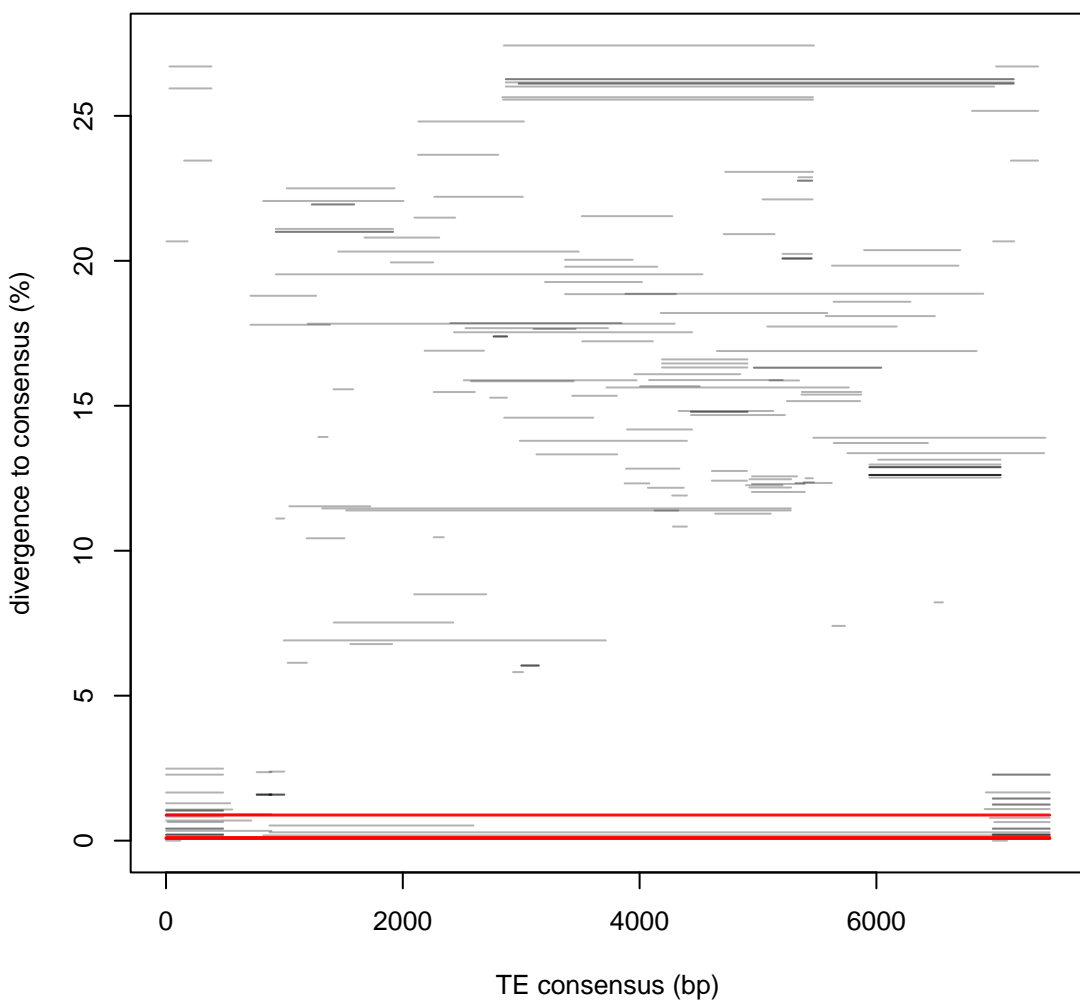
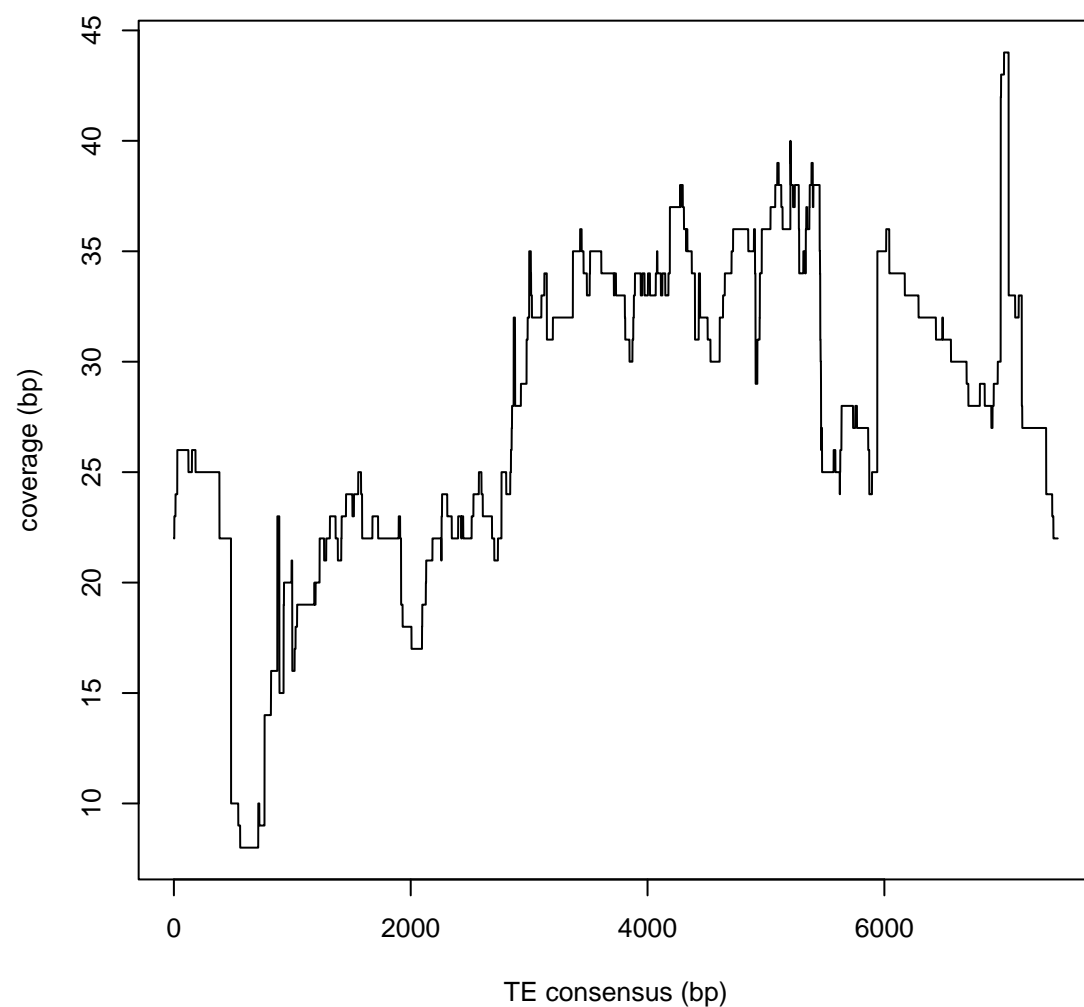


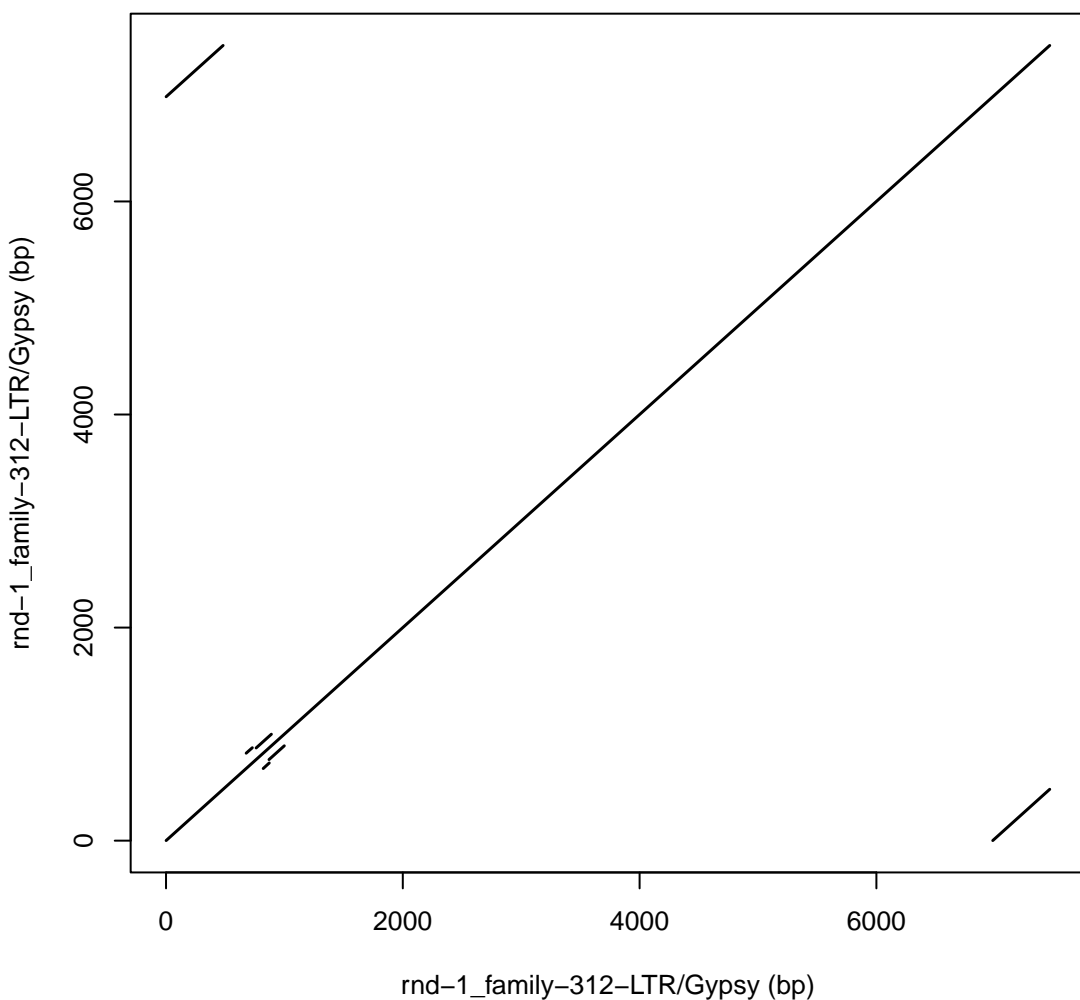
TE: rnd-1\_family-312-LTR/Gypsy  
consensus size: 7464bp; fragments: 196; full length: 4 ( $\geq 6717.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

