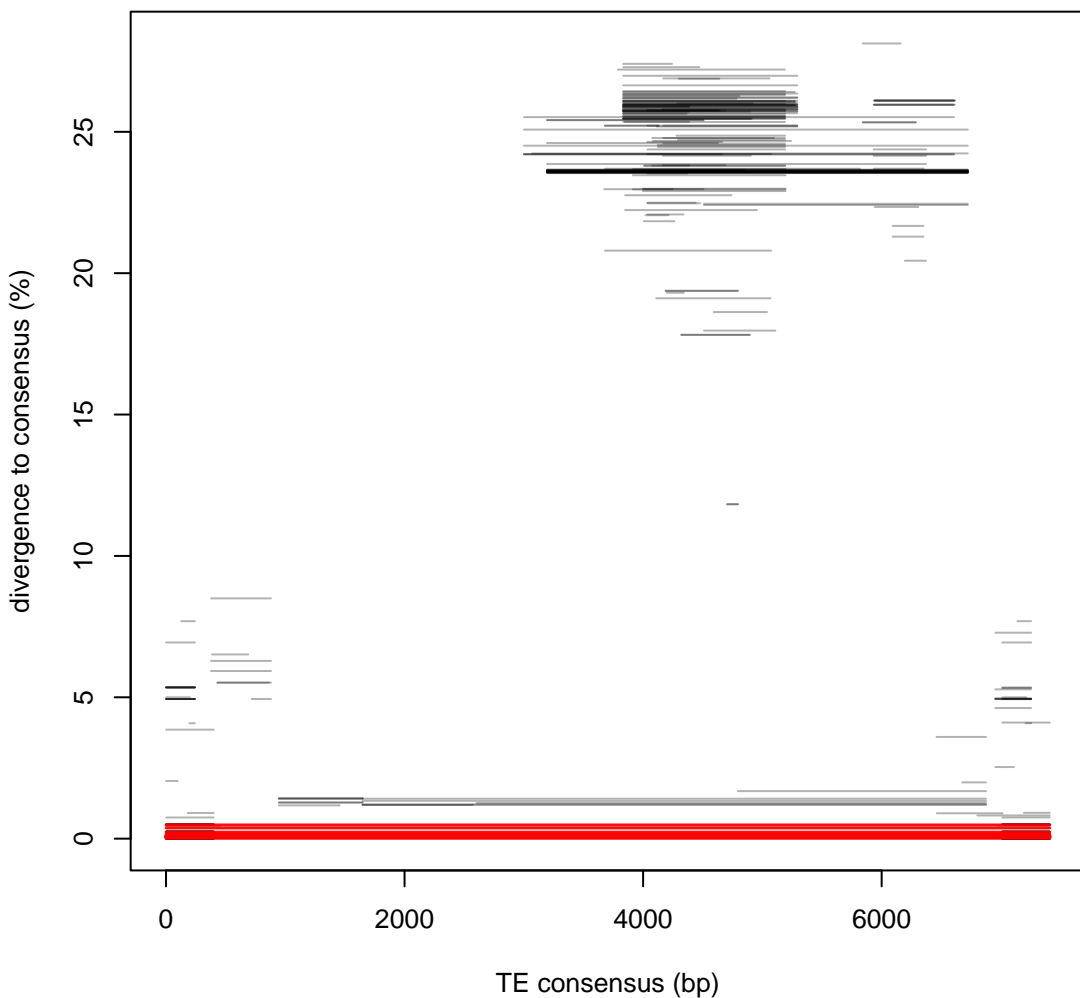
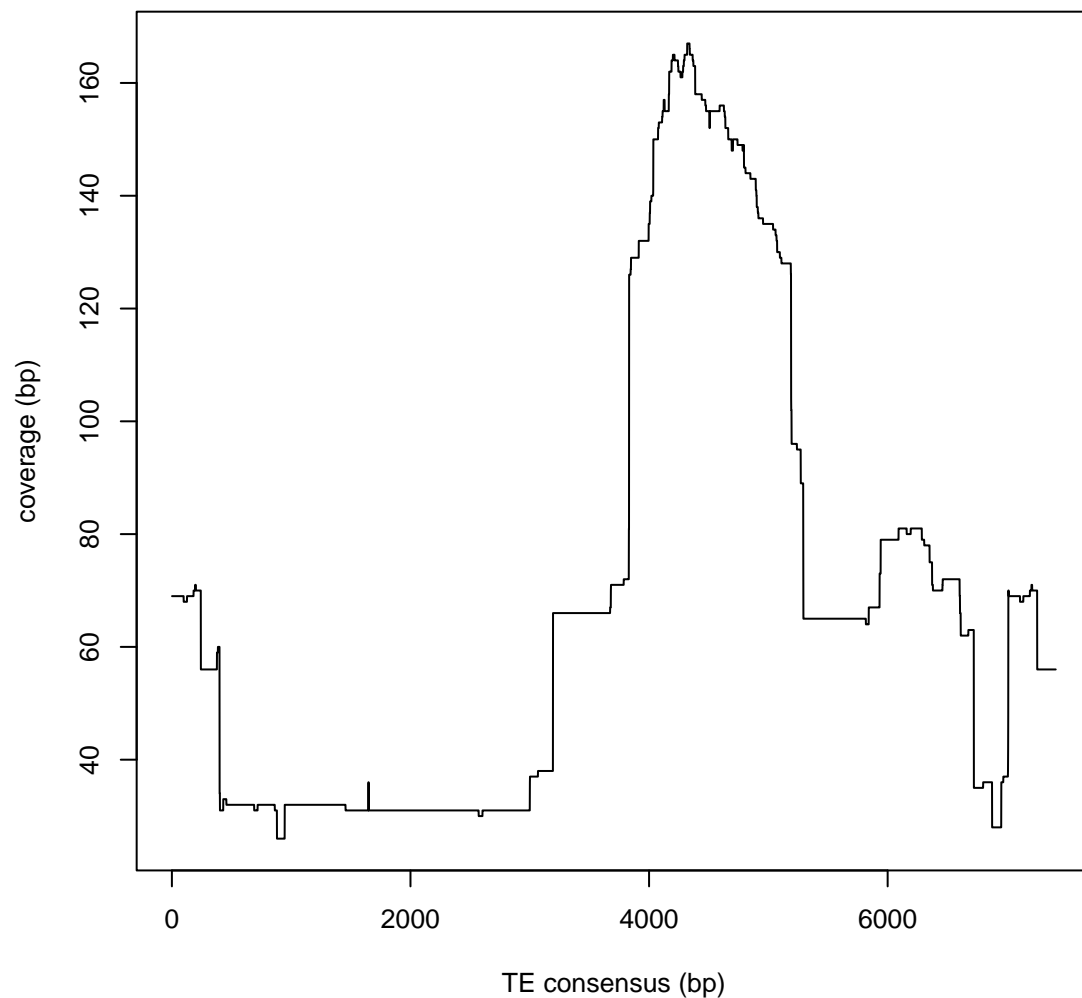


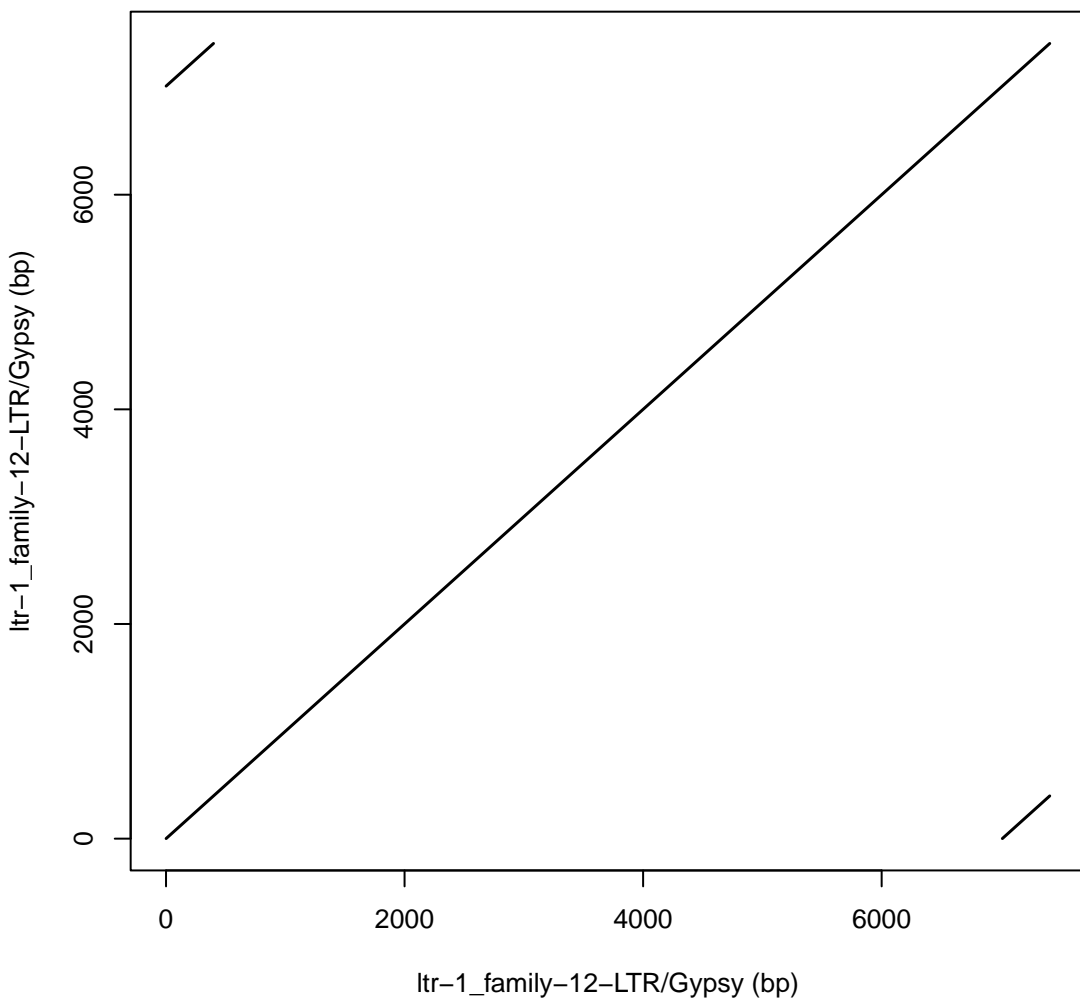
TE: ltr-1\_family-12-LTR/Gypsy  
consensus size: 7409bp; fragments: 307; full length: 26 (>=6668.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

