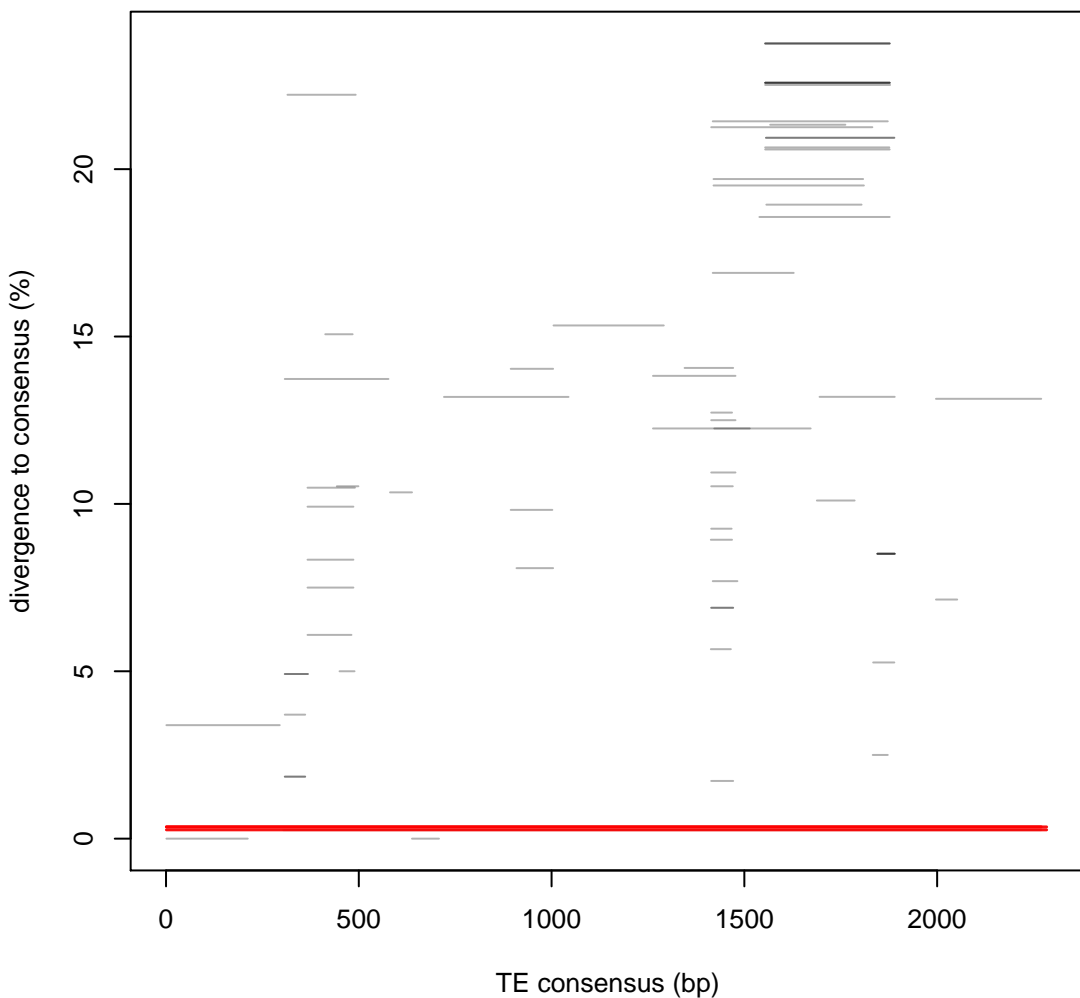
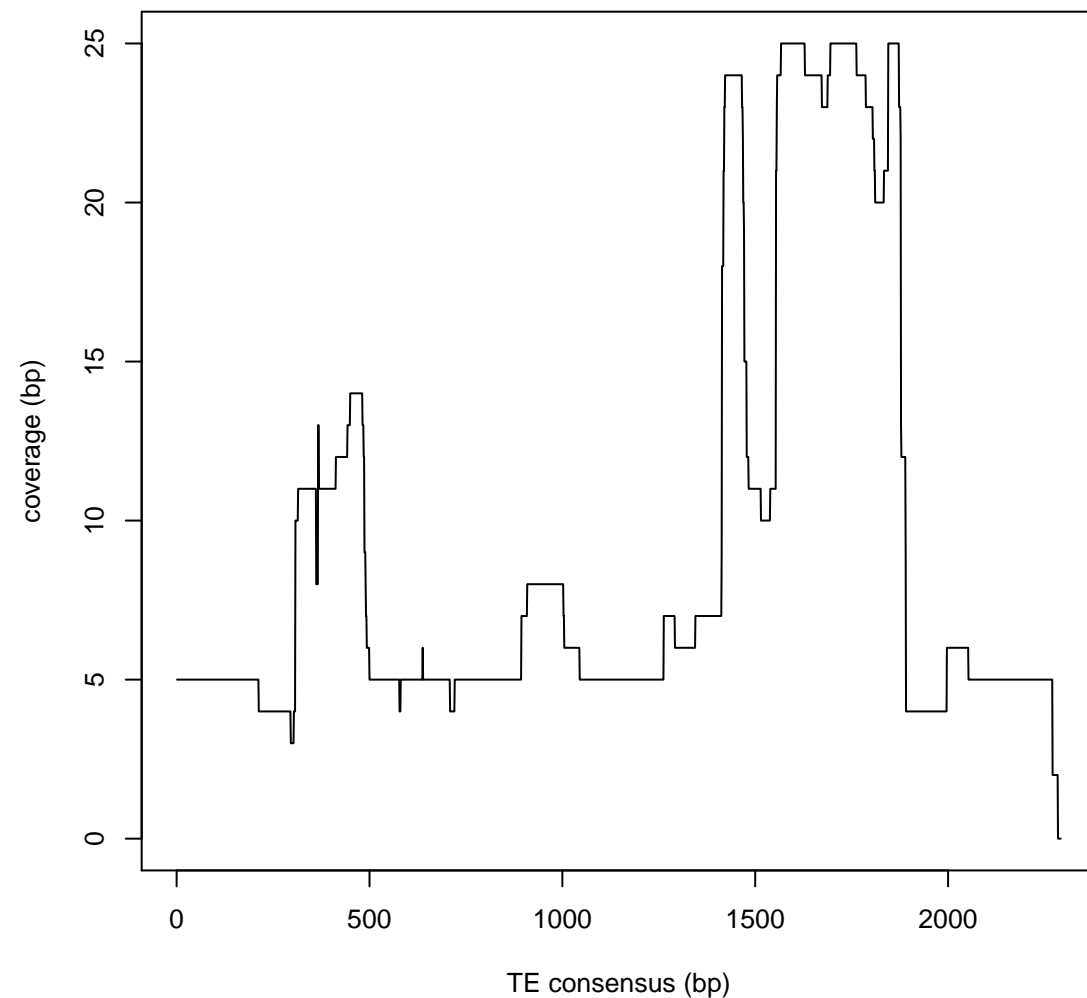


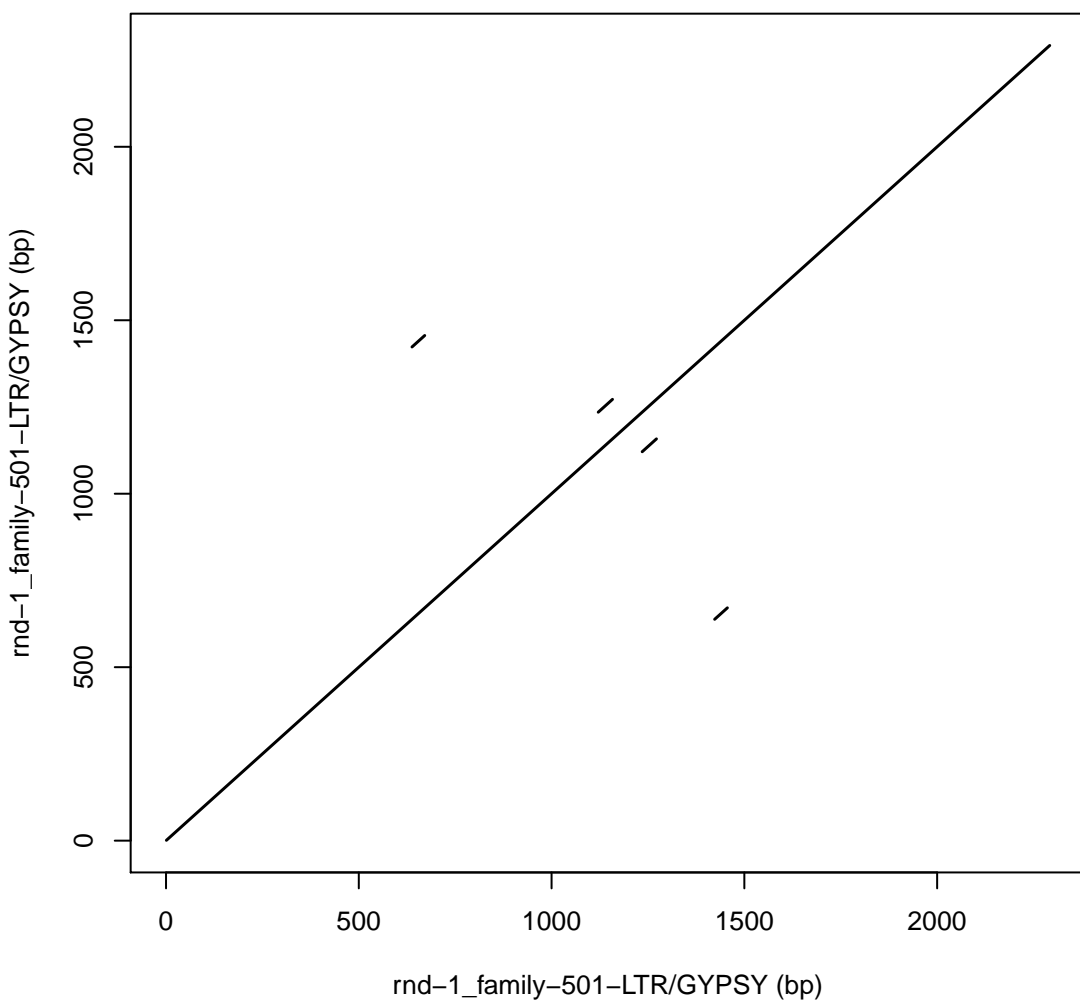
TE: rnd-1\_family-501-LTR/GYPSY  
consensus size: 2292bp; fragments: 73; full length: 3 (>=2062.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

