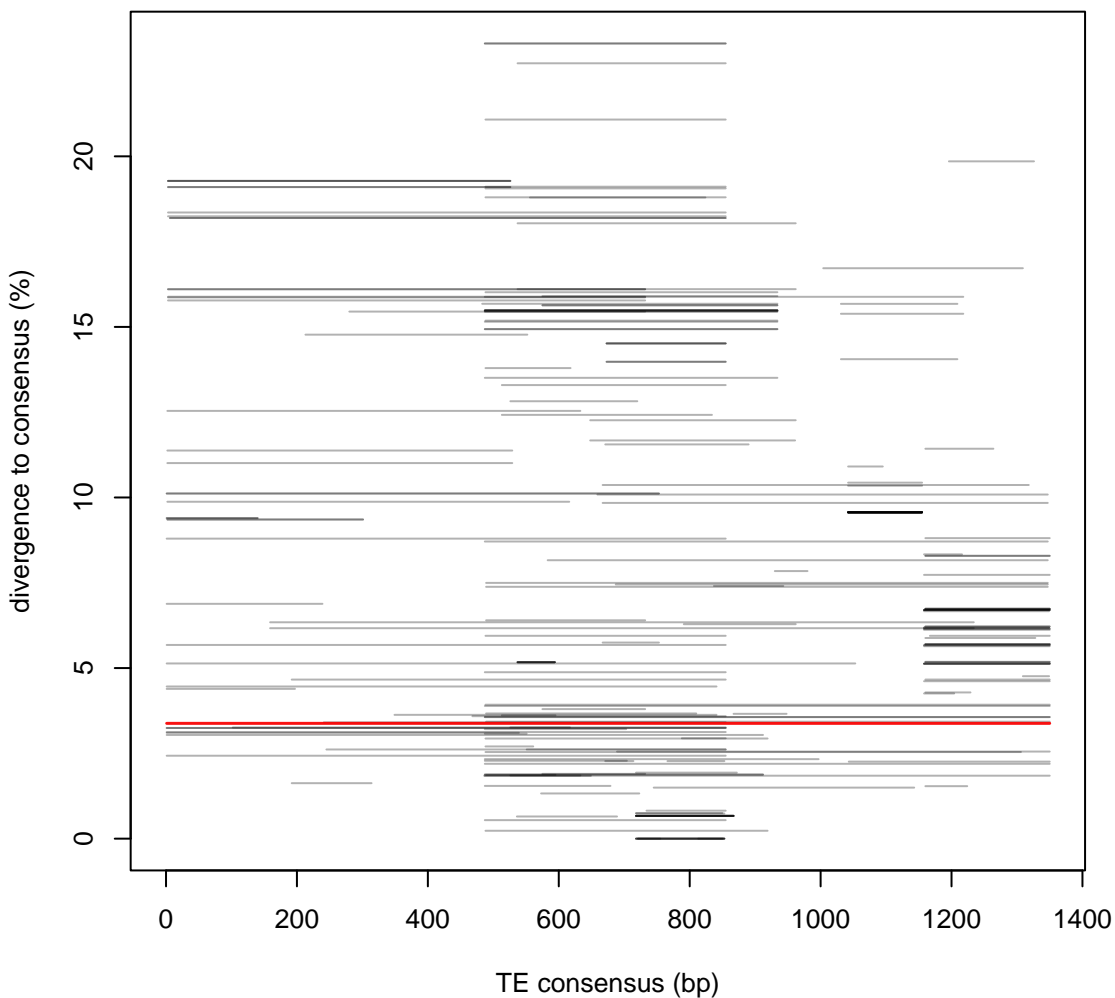
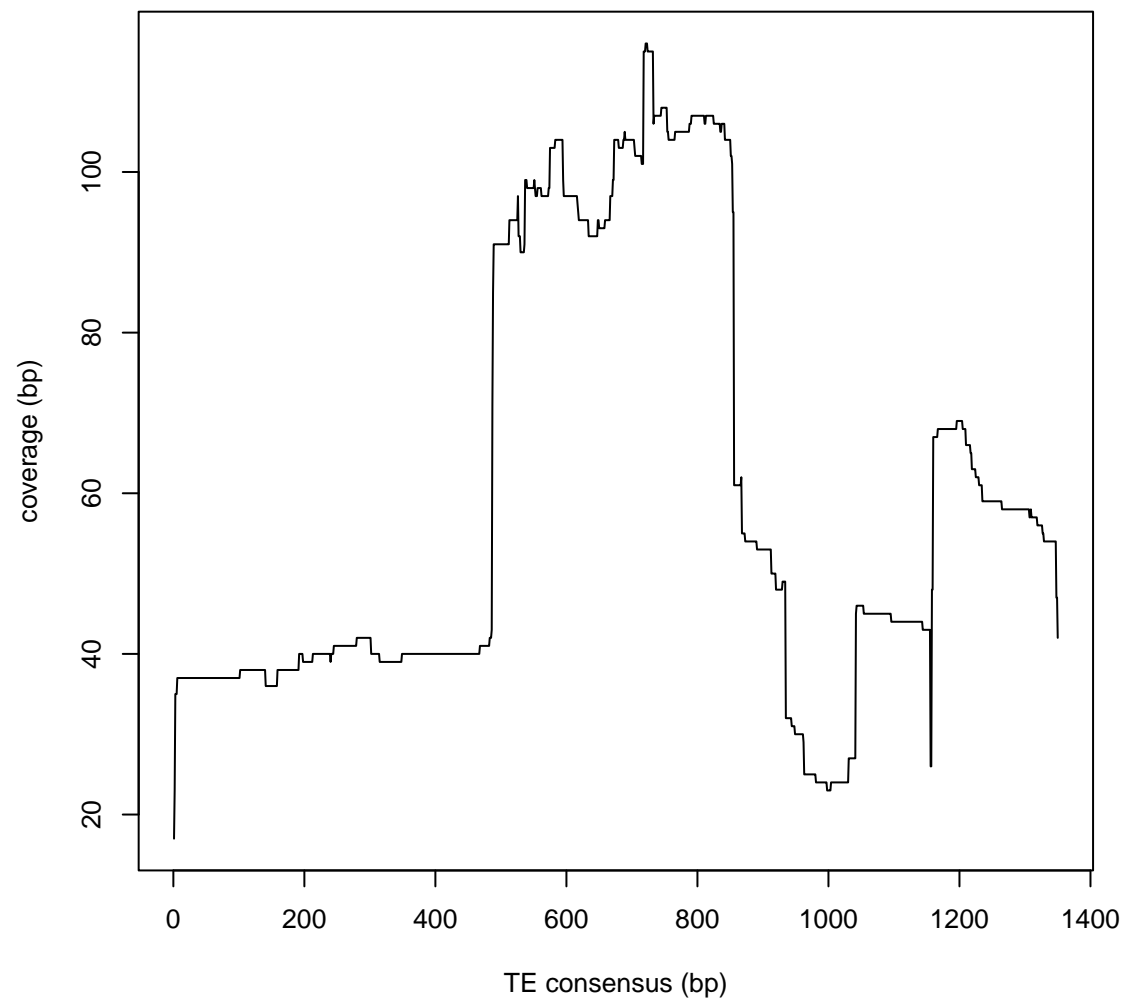


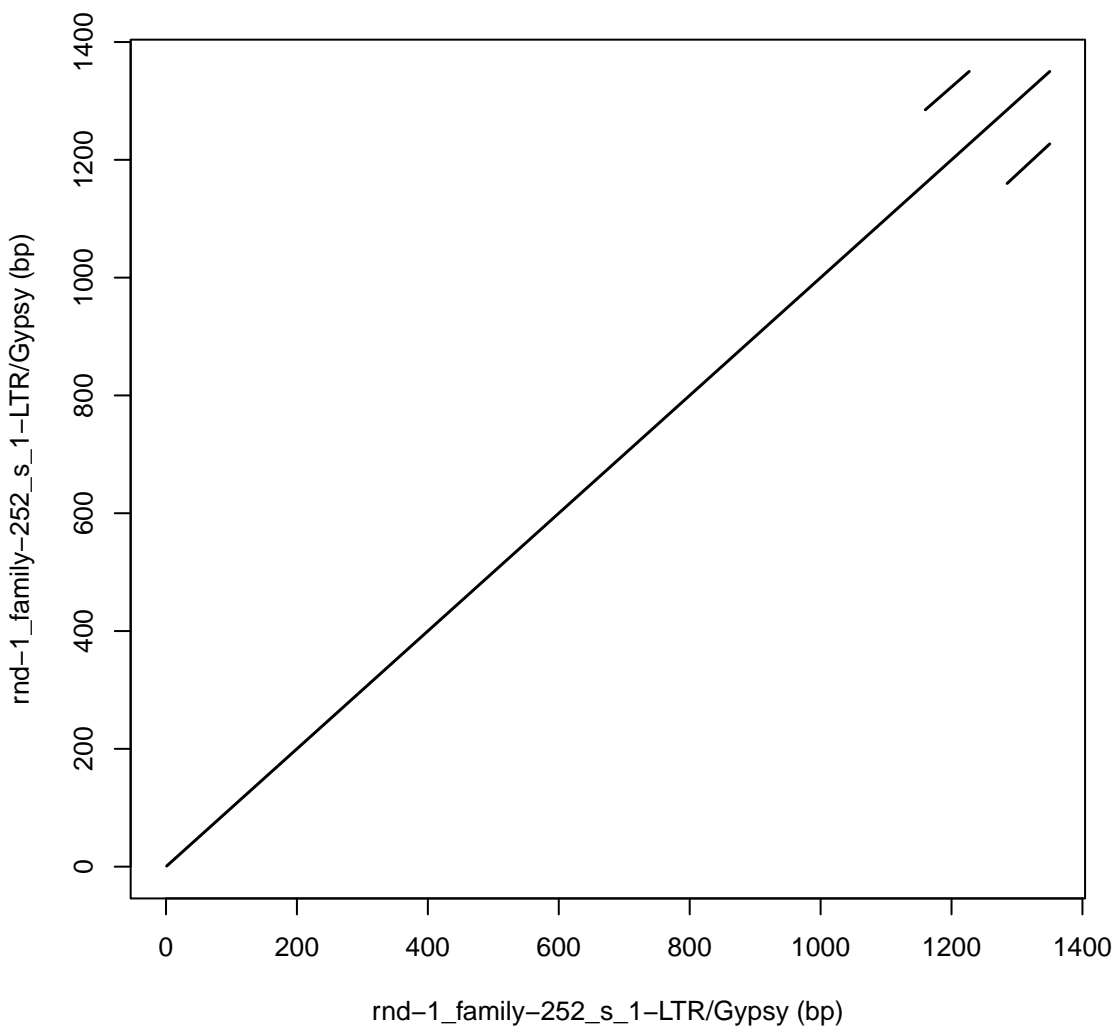
TE: rnd-1_family-252_s_1-LTR/Gypsy
consensus size: 1350bp; fragments: 229; full length: 1 (>=1215bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

