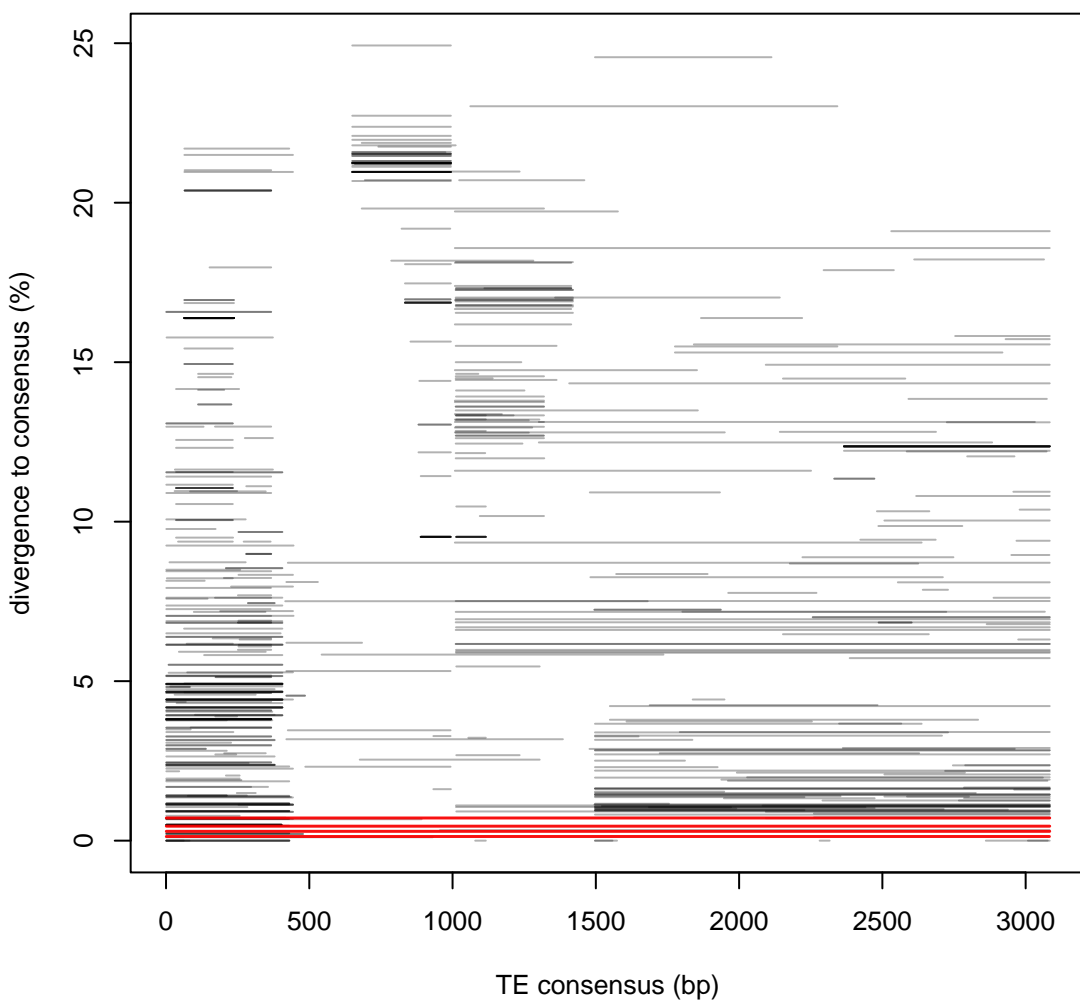
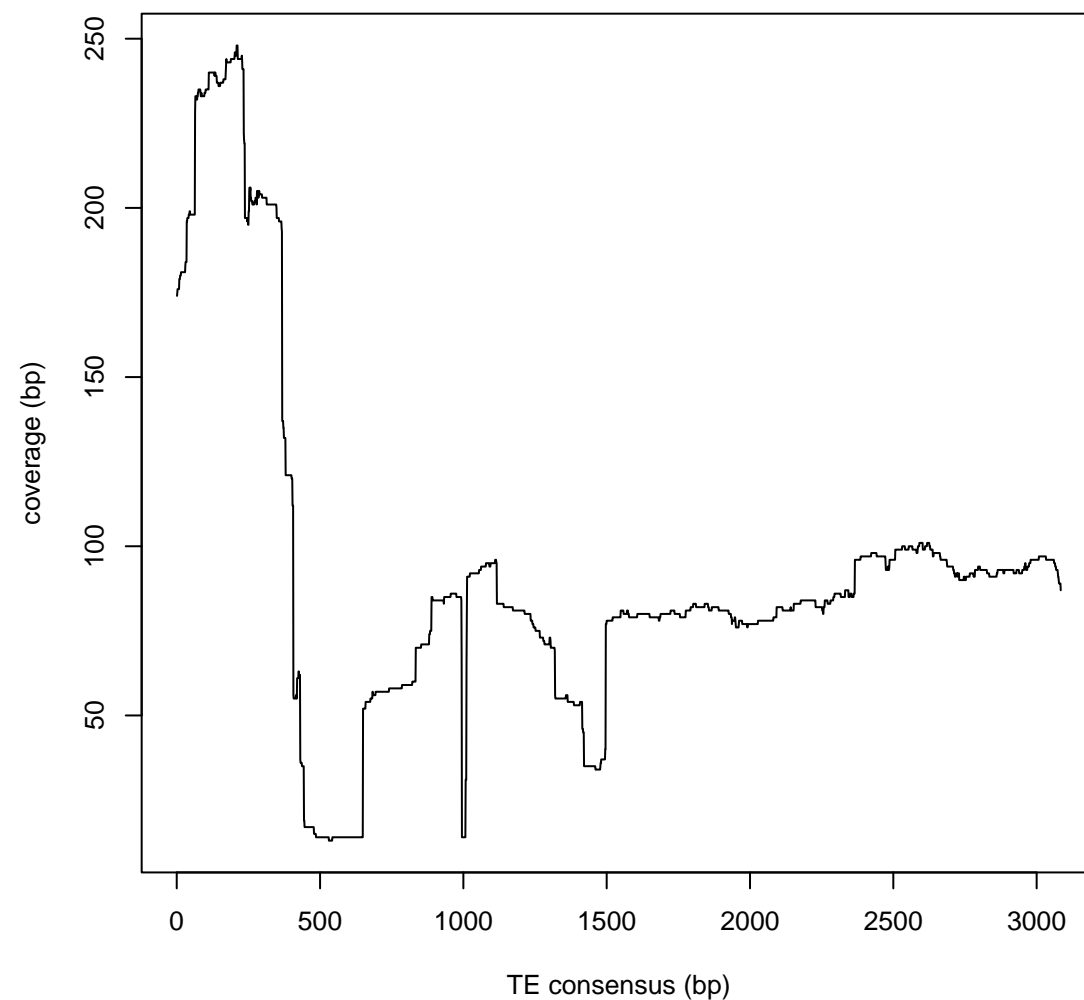


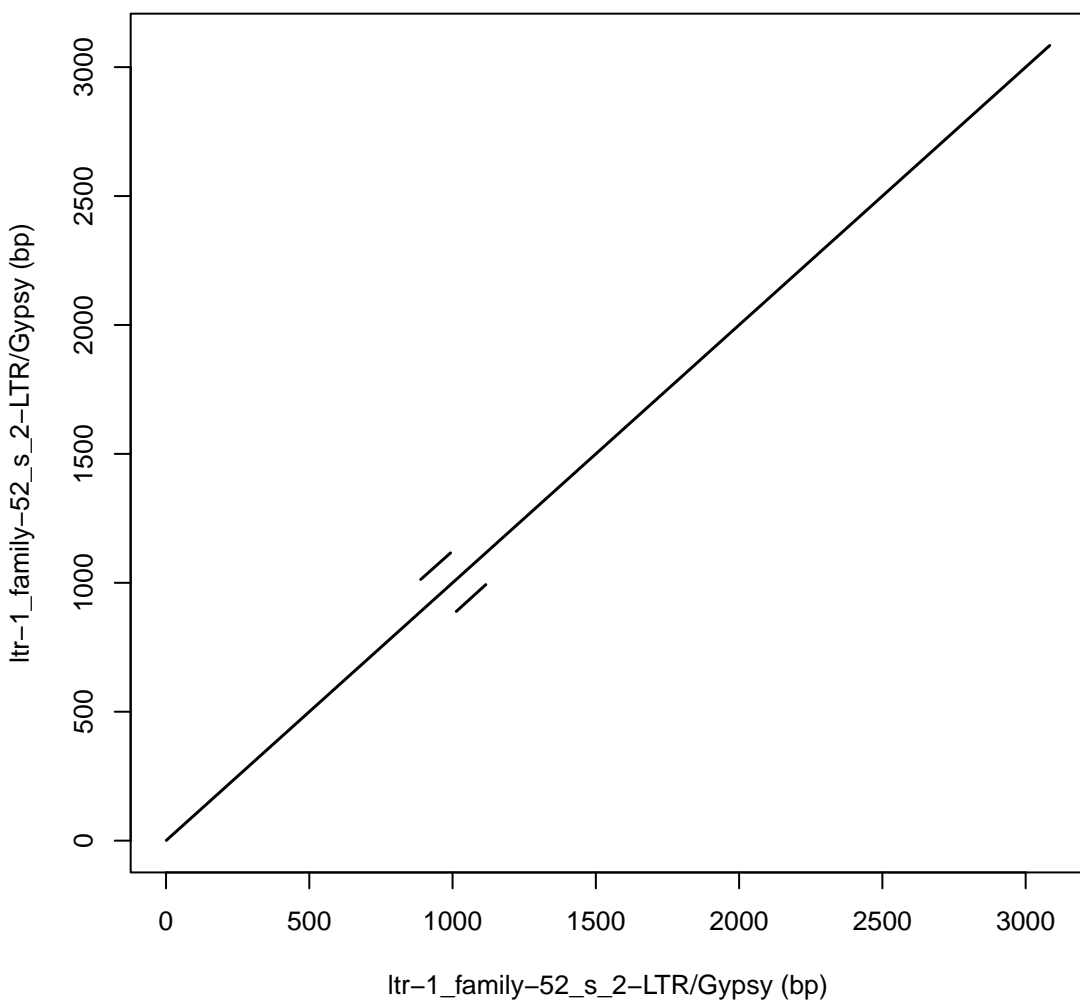
TE: ltr-1\_family-52\_s\_2-LTR/Gypsy  
consensus size: 3084bp; fragments: 590; full length: 4 ( $\geq 2775.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

