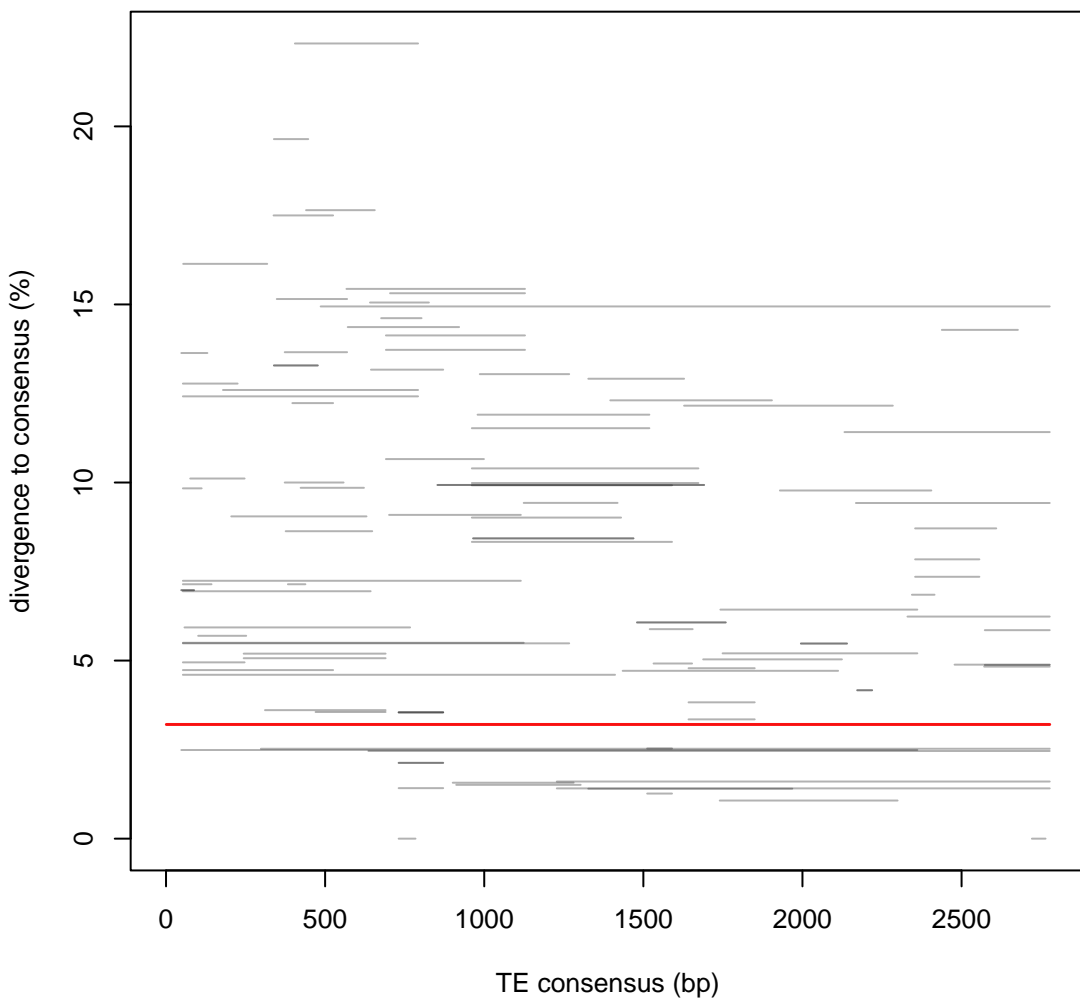
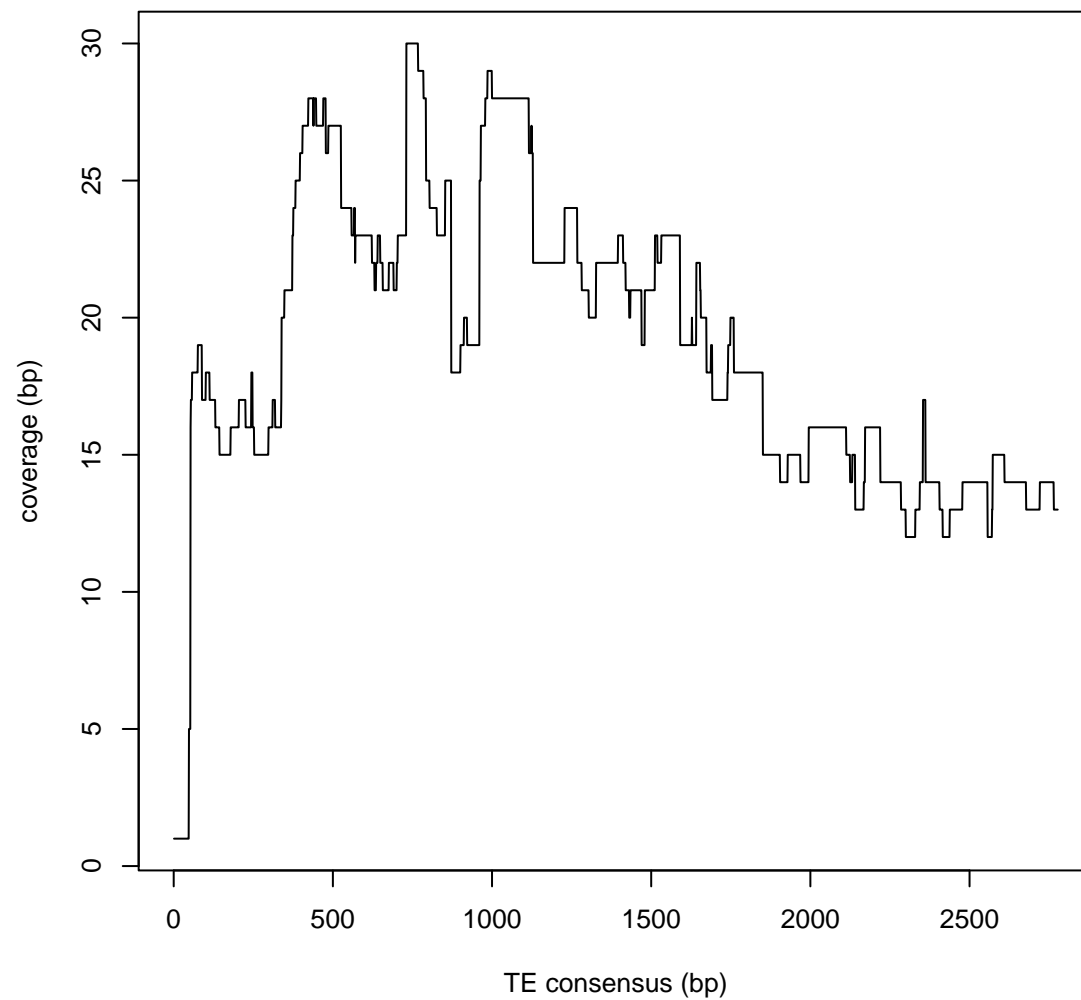


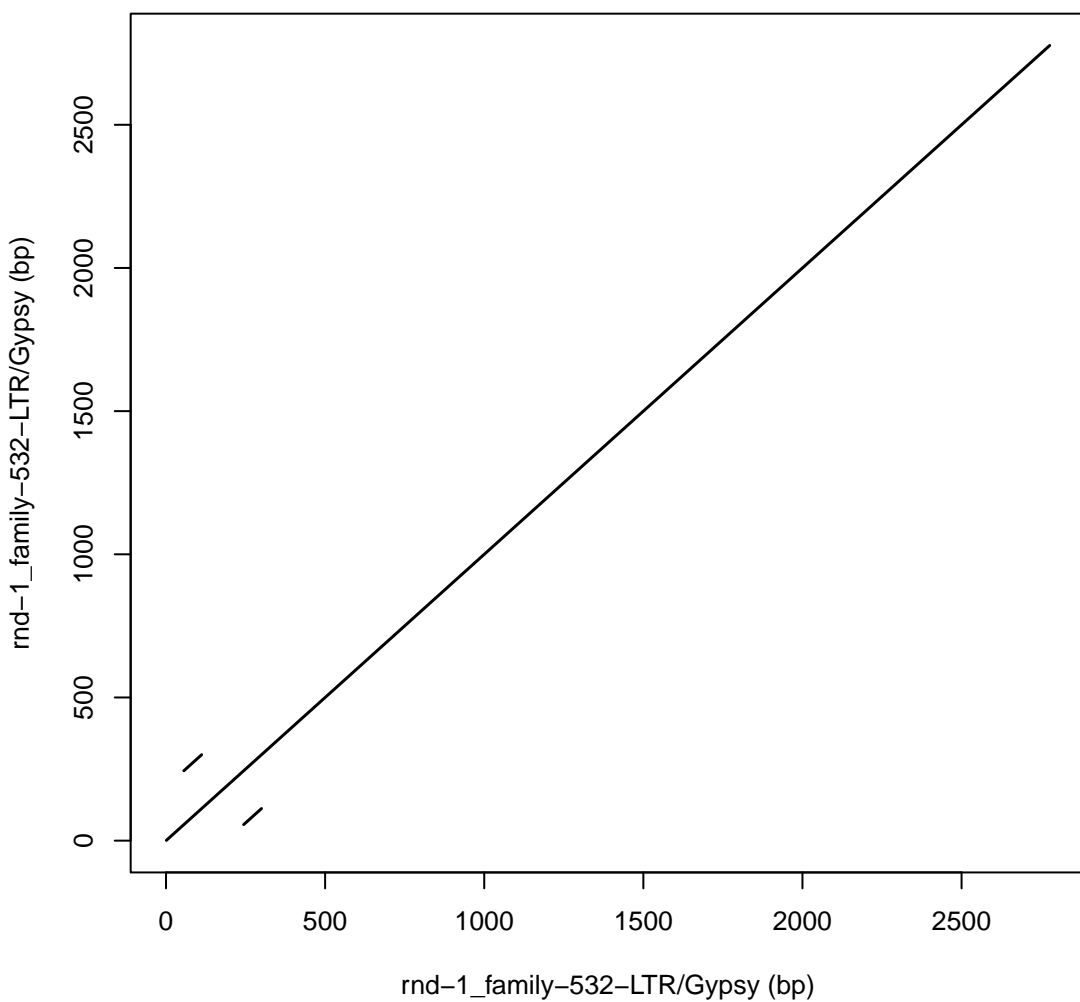
TE: rnd-1\_family-532-LTR/Gypsy  
consensus size: 2777bp; fragments: 112; full length: 1 (>=2499.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

