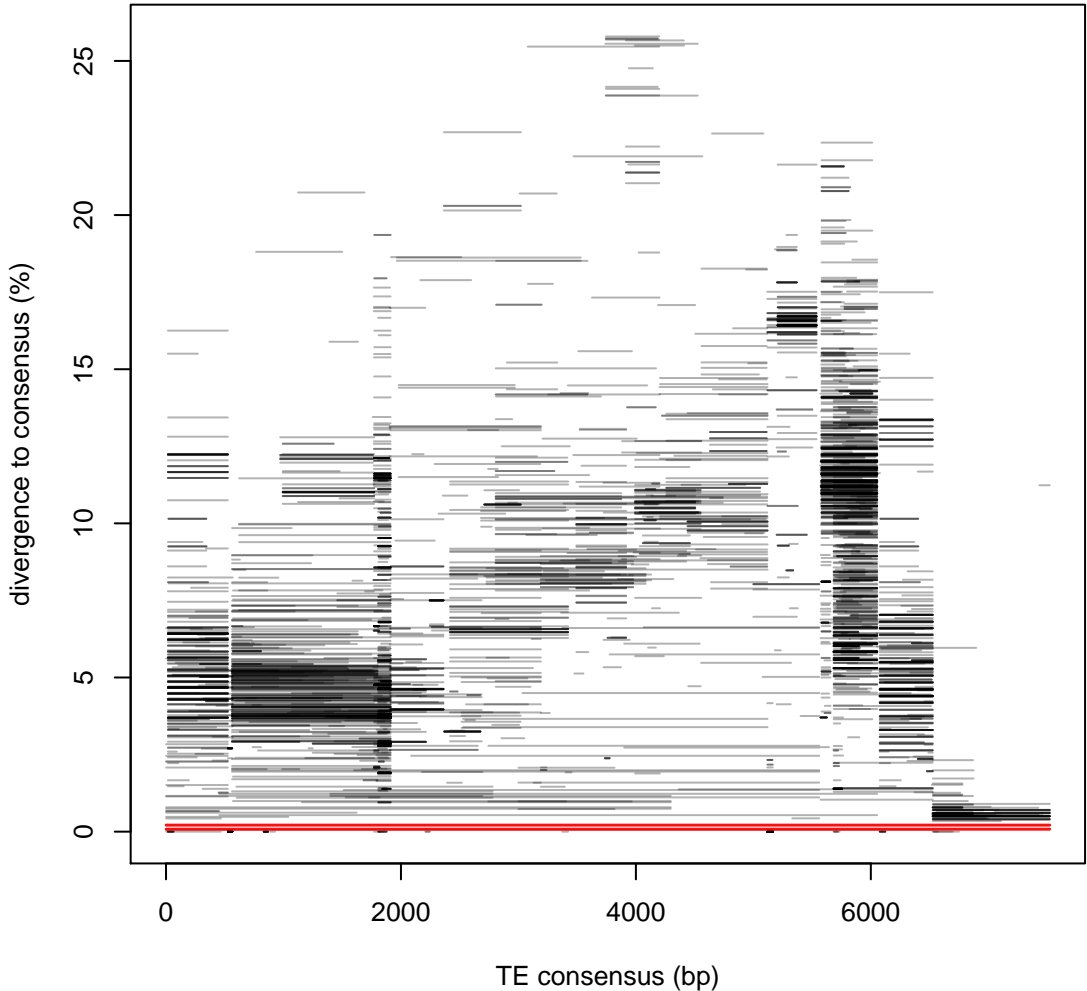
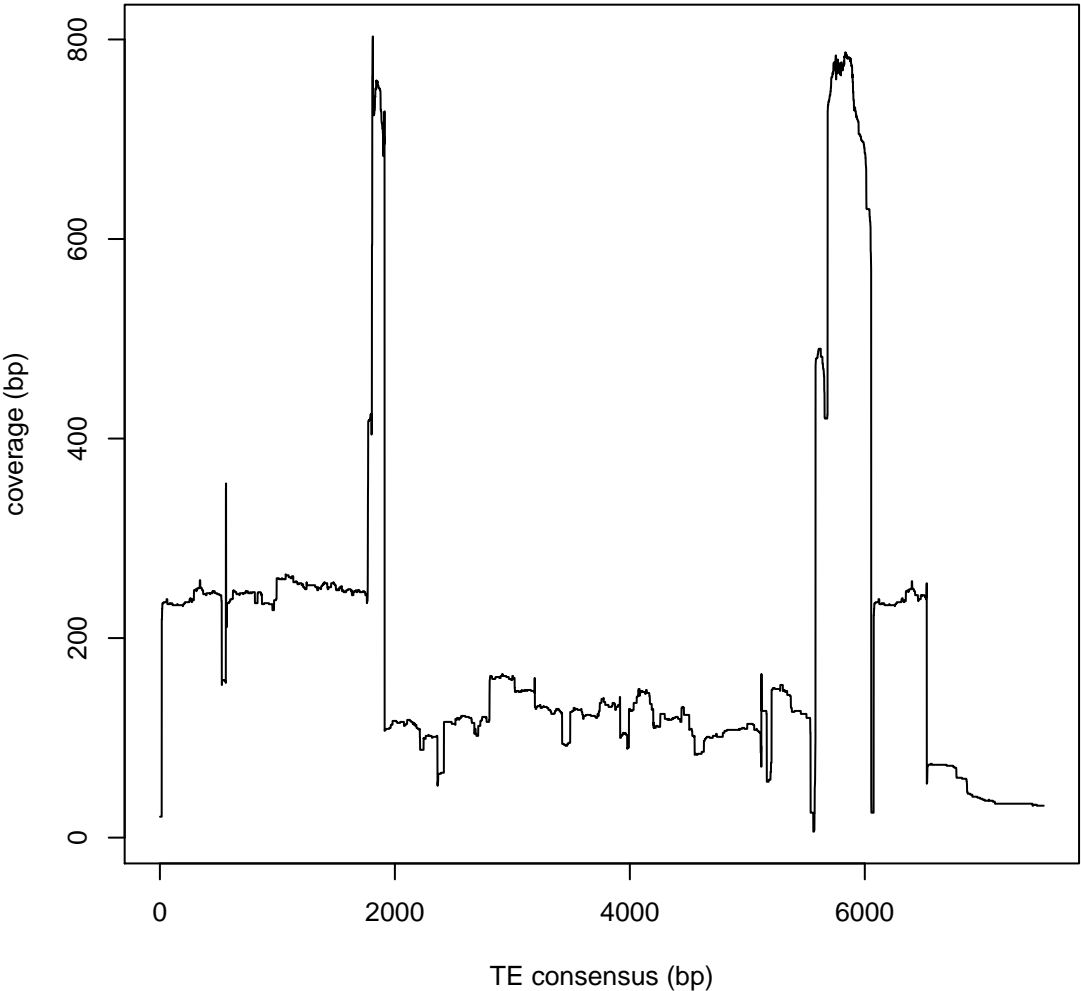


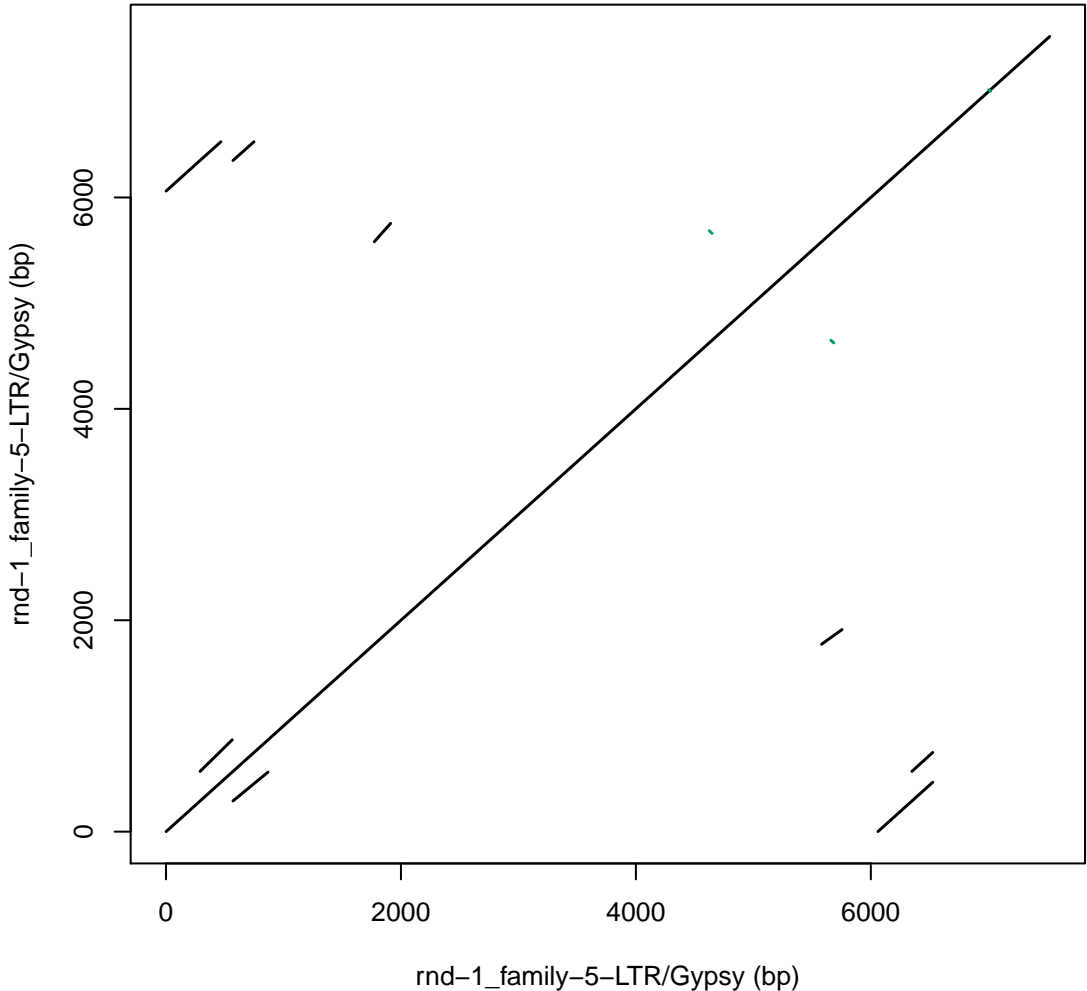
TE: rnd-1\_family-5-LTR/Gypsy  
consensus size: 7523bp; fragments: 3515; full length: 2 (>=6770.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

