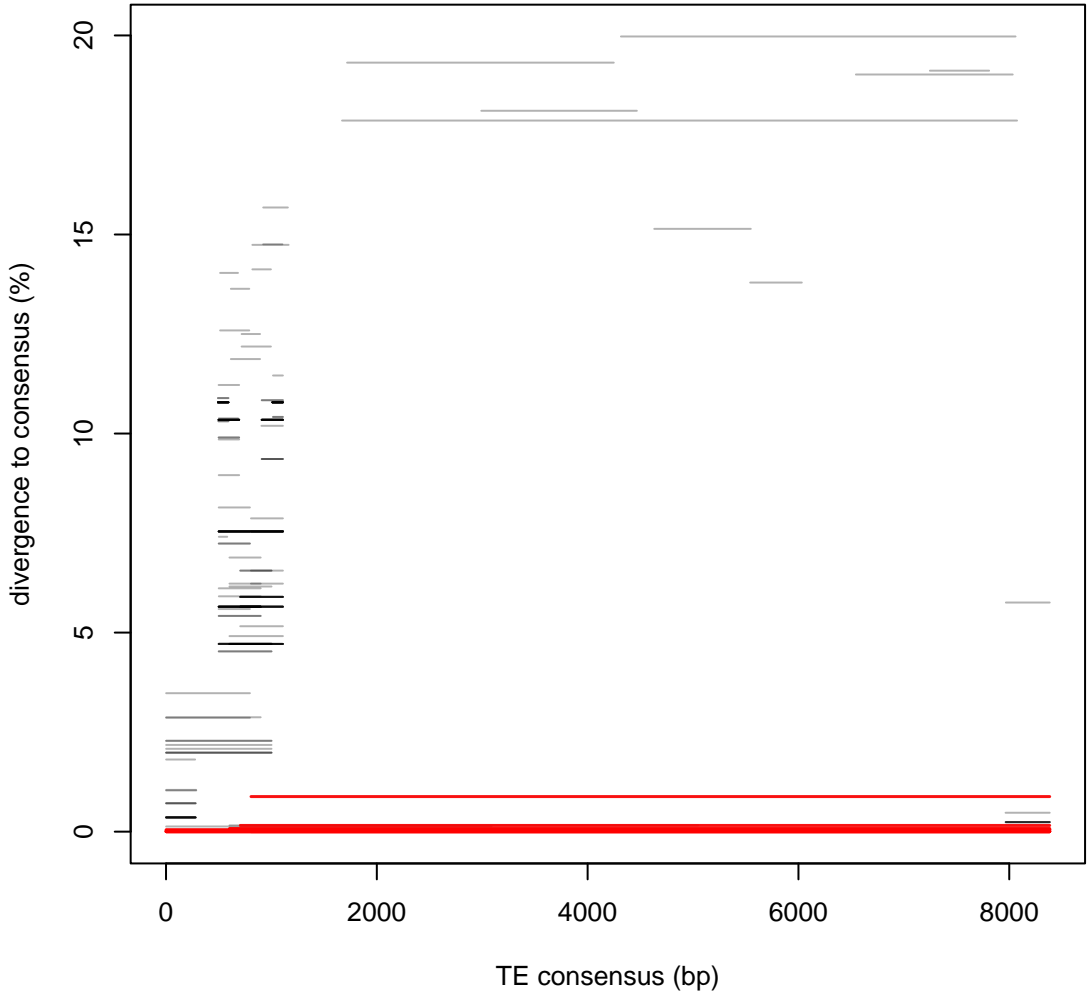
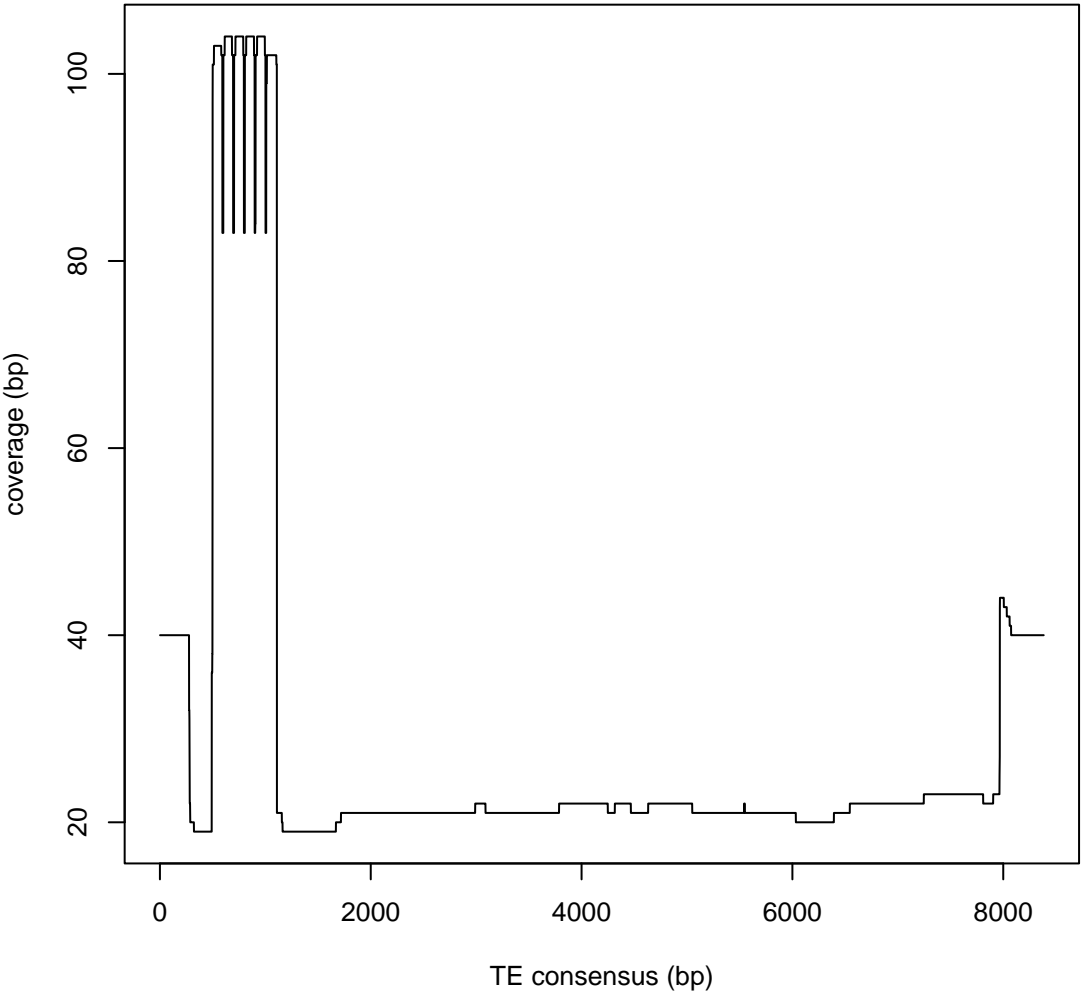


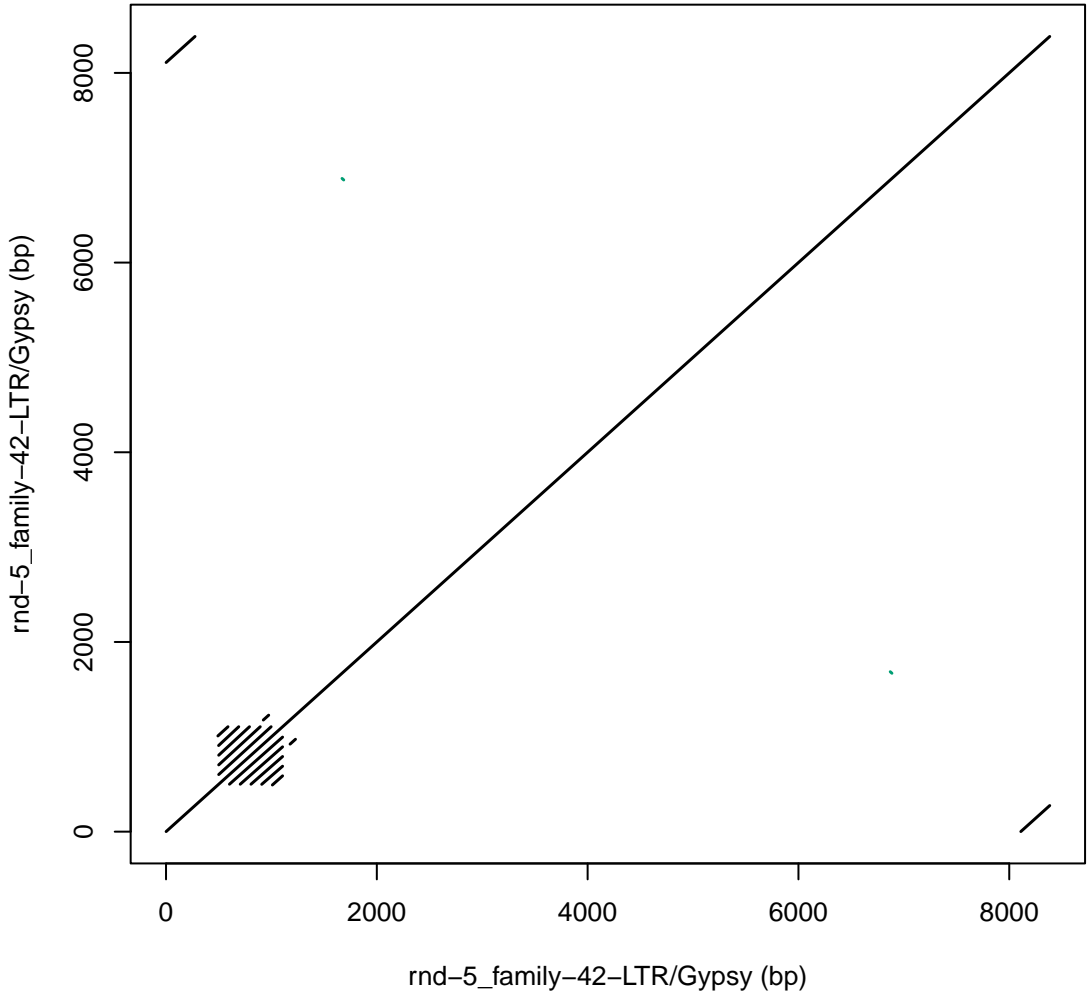
TE: rnd-5\_family-42-LTR/Gypsy  
consensus size: 8384bp; fragments: 259; full length: 17 (>=7545.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

