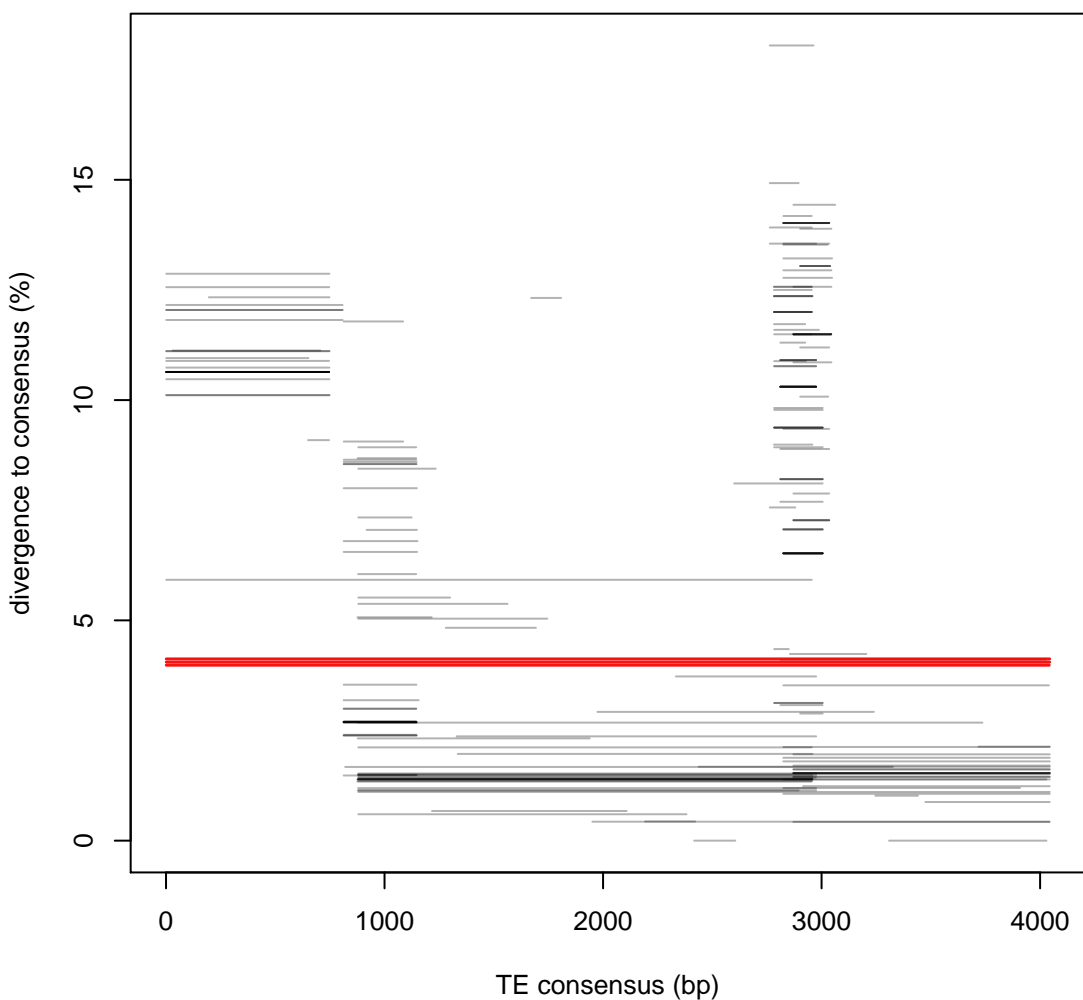
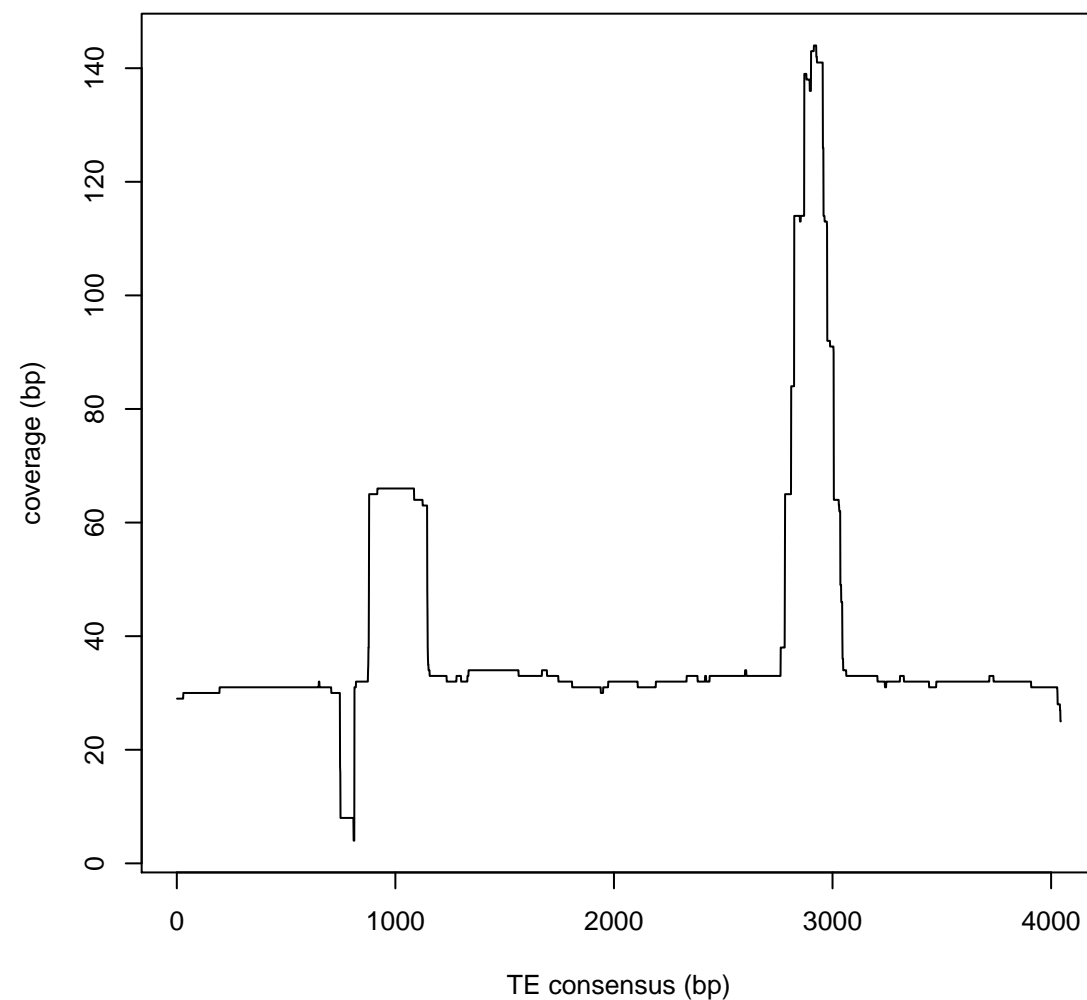


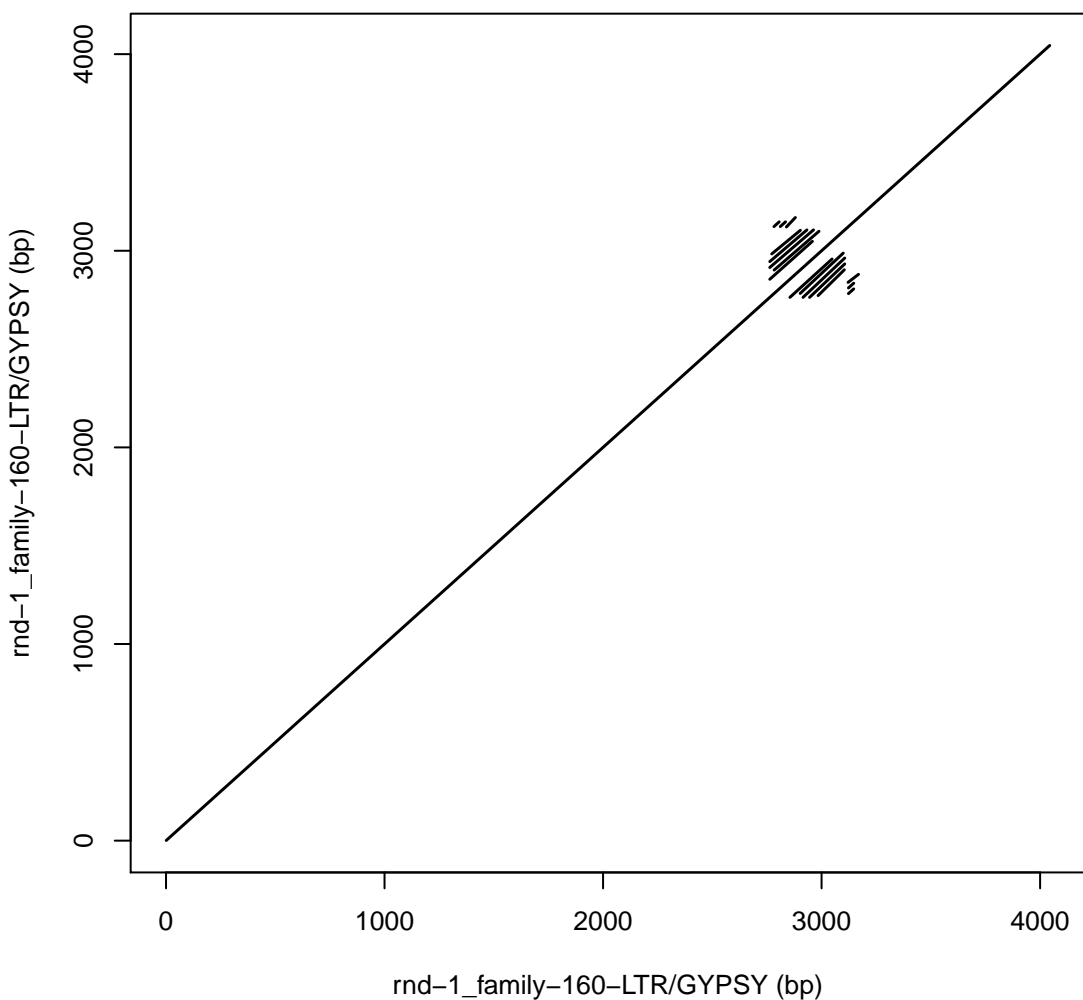
TE: rnd-1_family-160-LTR/GYPSY
consensus size: 4044bp; fragments: 225; full length: 3 (>=3639.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

