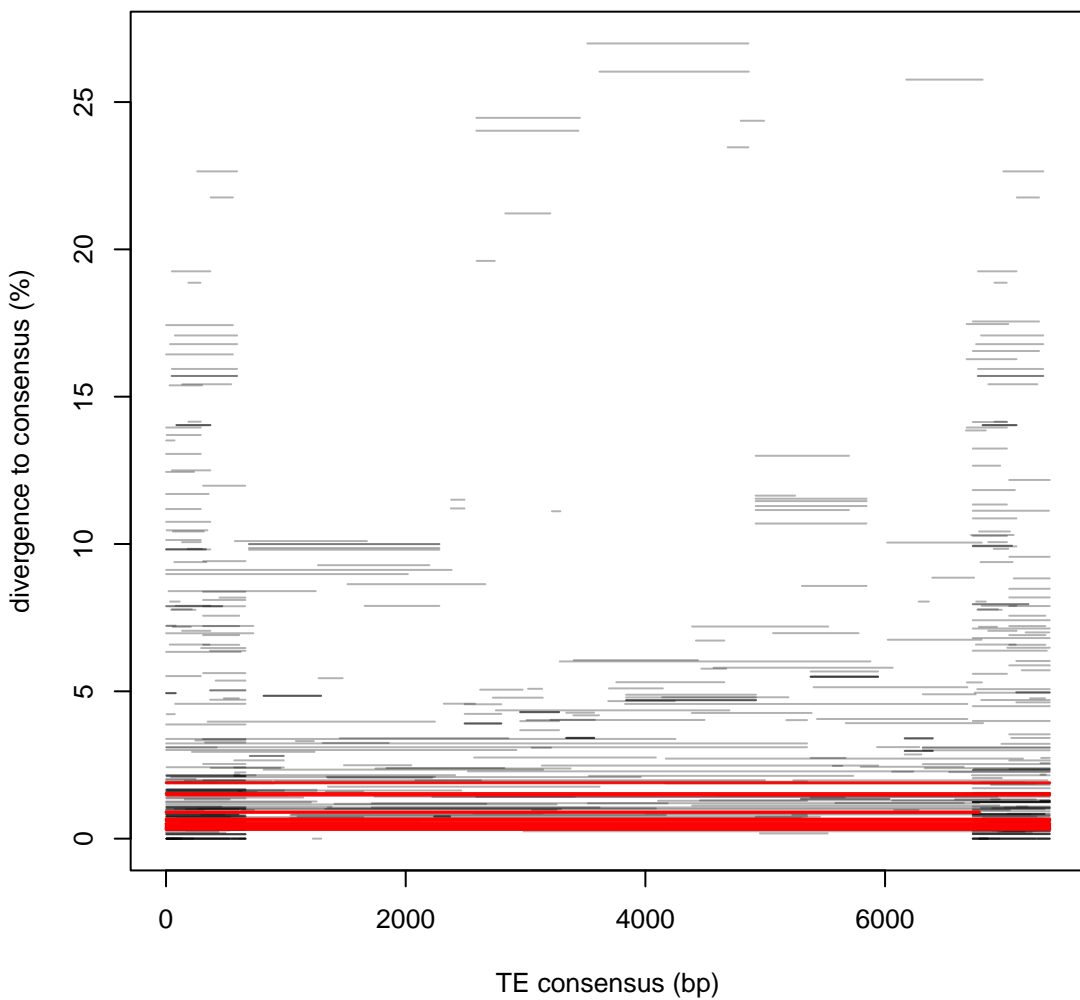
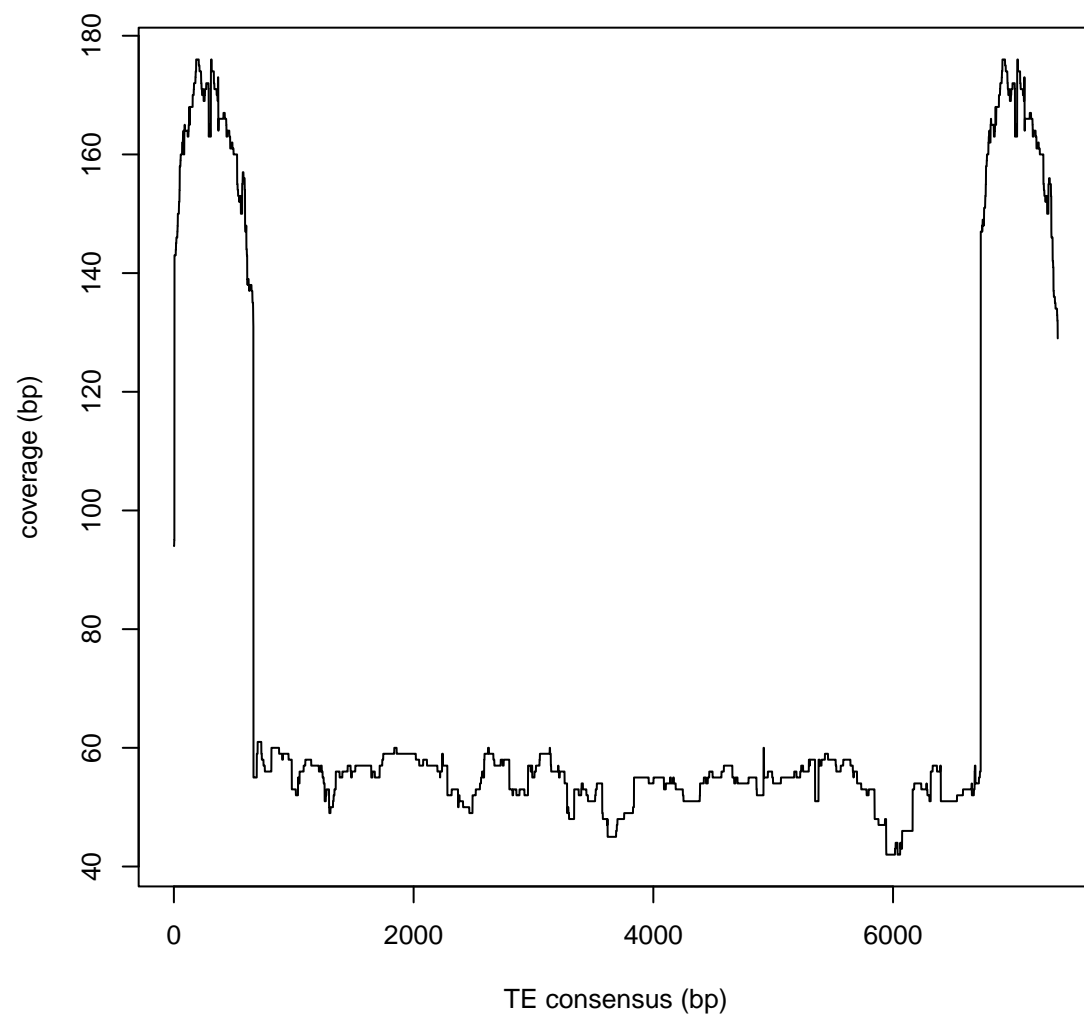


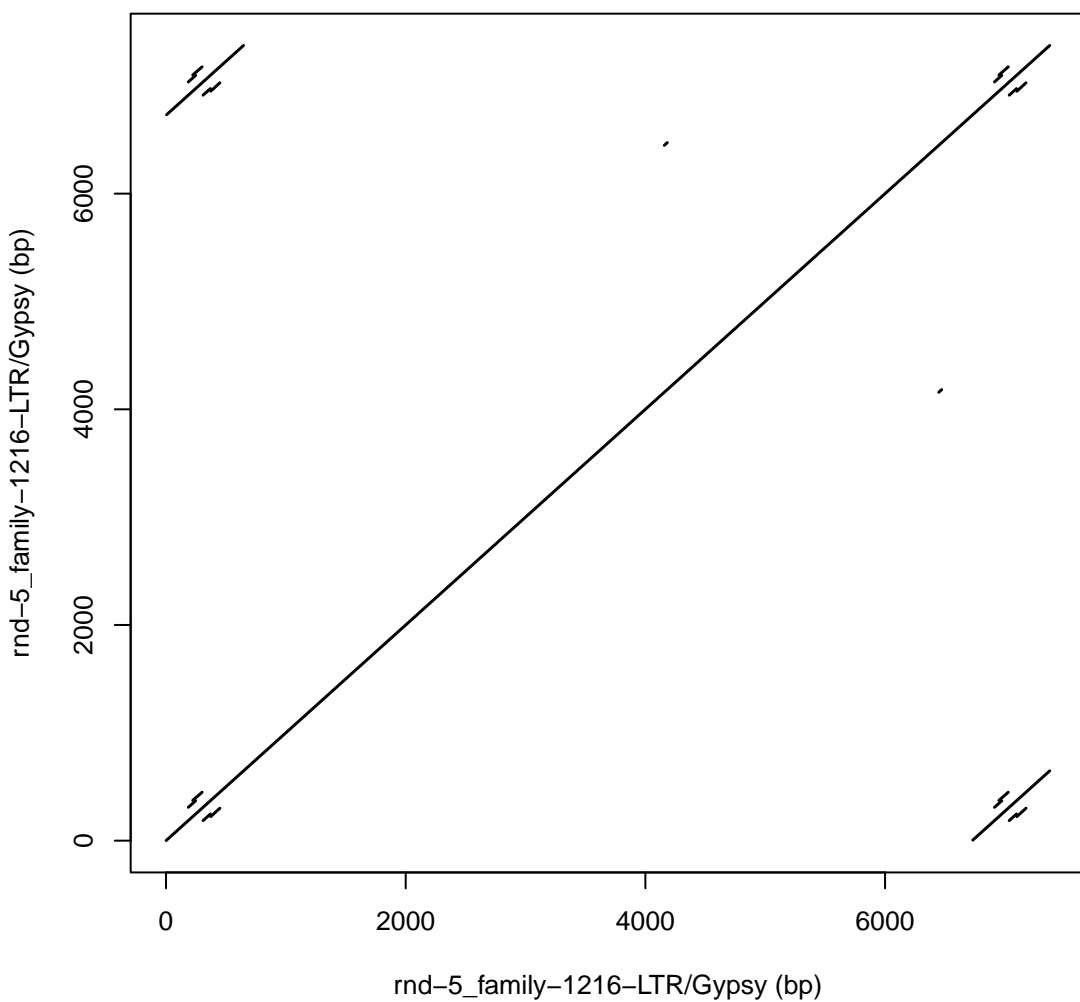
TE: rnd-5_family-1216-LTR/Gypsy
consensus size: 7374bp; fragments: 583; full length: 16 (>=6636.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

