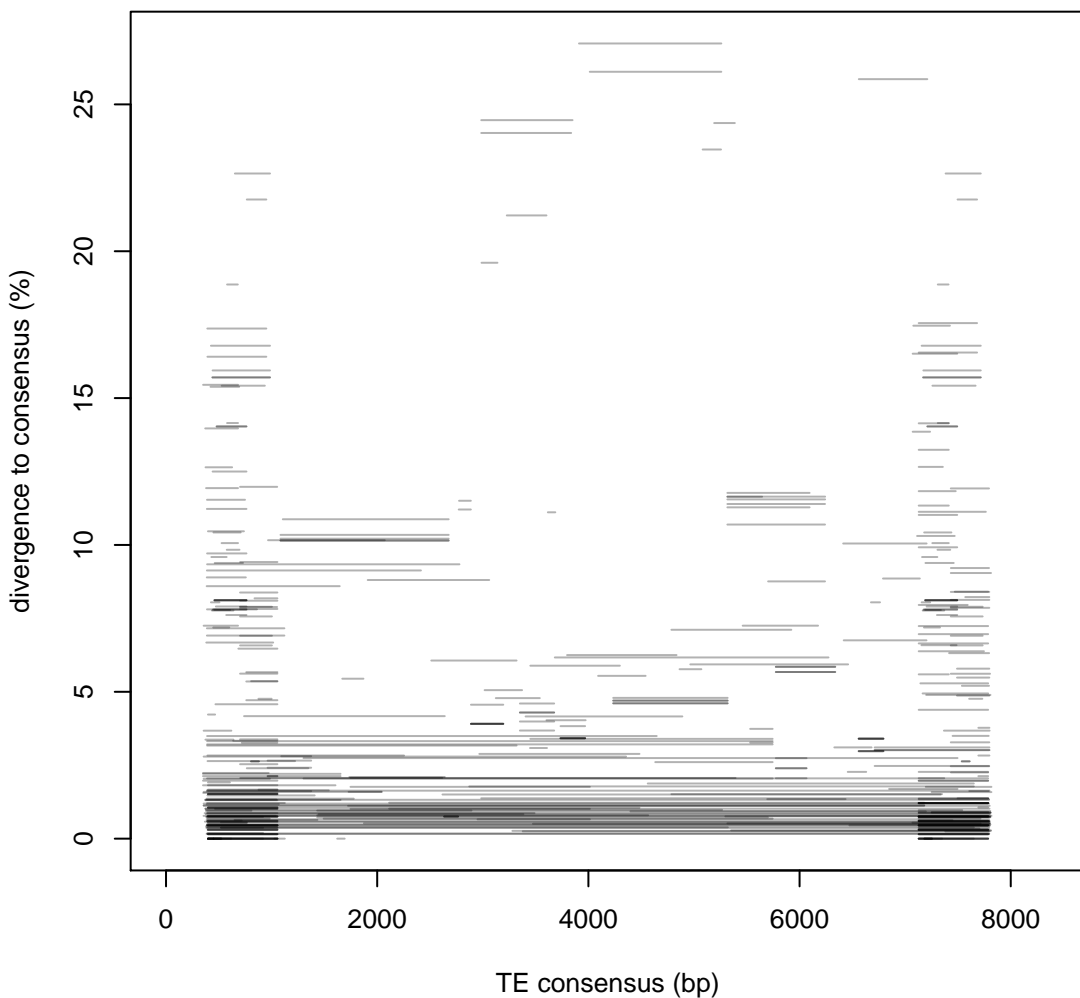
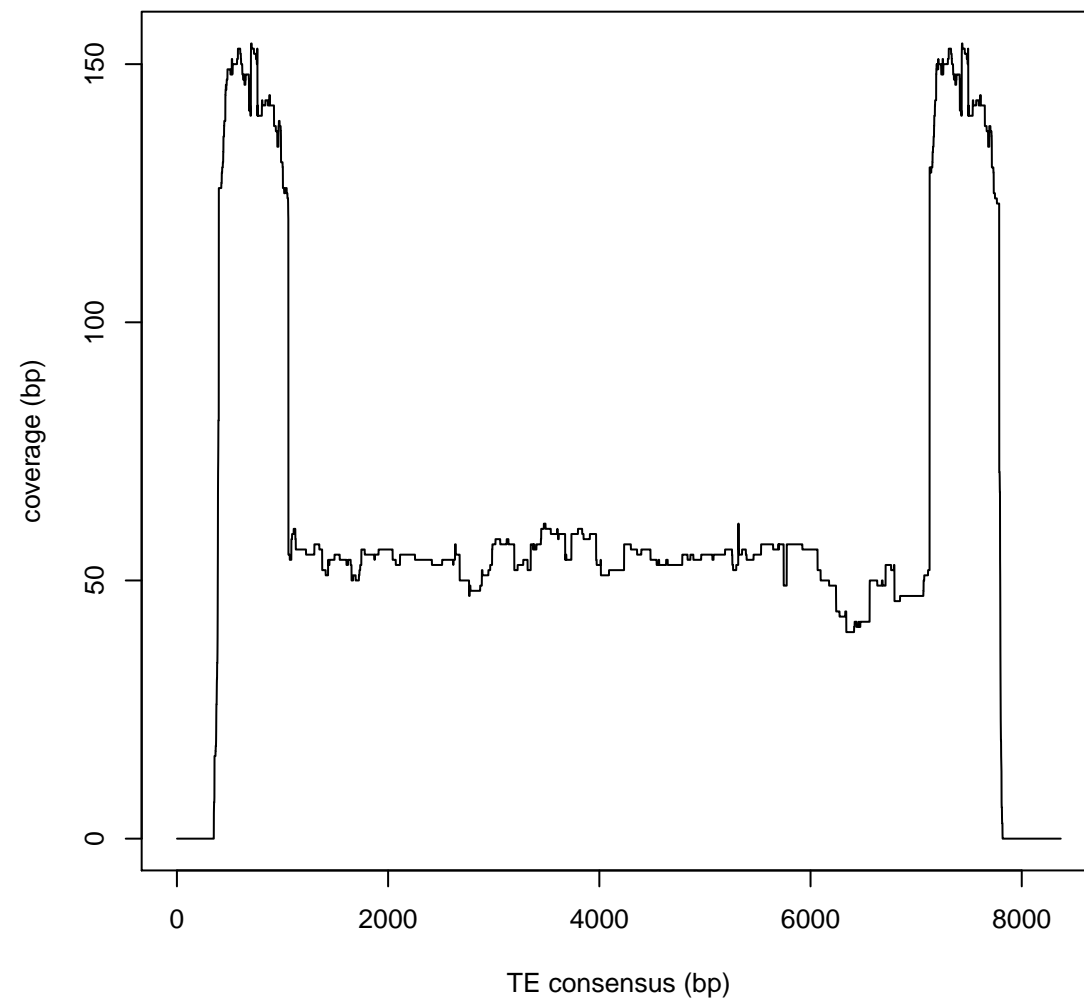


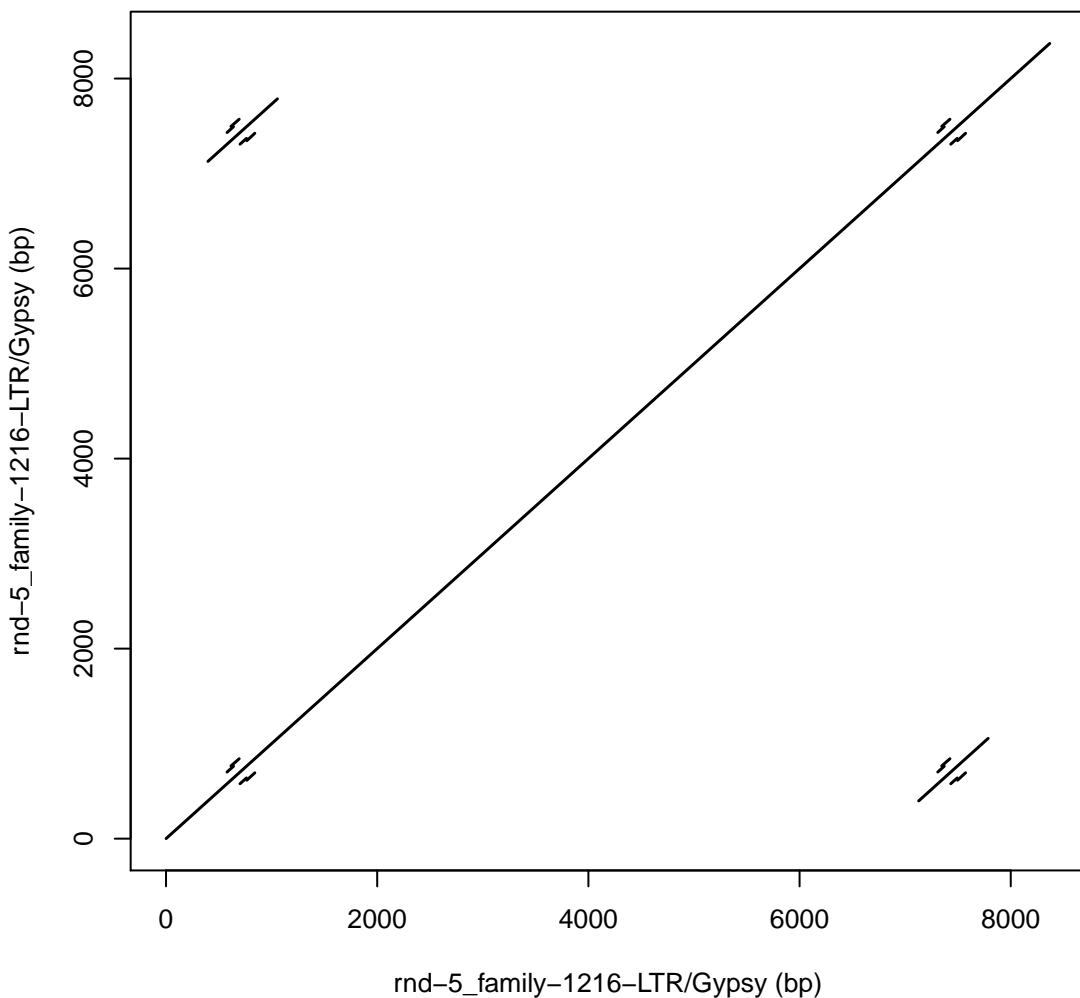
TE: rnd-5_family-1216-LTR/Gypsy
consensus size: 8369bp; fragments: 482; full length: 0 (>=7532.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

