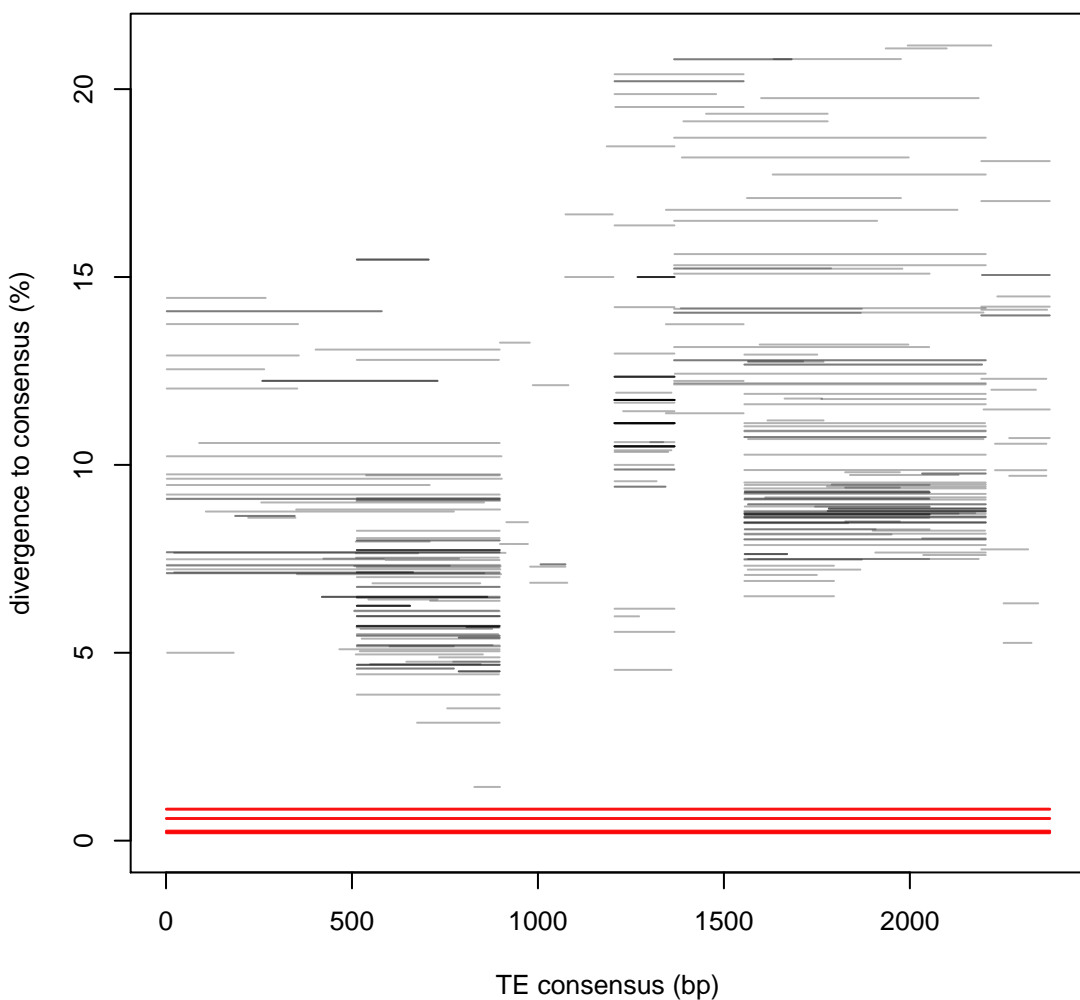
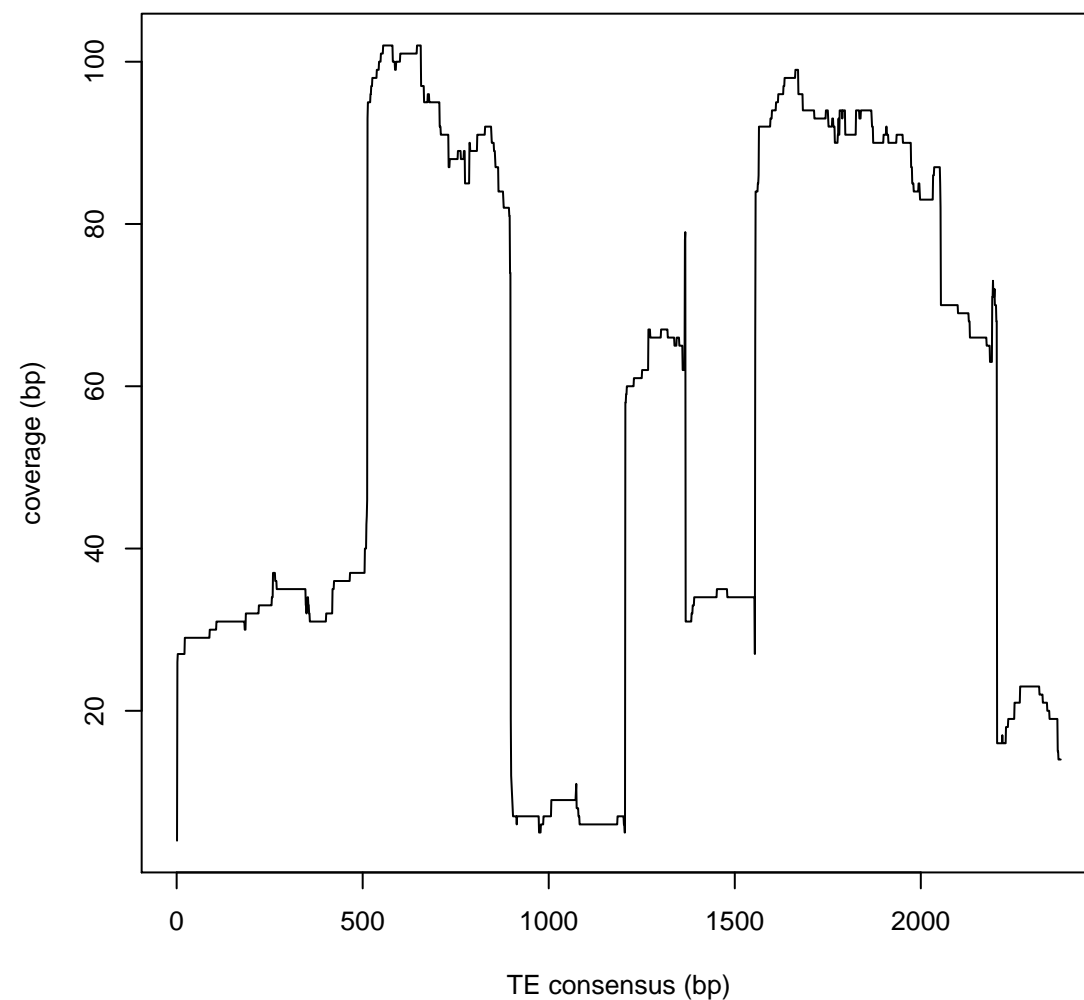


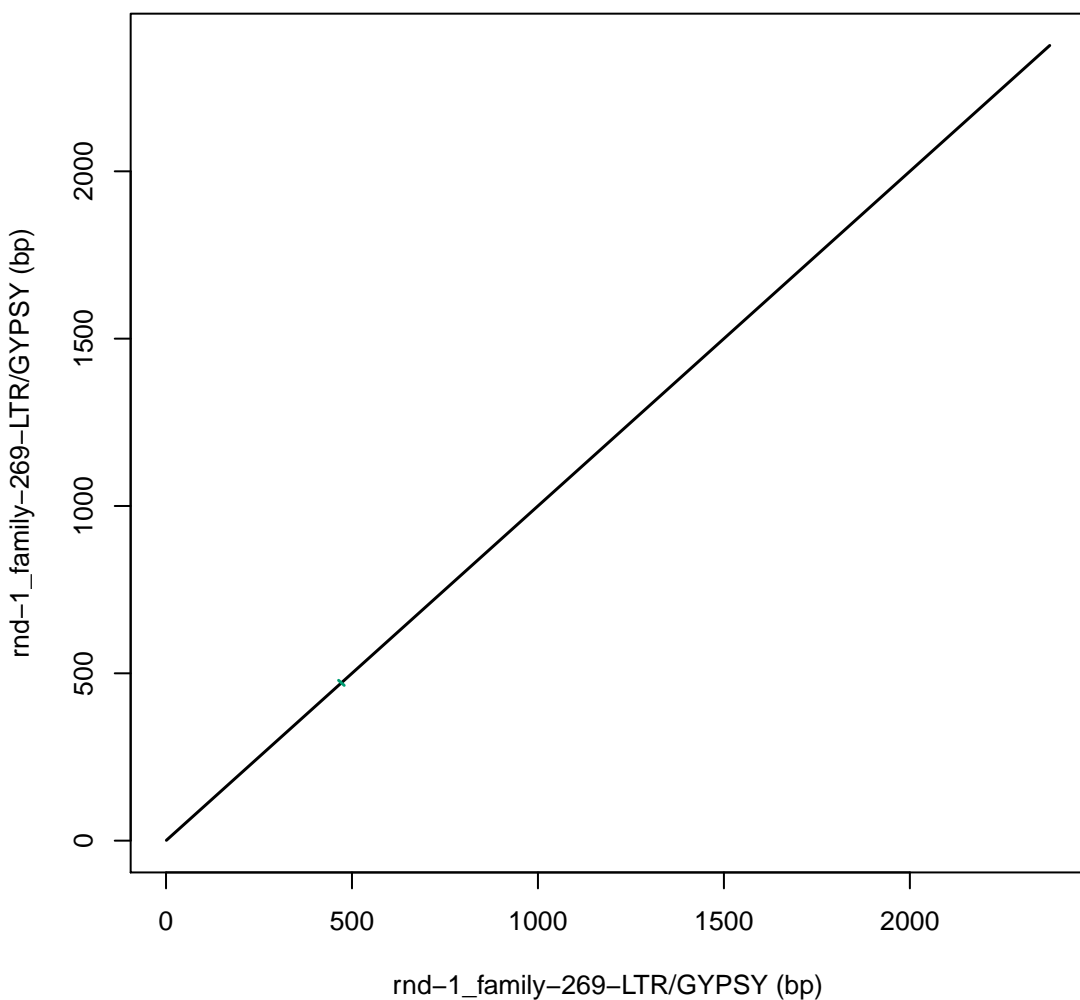
TE: rnd-1\_family-269-LTR/GYPSY  
consensus size: 2376bp; fragments: 339; full length: 4 ( $\geq 2138.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

