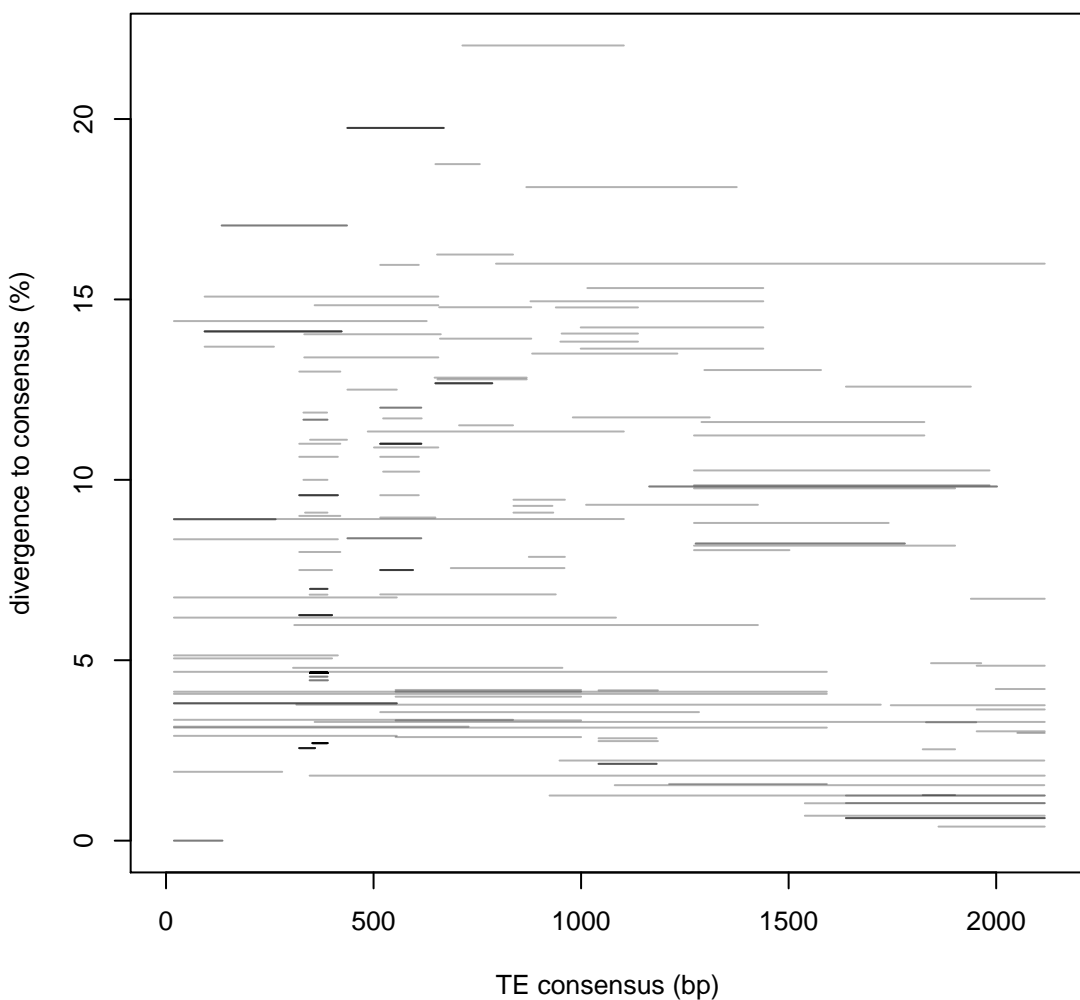
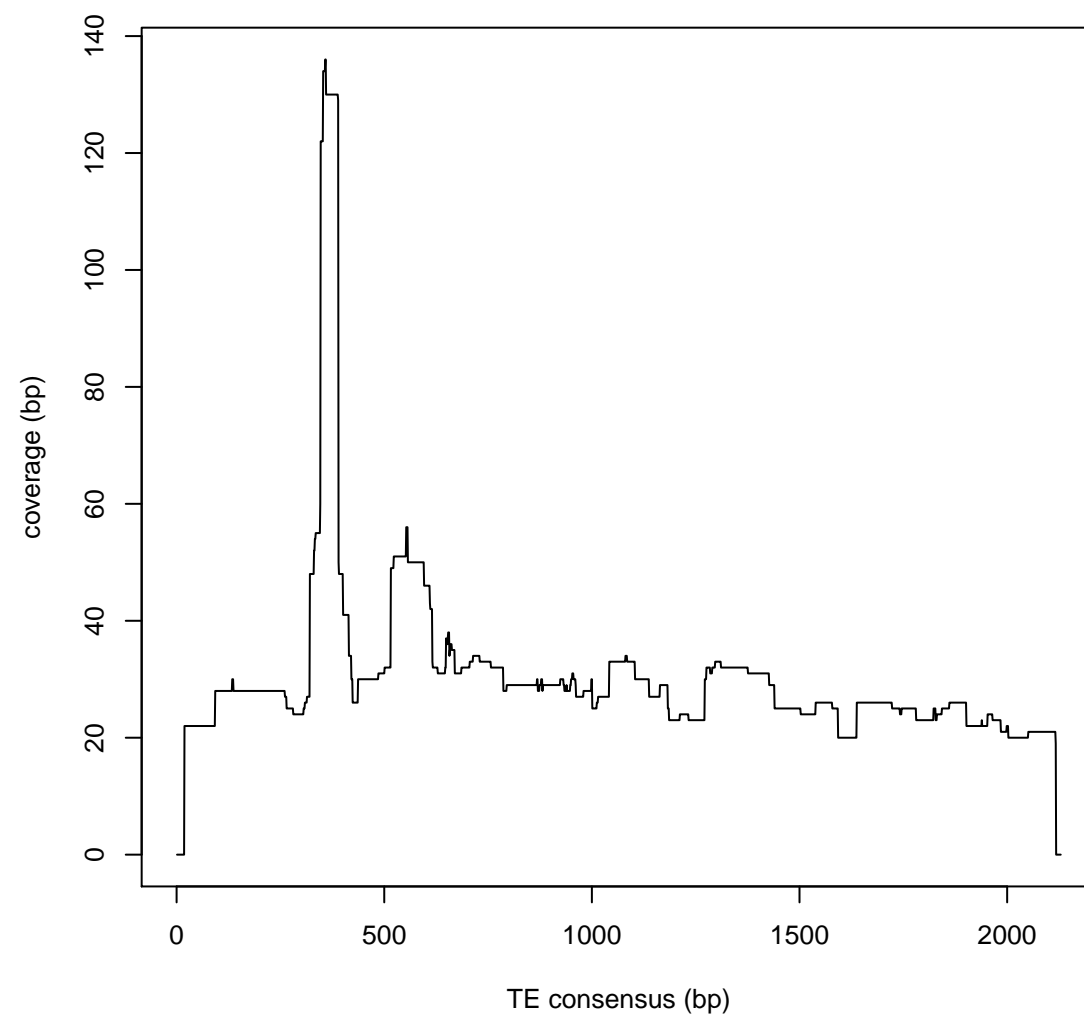


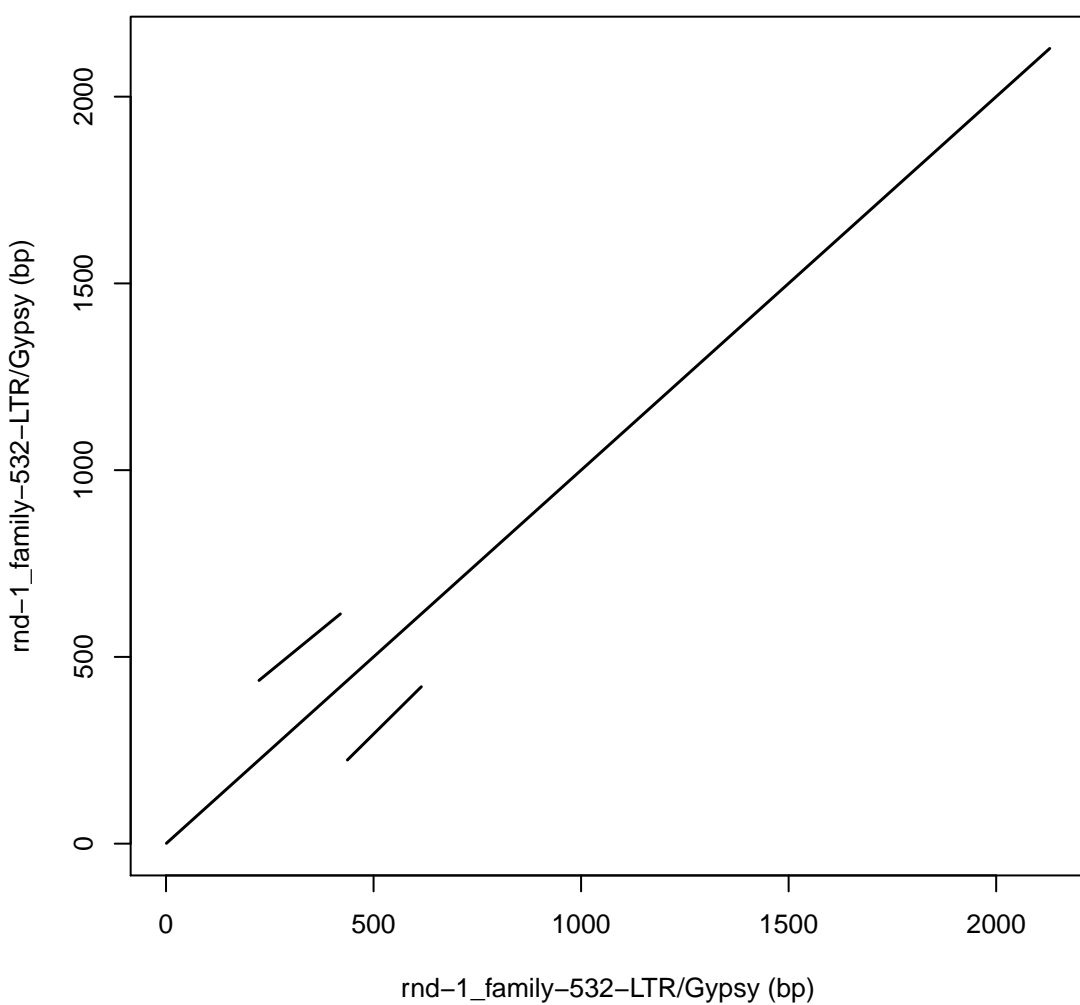
TE: rnd-1\_family-532-LTR/Gypsy  
consensus size: 2129bp; fragments: 247; full length: 0 (>=1916.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

