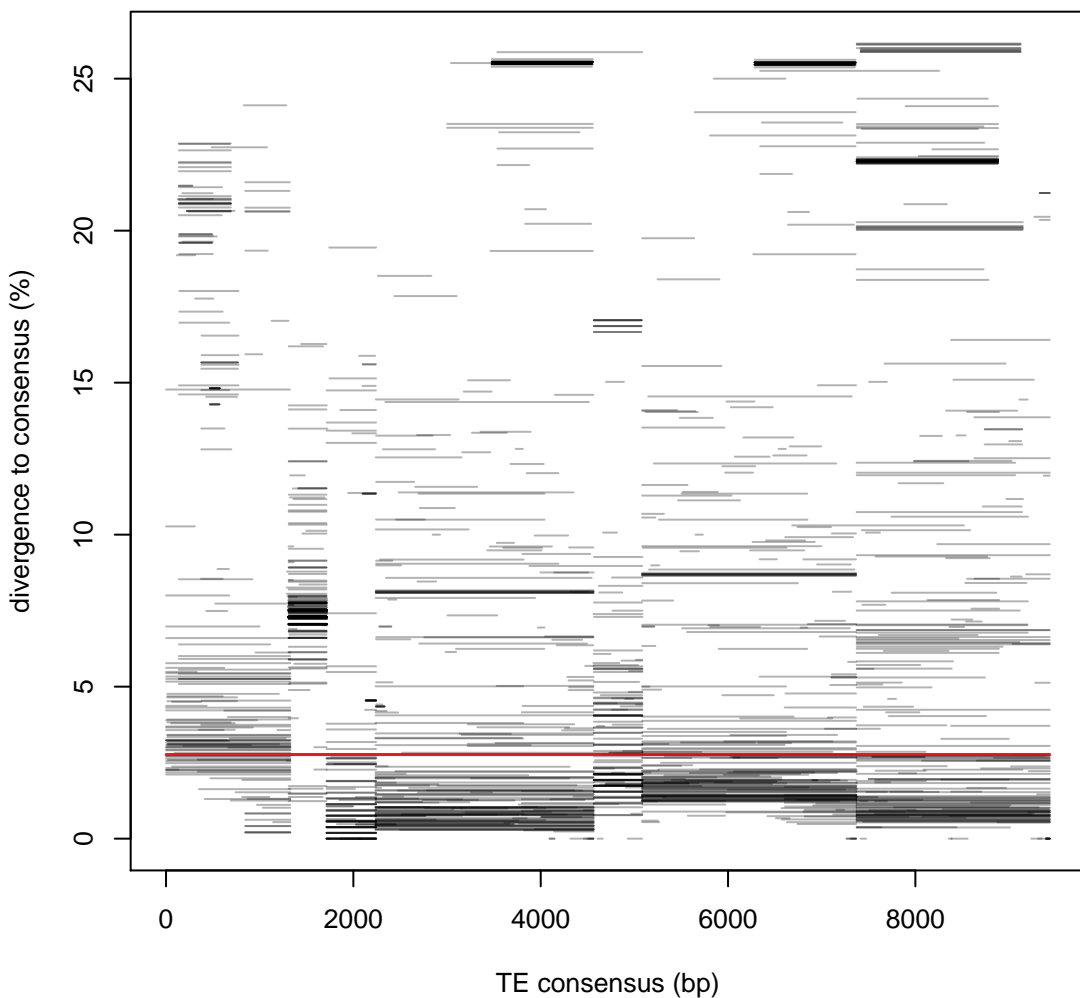
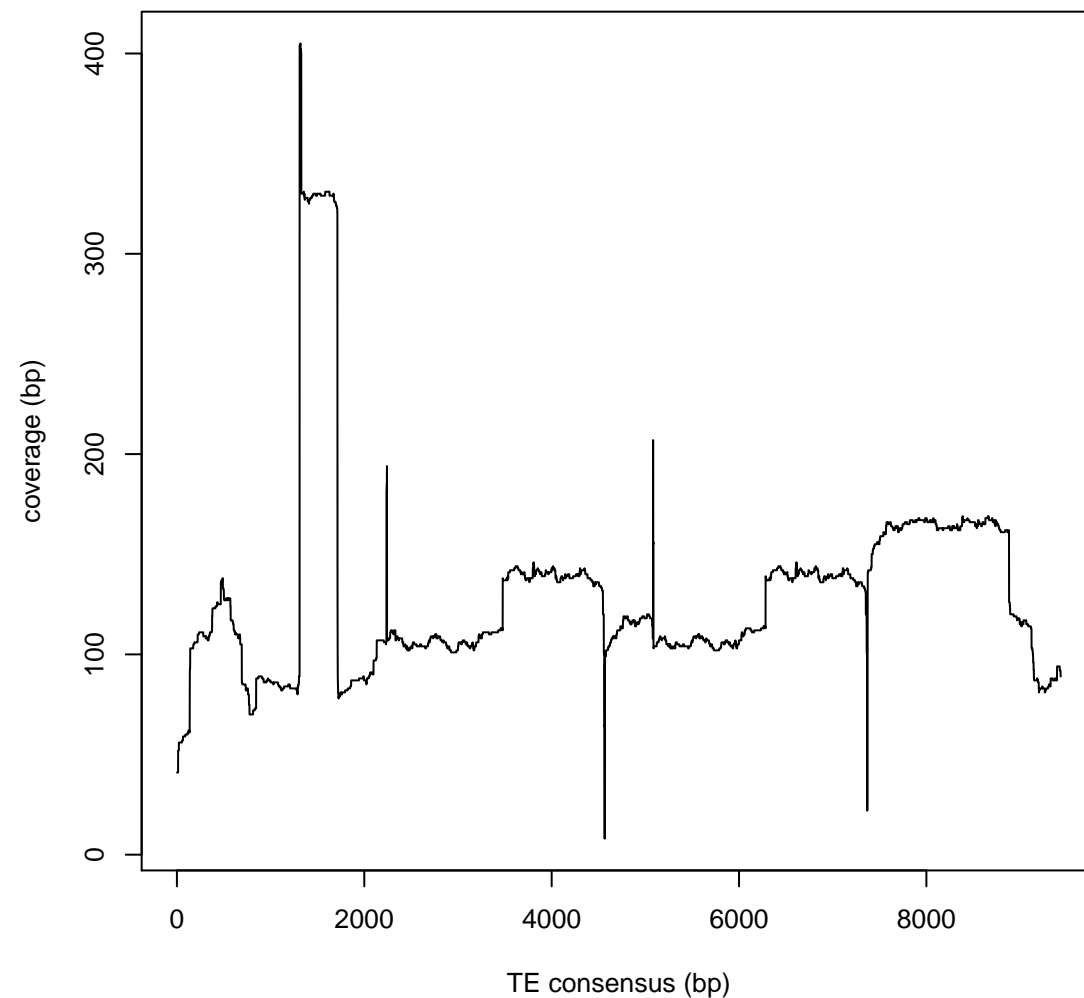


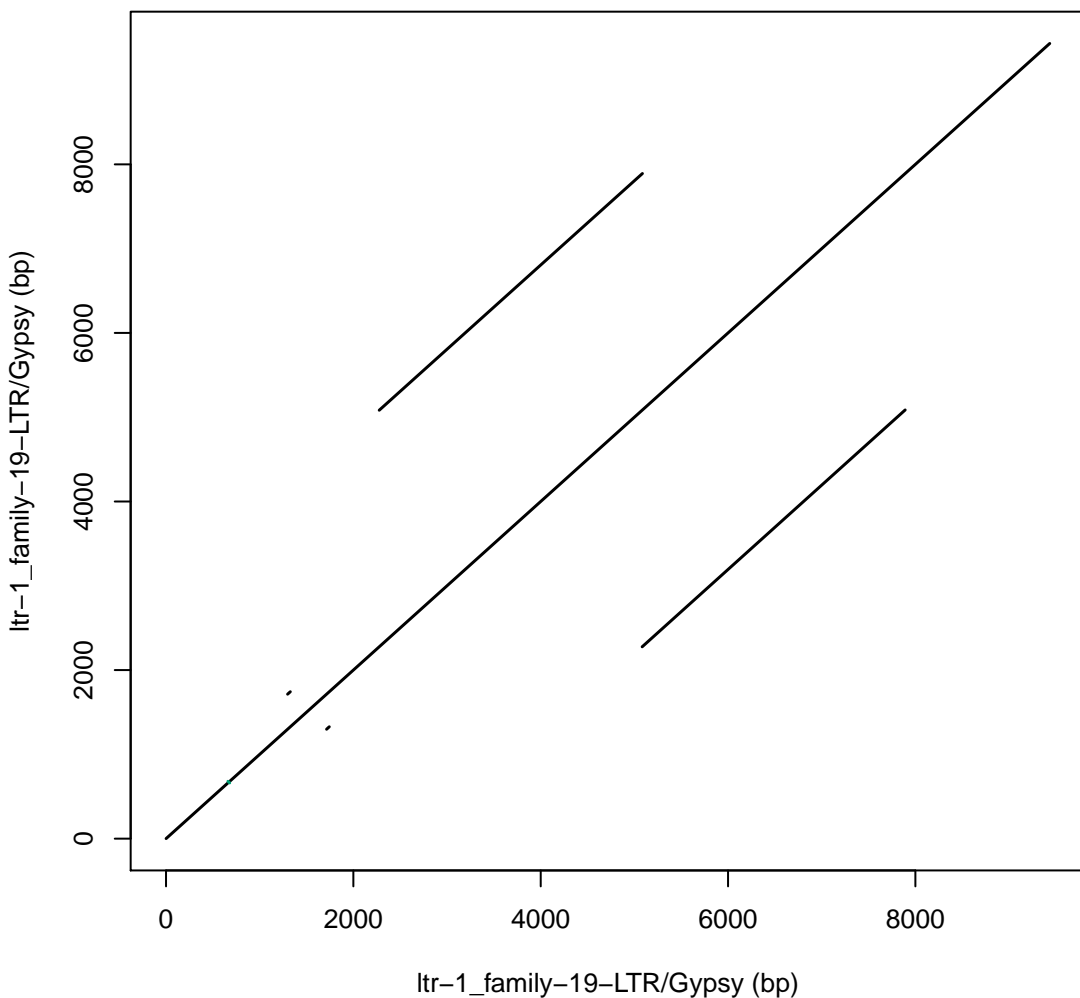
TE: ltr-1_family-19-LTR/Gypsy
consensus size: 9434bp; fragments: 1510; full length: 1 (>=8490.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

