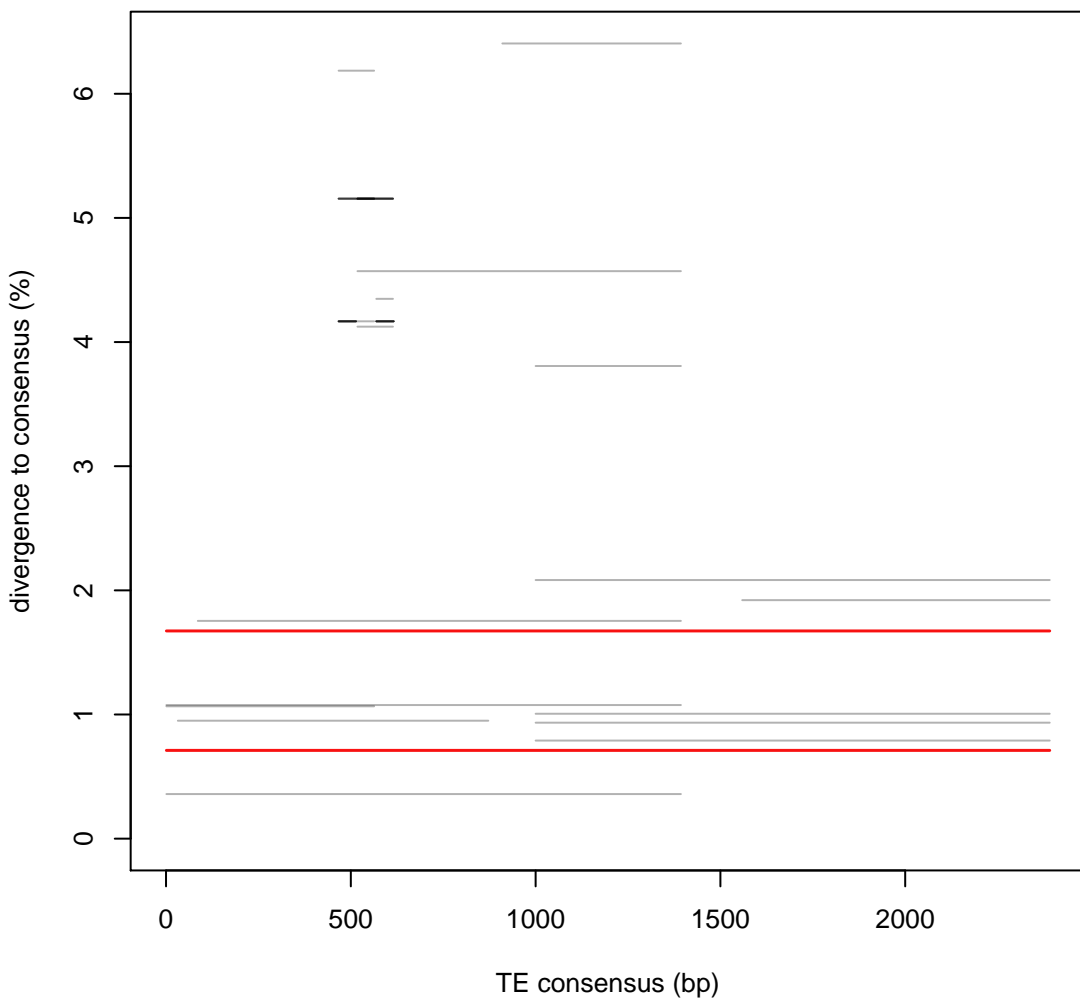
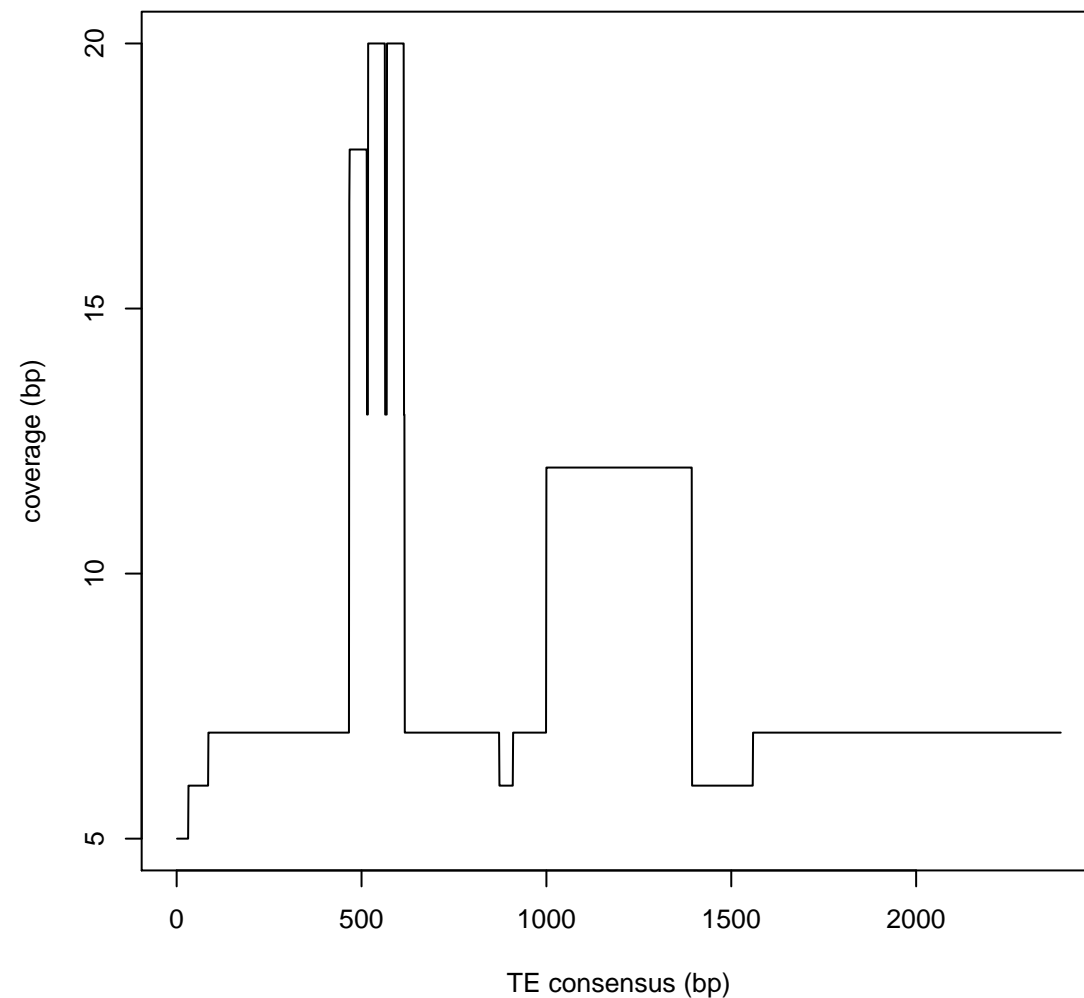


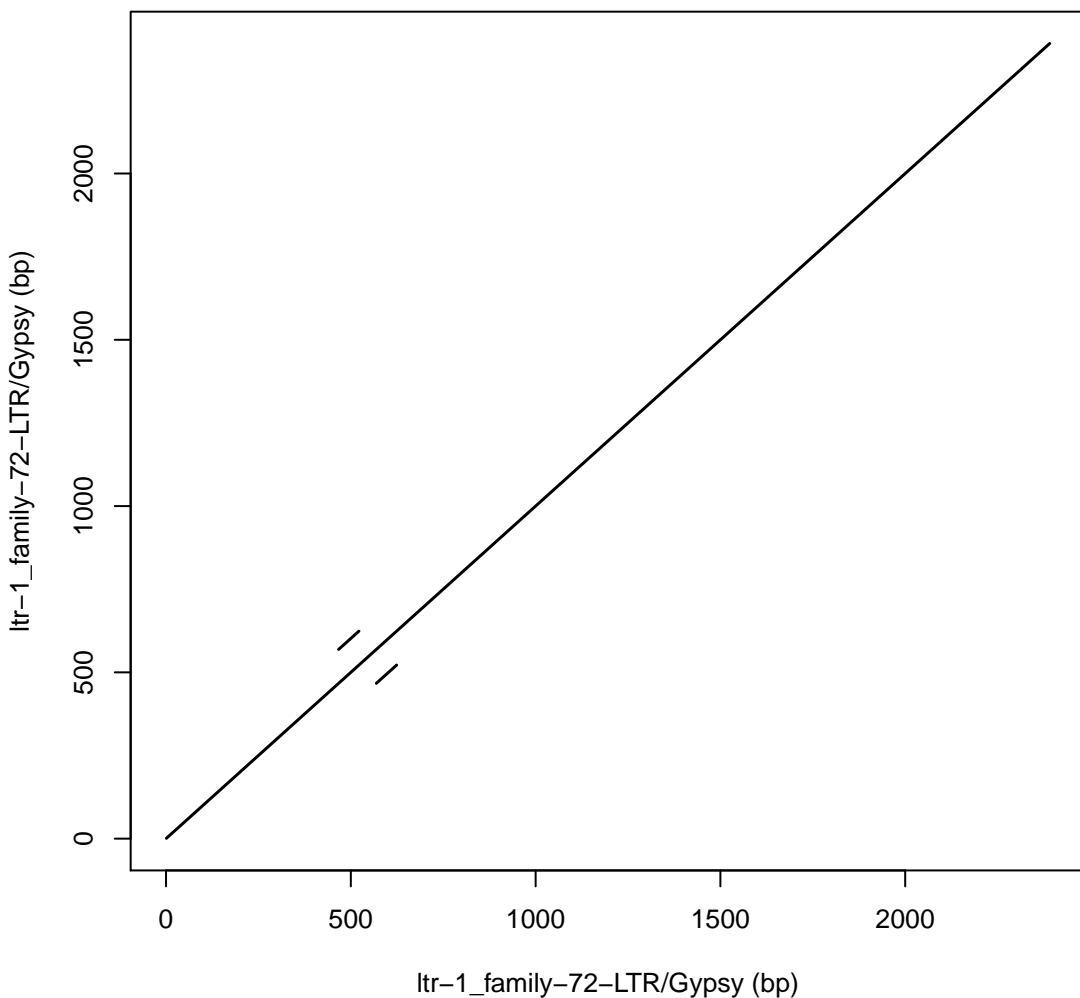
TE: ltr-1\_family-72-LTR/Gypsy  
consensus size: 2391bp; fragments: 39; full length: 2 ( $\geq 2151.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

