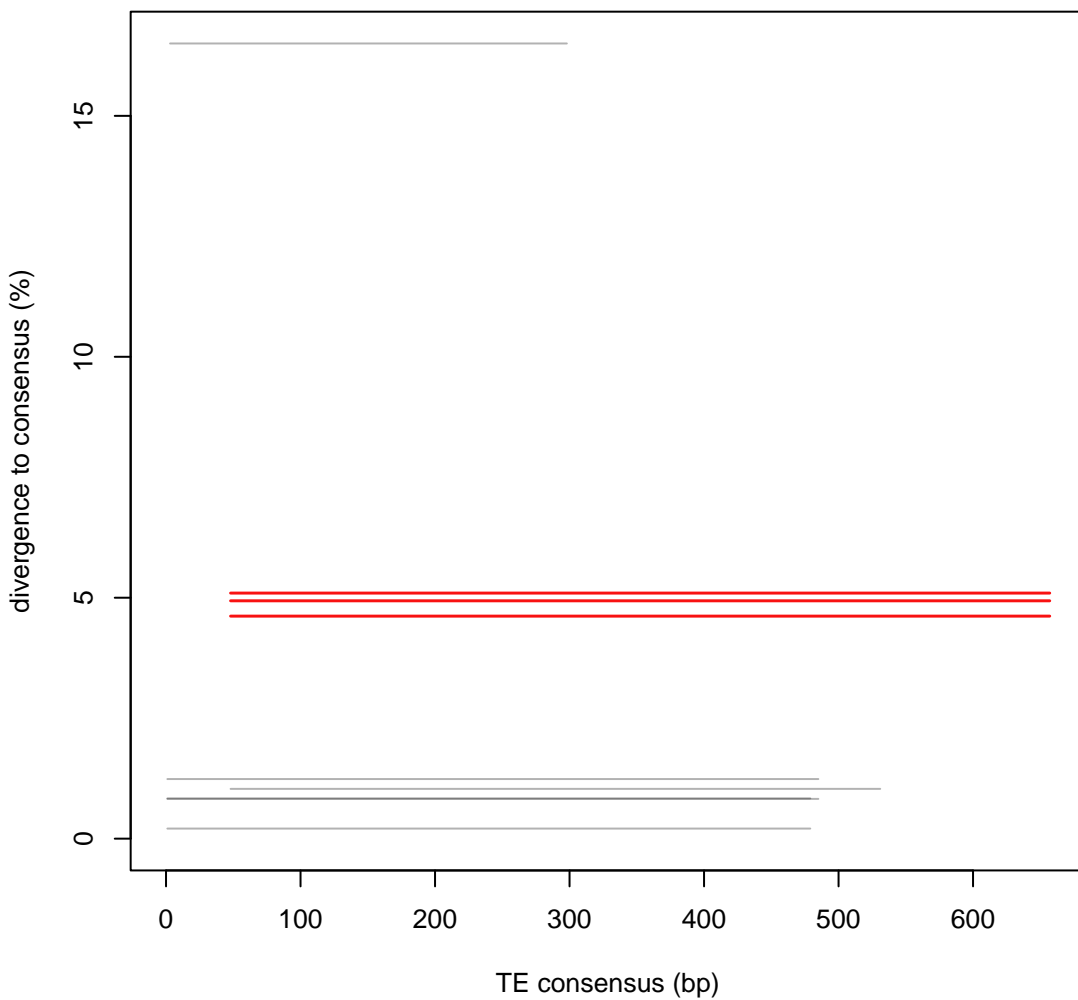
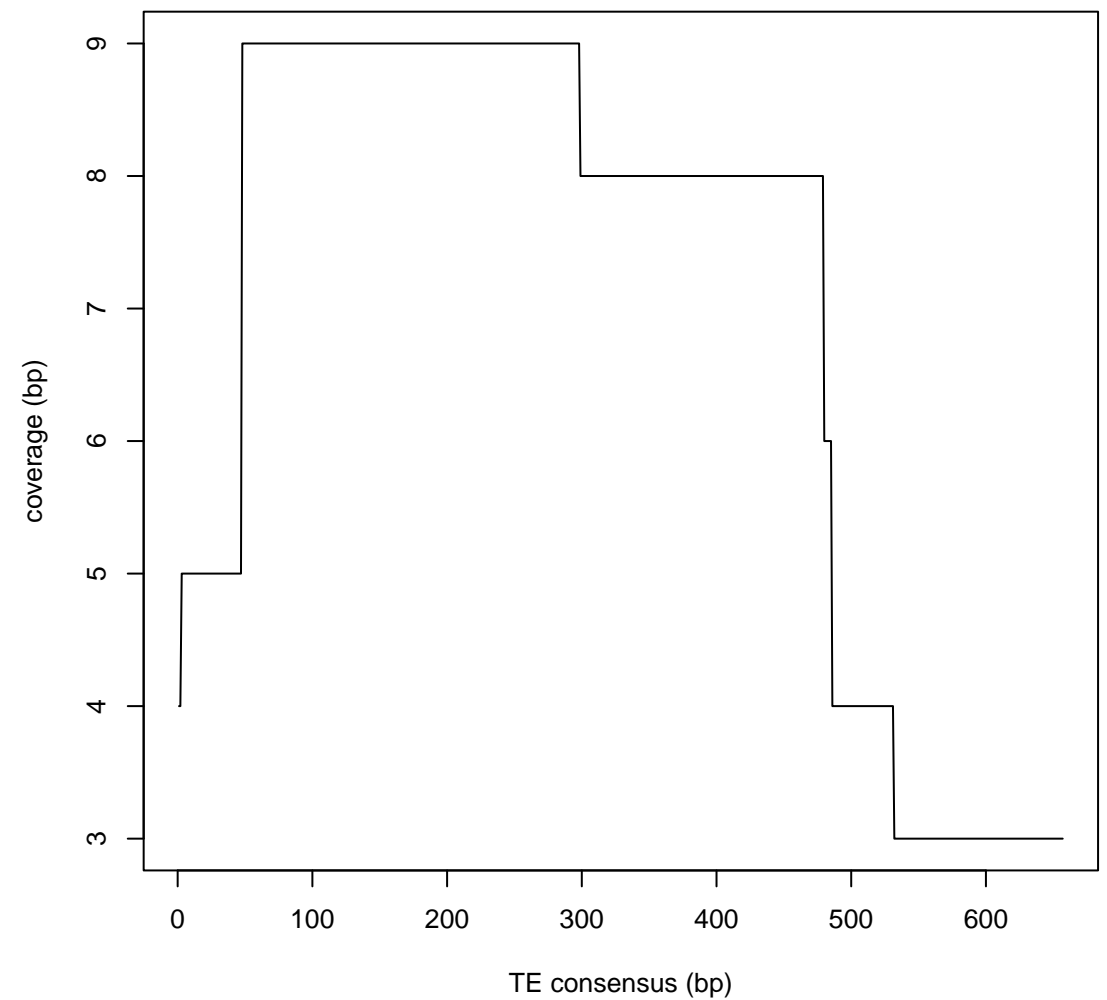


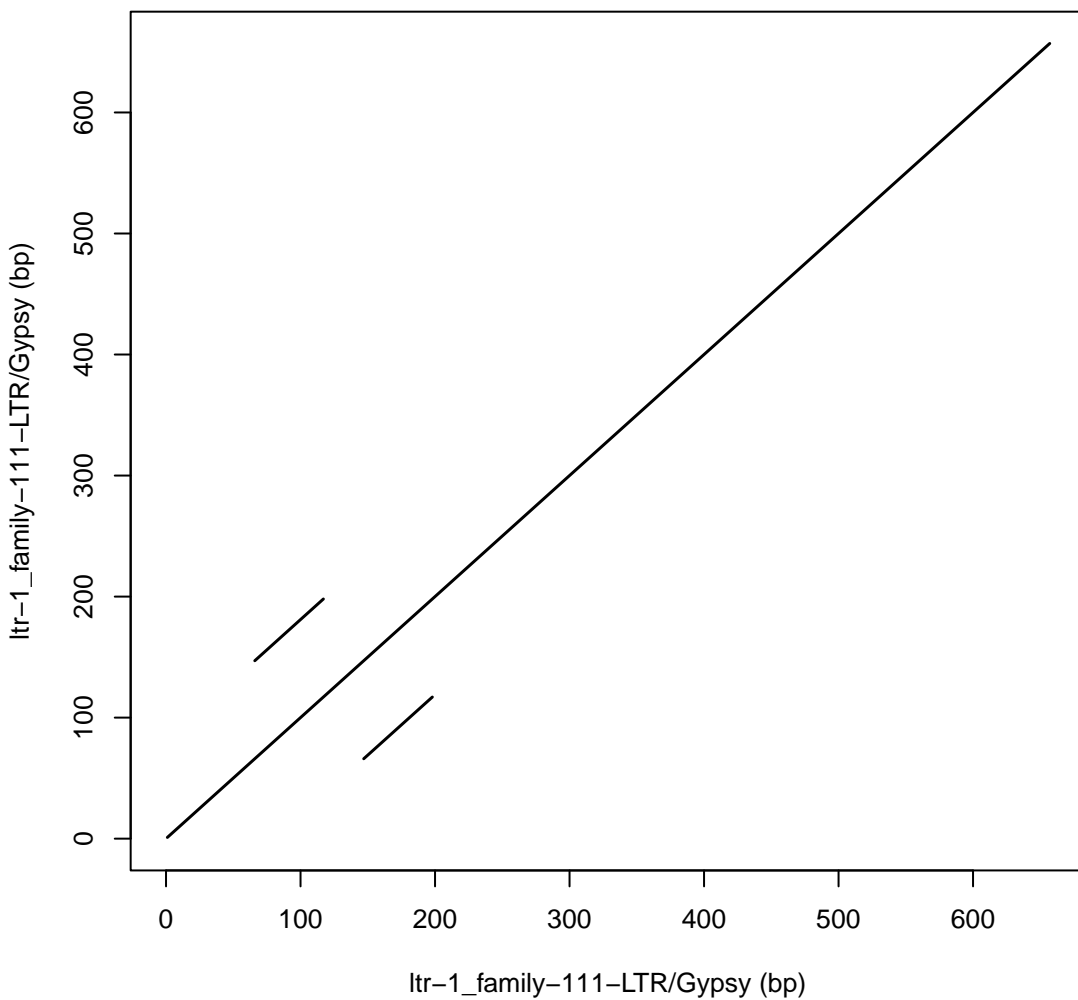
TE: ltr-1\_family-111-LTR/Gypsy  
consensus size: 657bp; fragments: 9; full length: 3 (>=591.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

