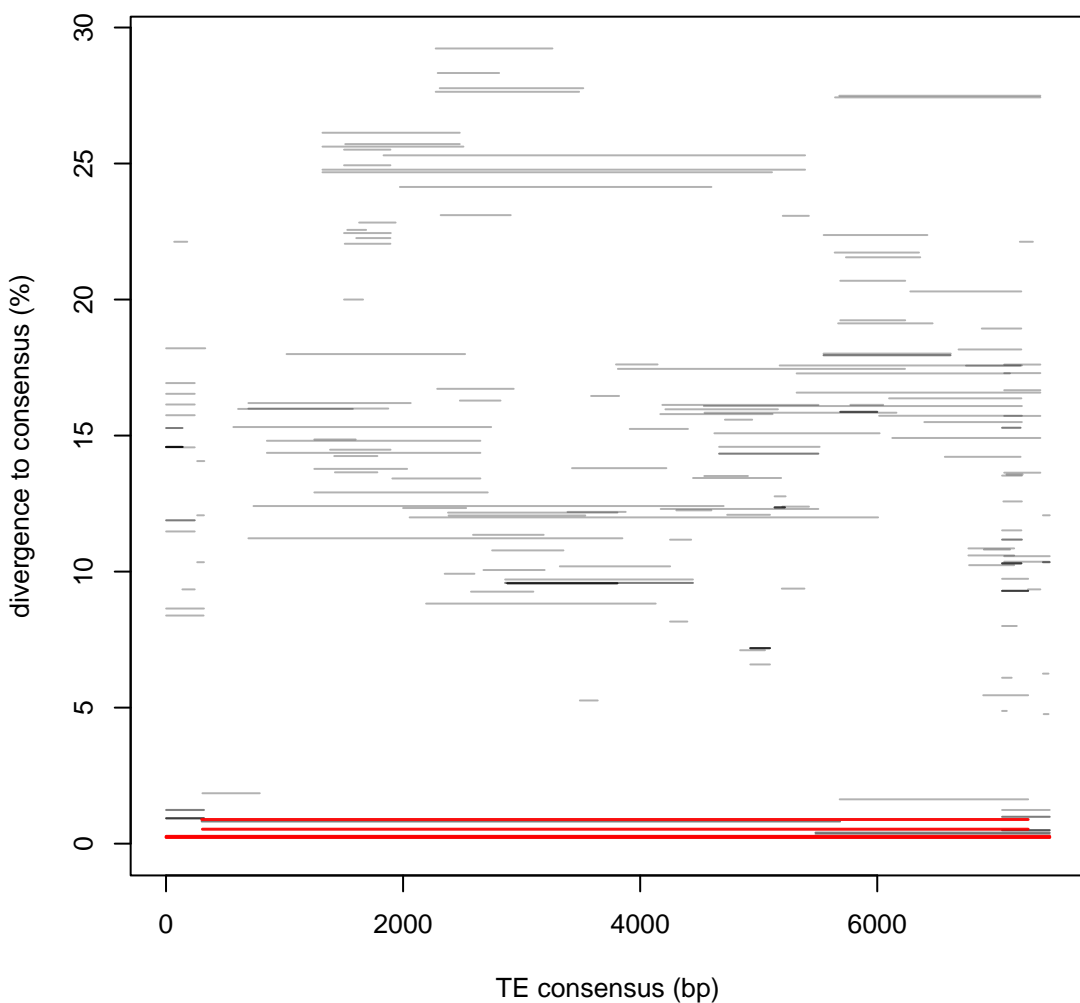
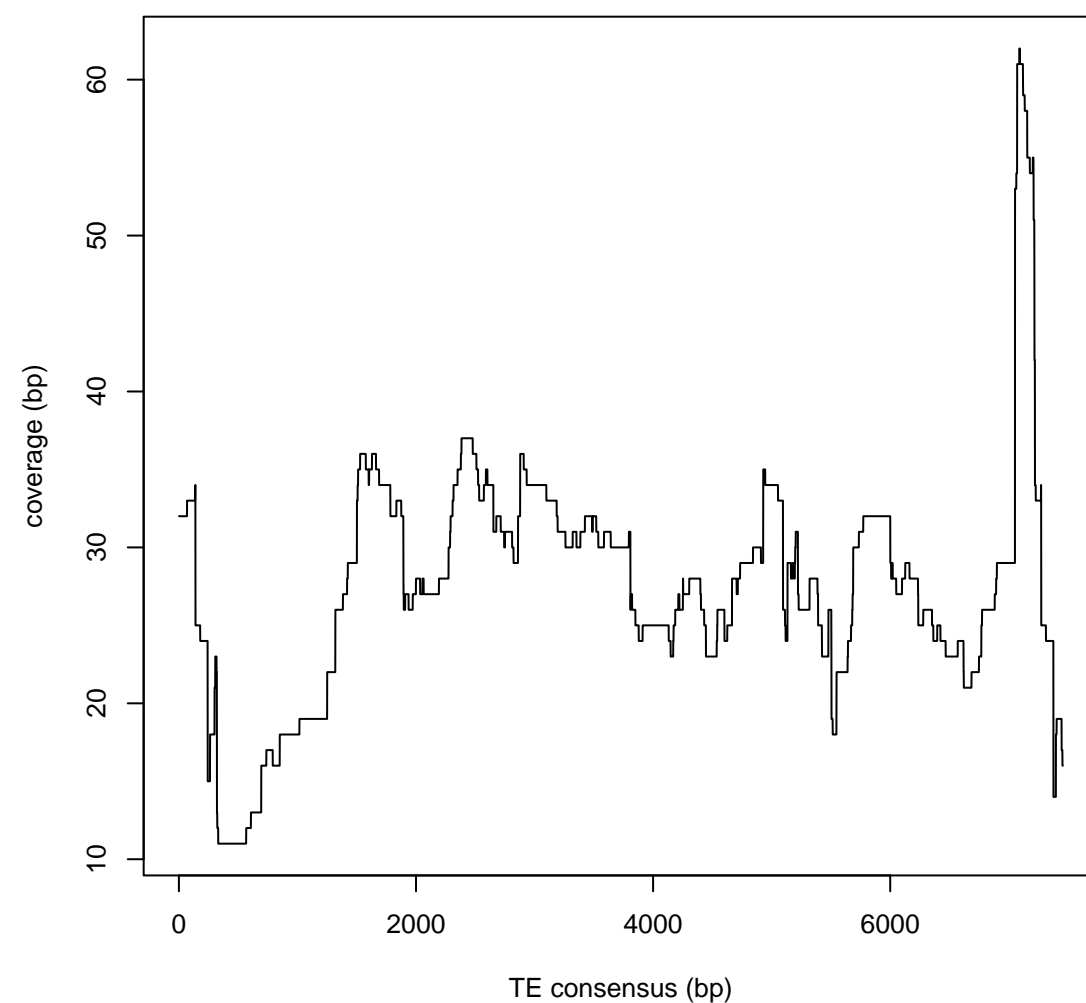


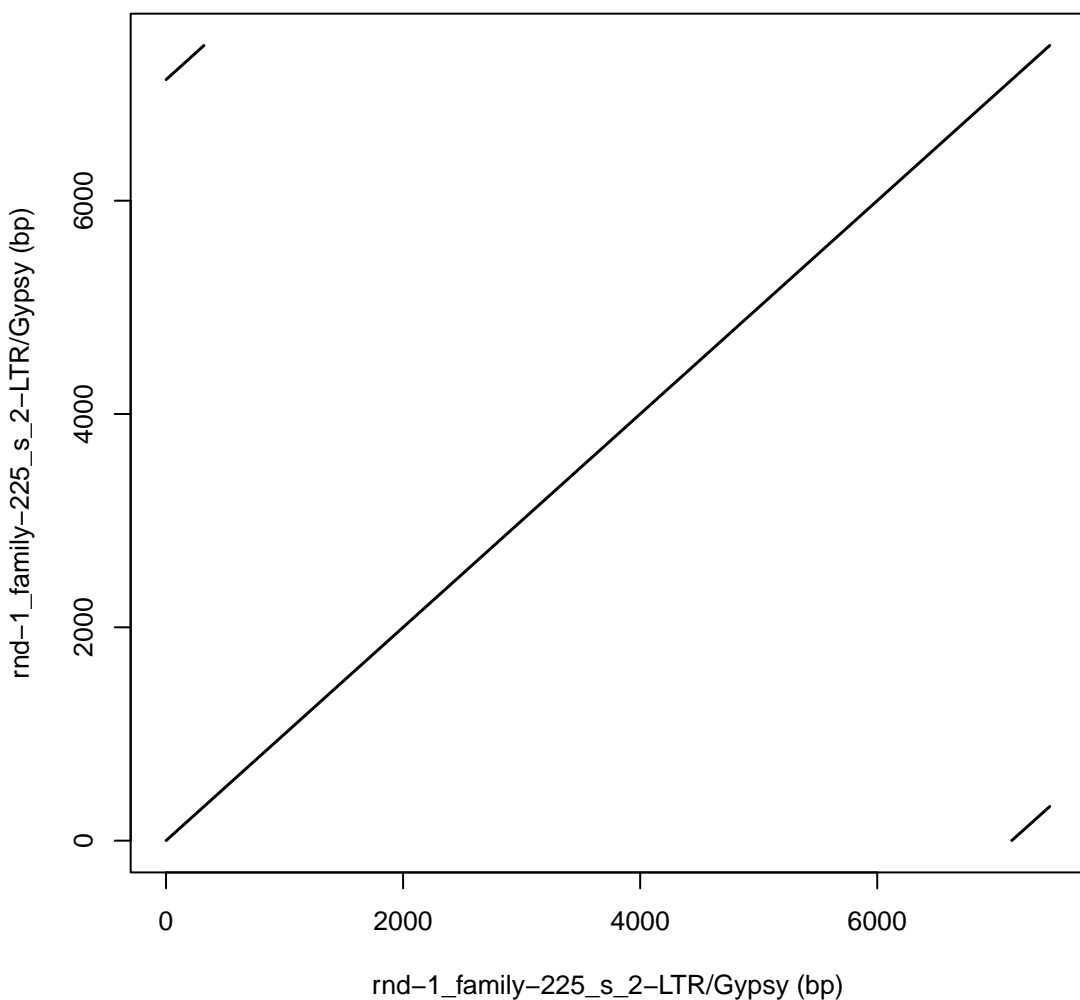
TE: rnd-1\_family-225\_s\_2-LTR/Gypsy  
consensus size: 7455bp; fragments: 213; full length: 5 ( $\geq 6709.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

