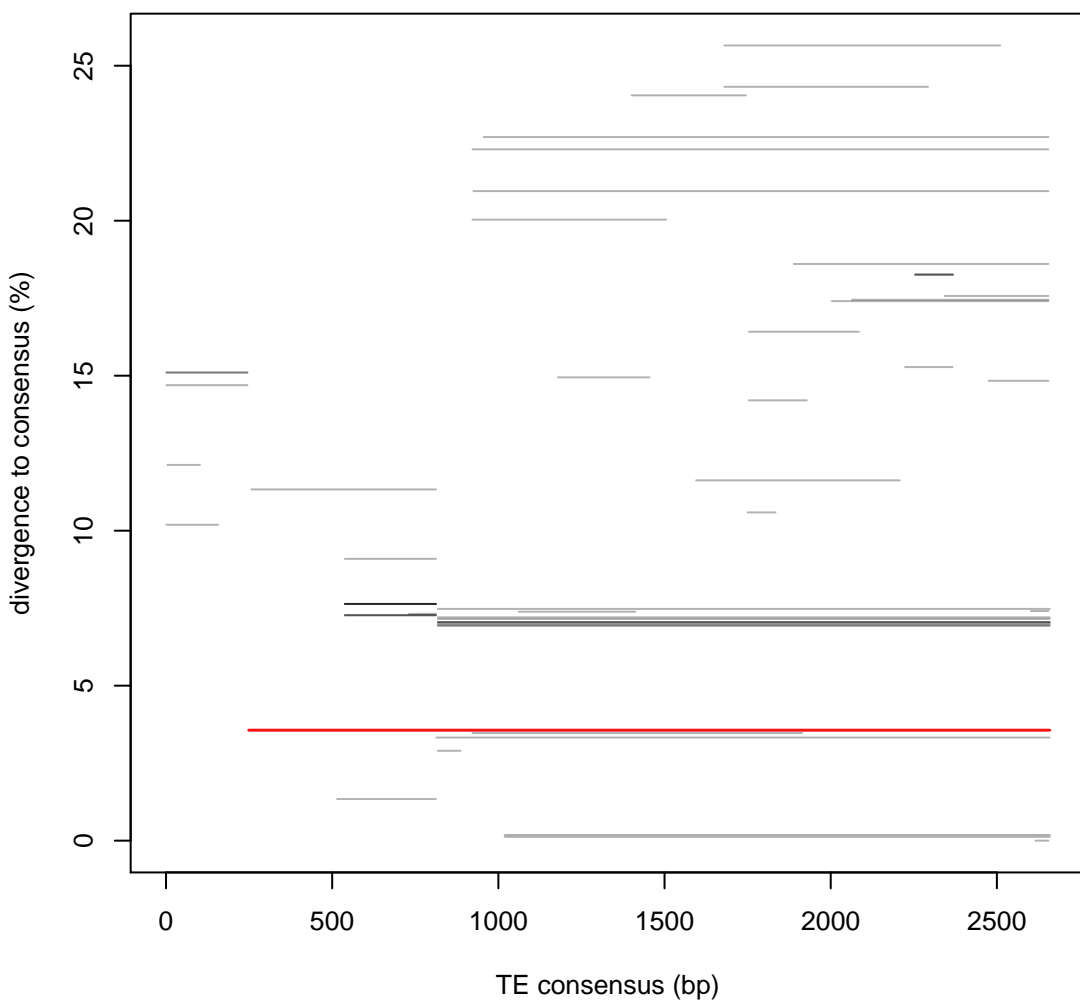
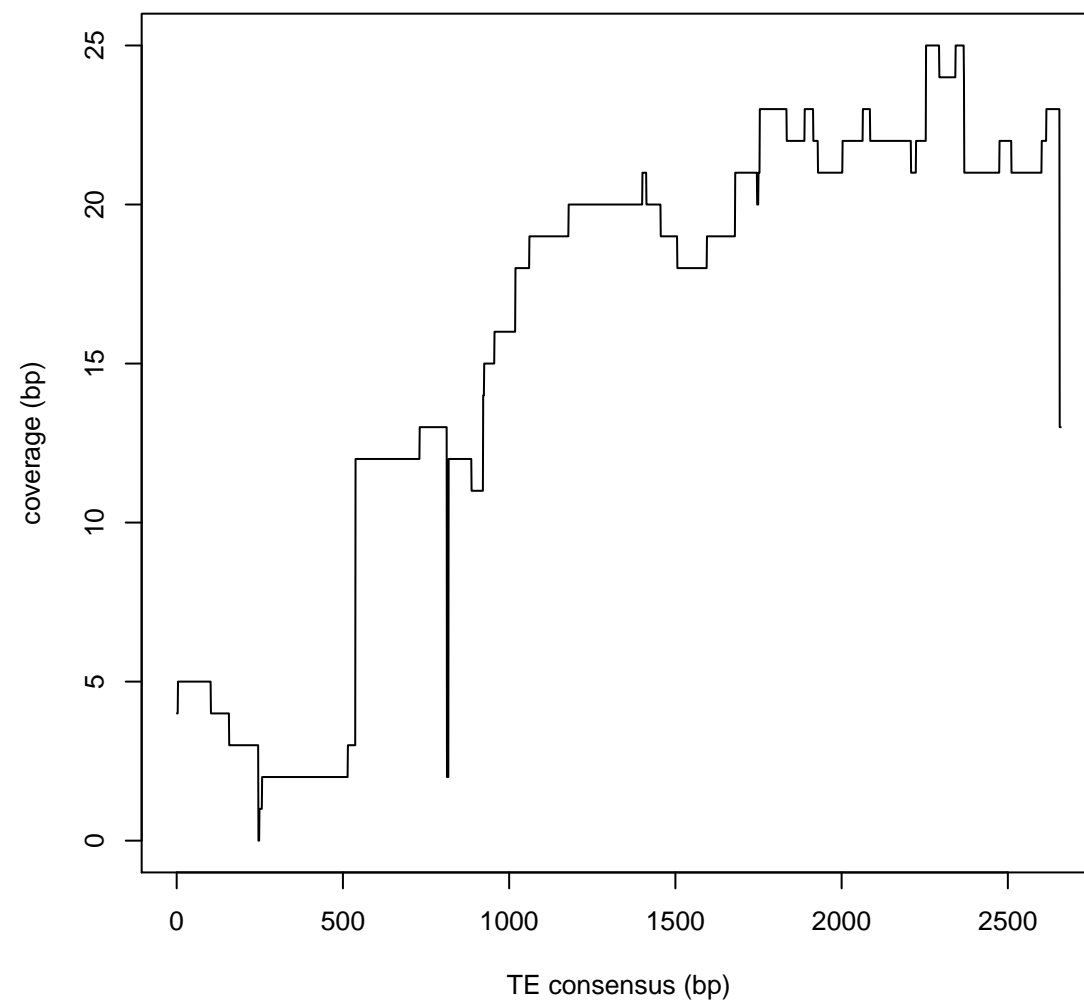


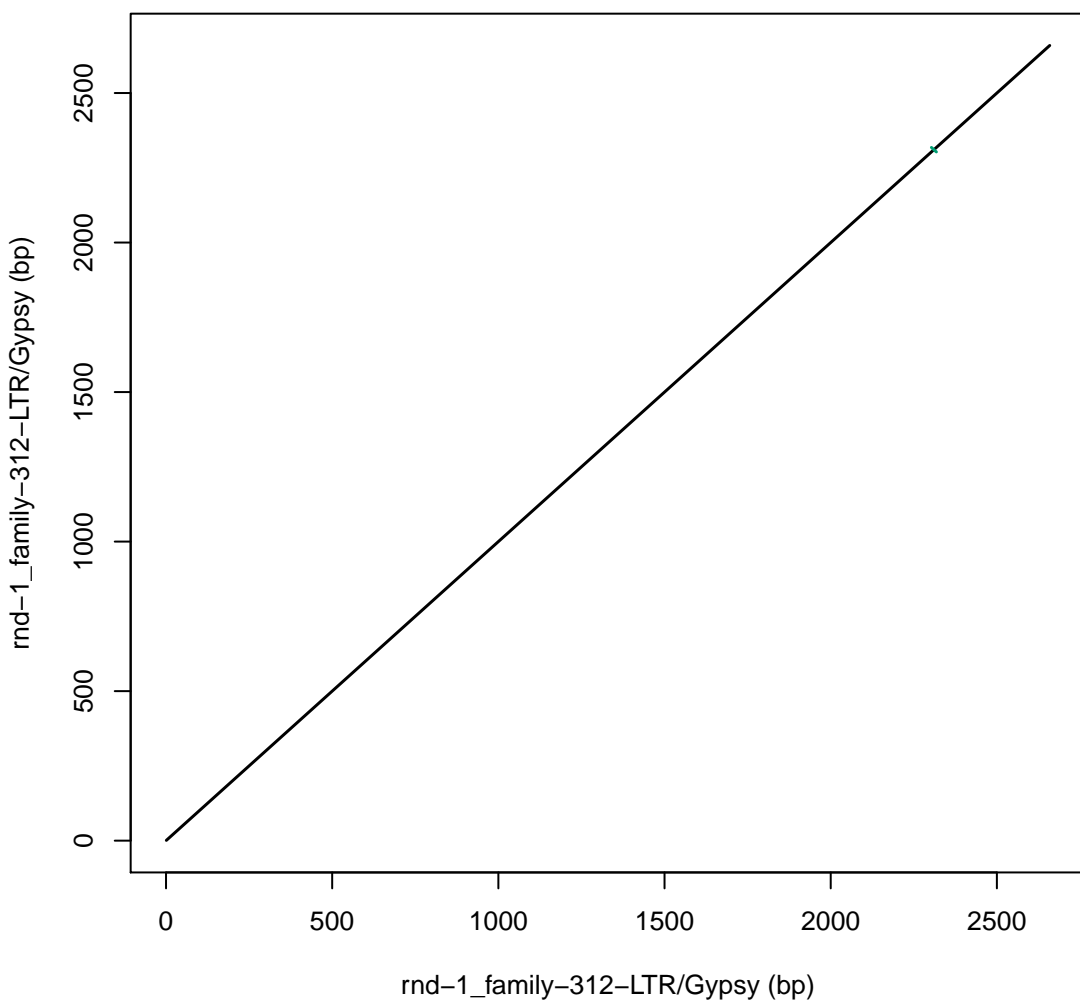
TE: rnd-1\_family-312-LTR/Gypsy  
consensus size: 2659bp; fragments: 56; full length: 1 (>=2393.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

