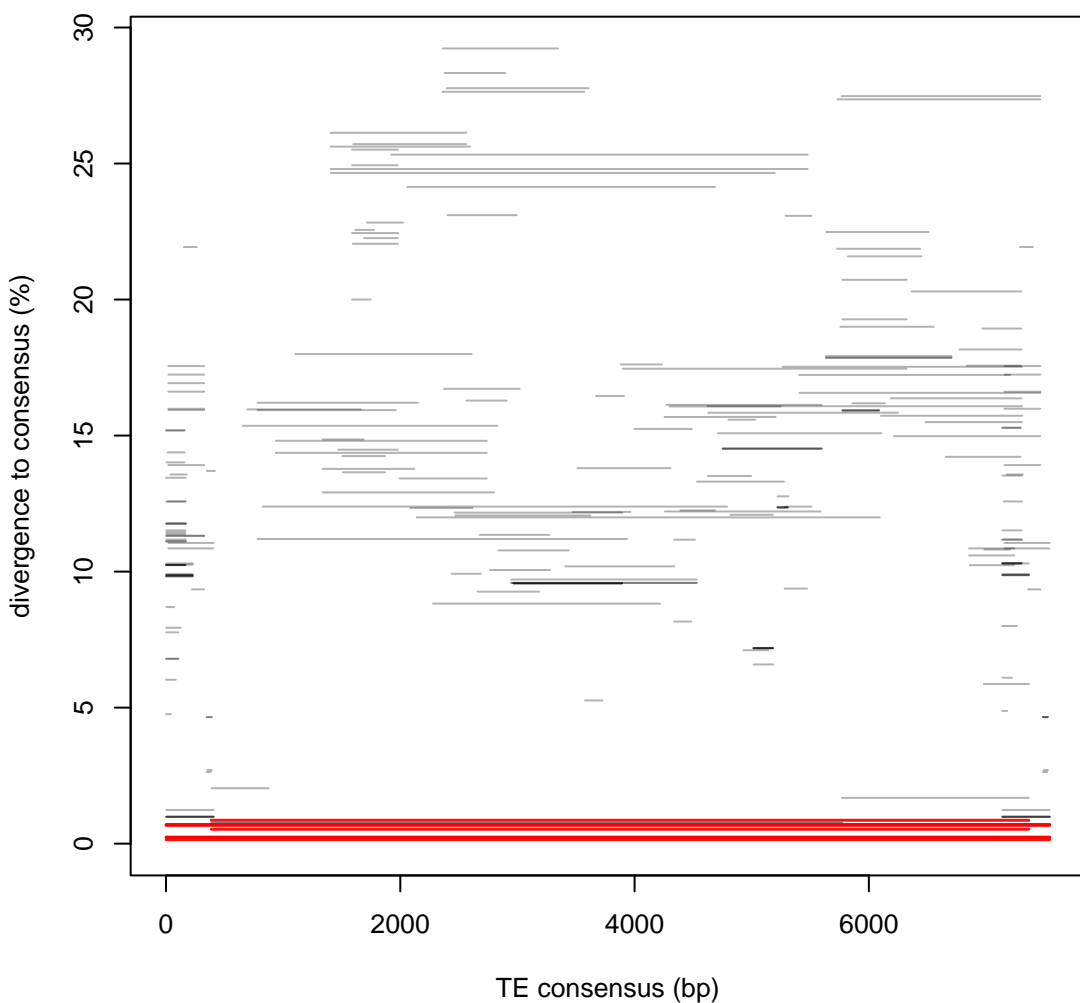
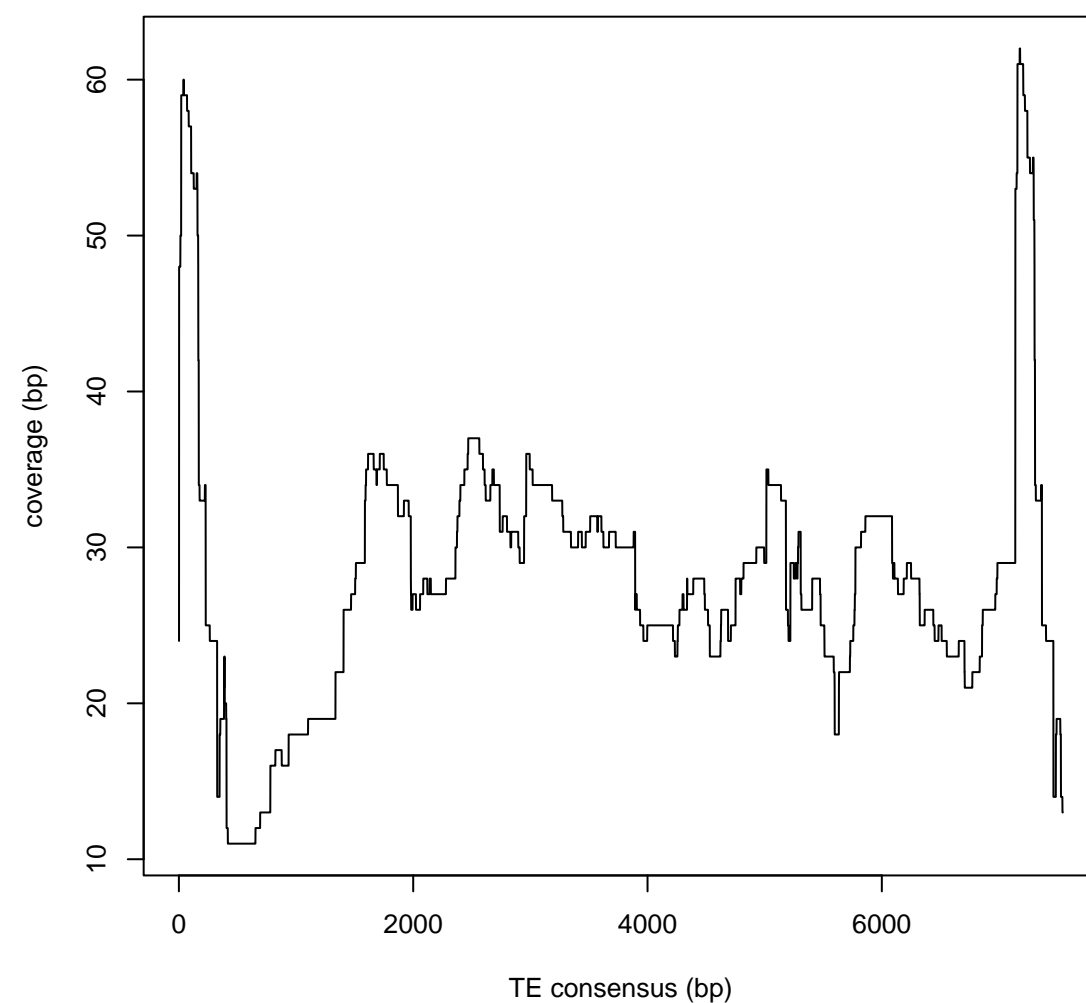


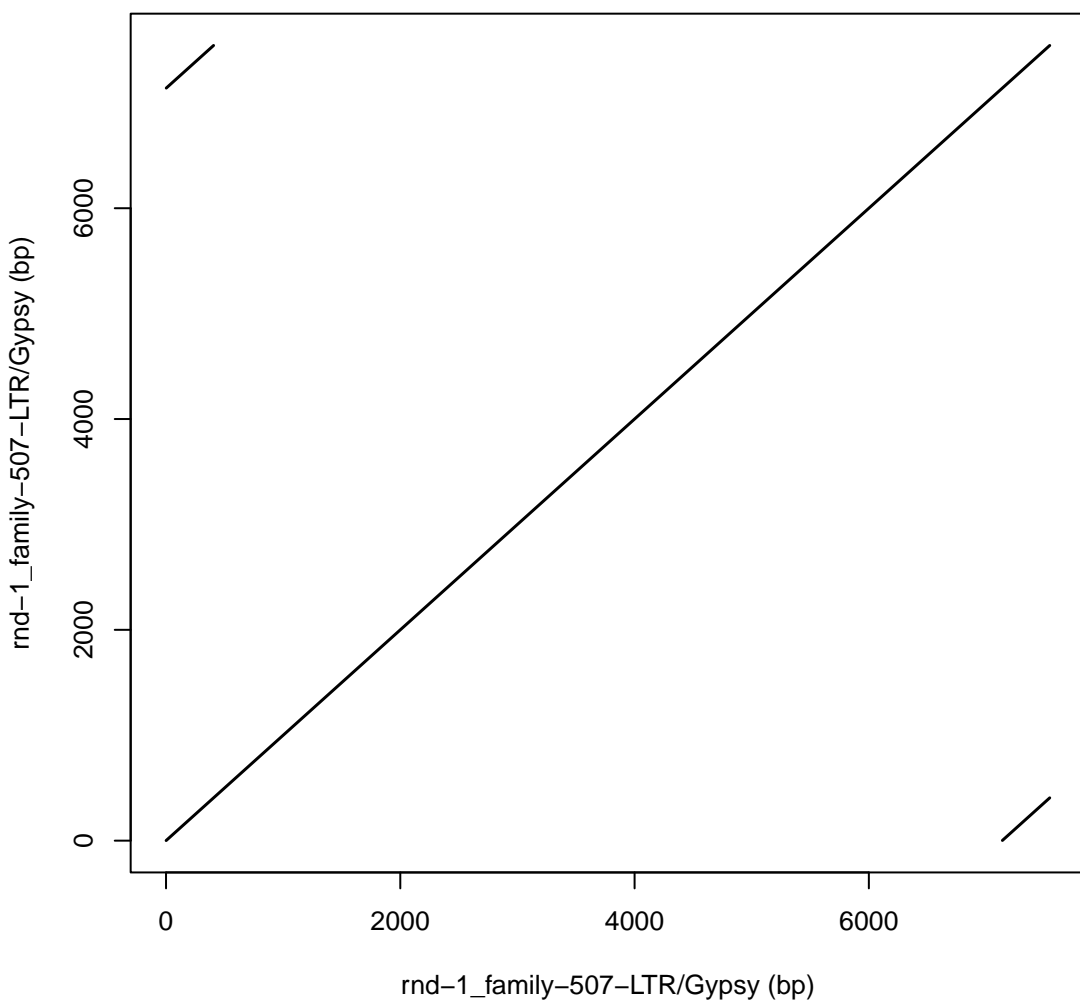
TE: rnd-1\_family-507-LTR/Gypsy  
consensus size: 7544bp; fragments: 240; full length: 8 ( $\geq 6789.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

