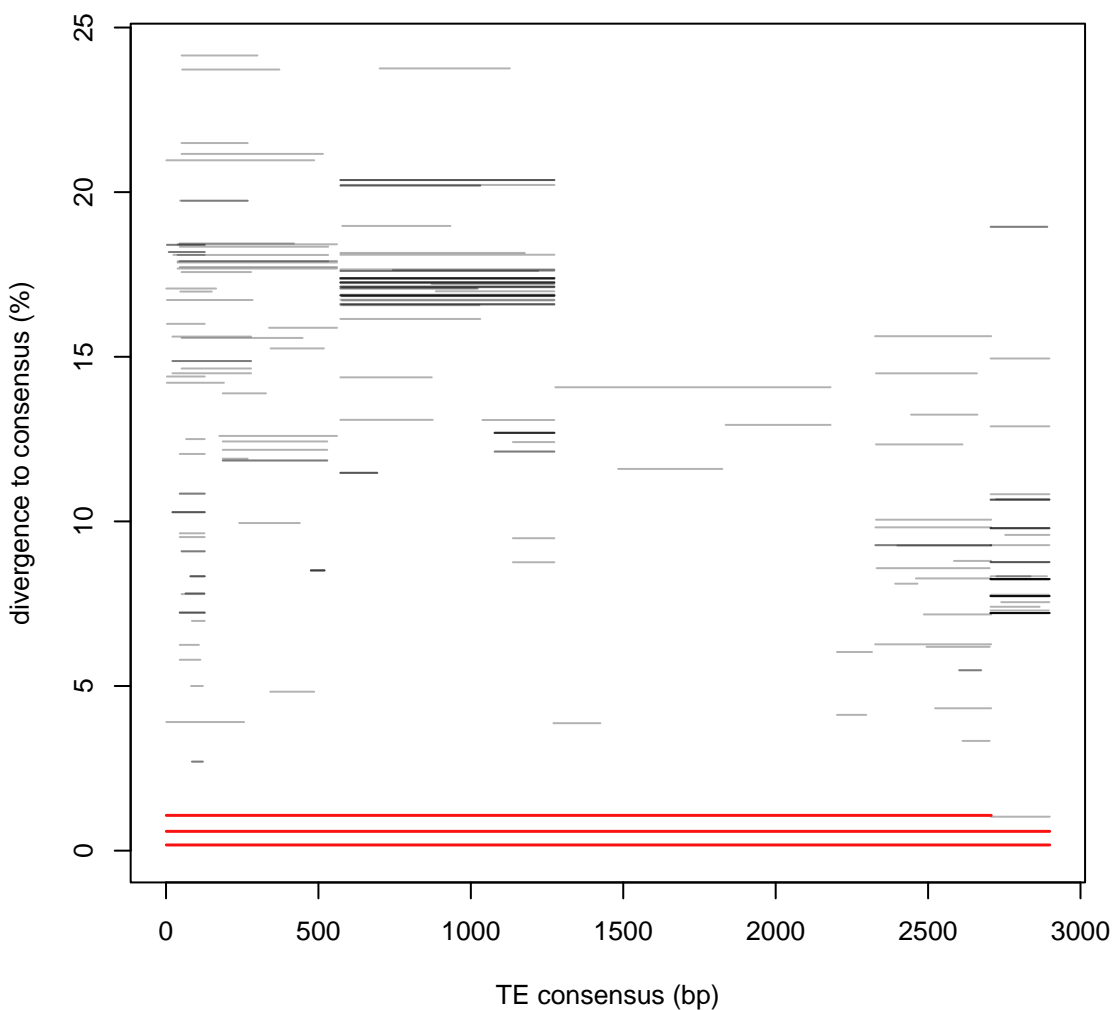
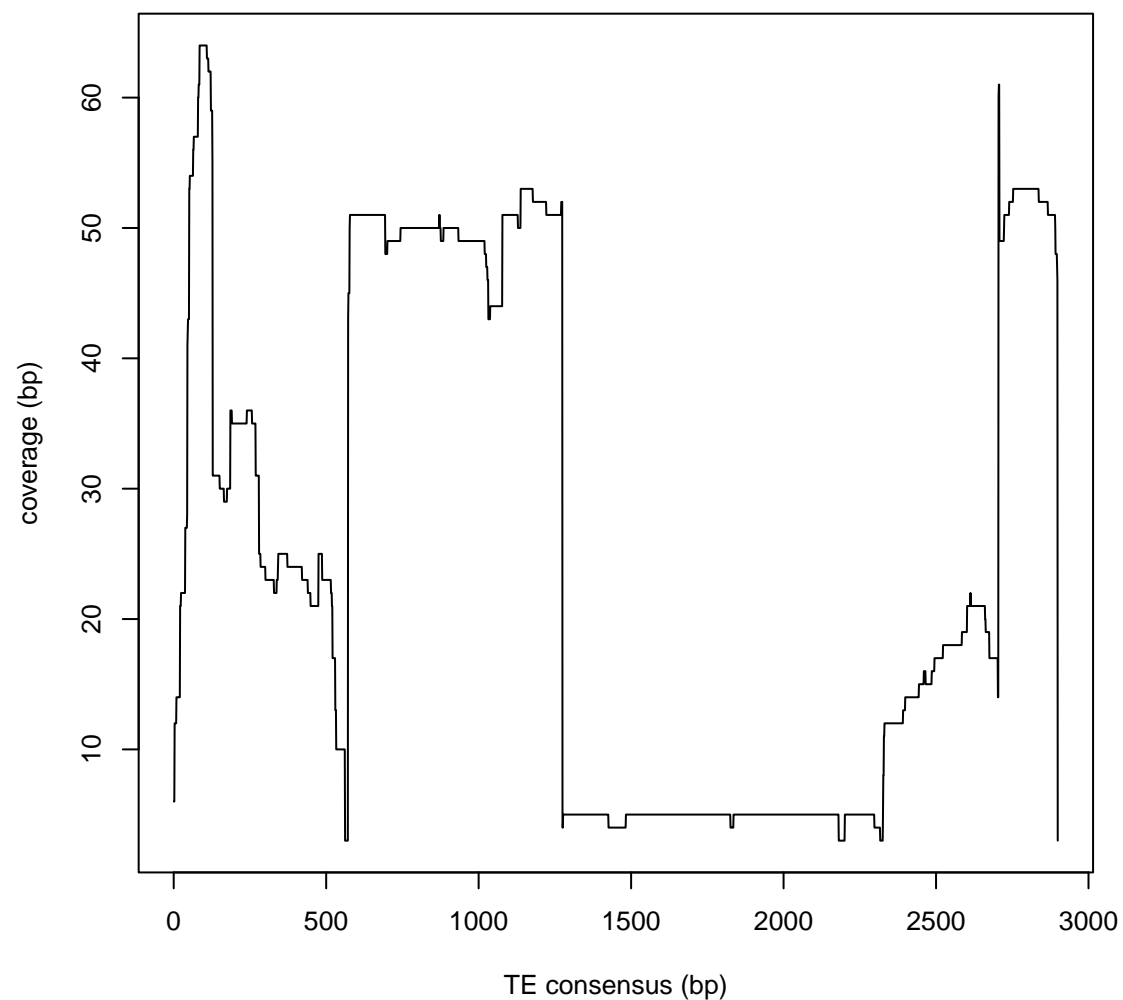


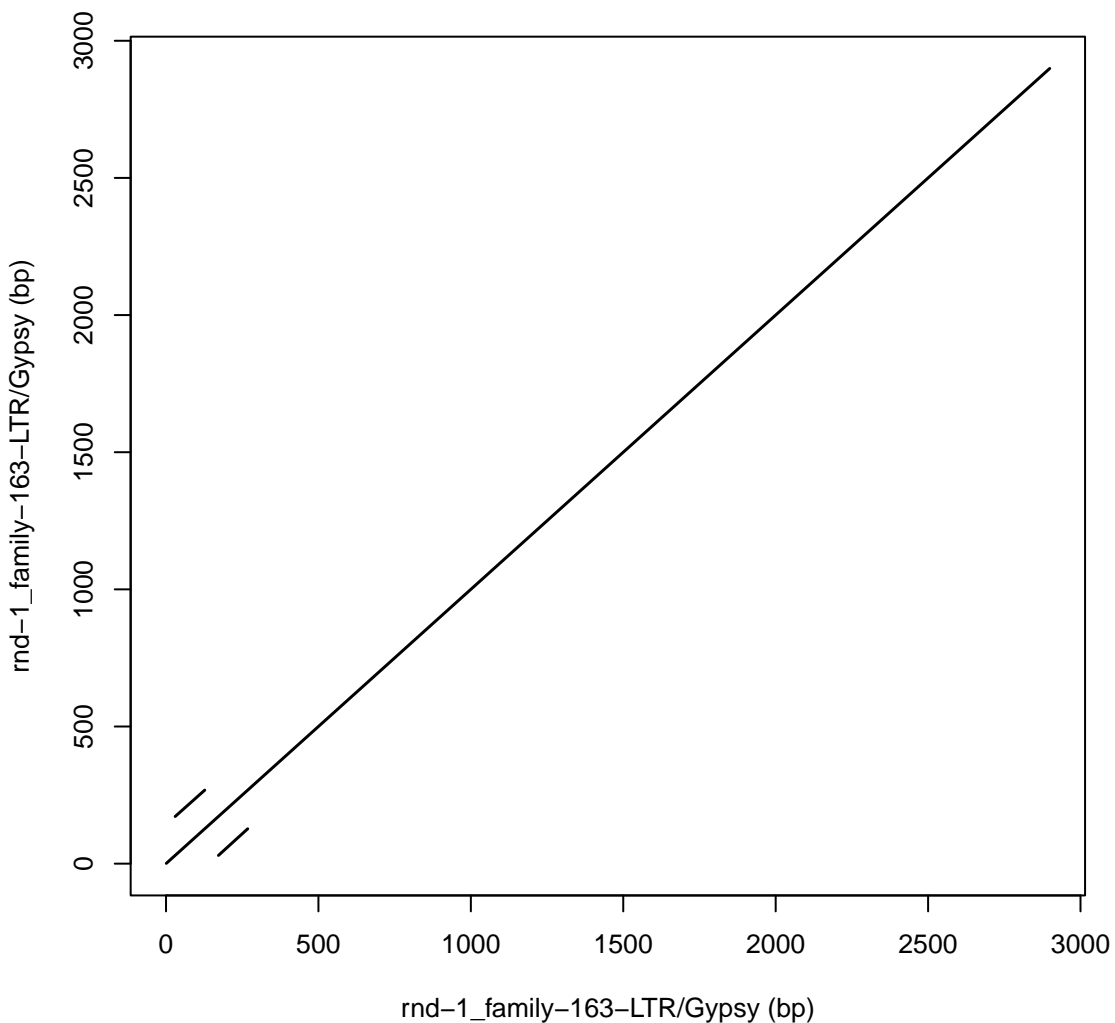
TE: rnd-1\_family-163-LTR/Gypsy  
consensus size: 2899bp; fragments: 219; full length: 3 (>=2609.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

