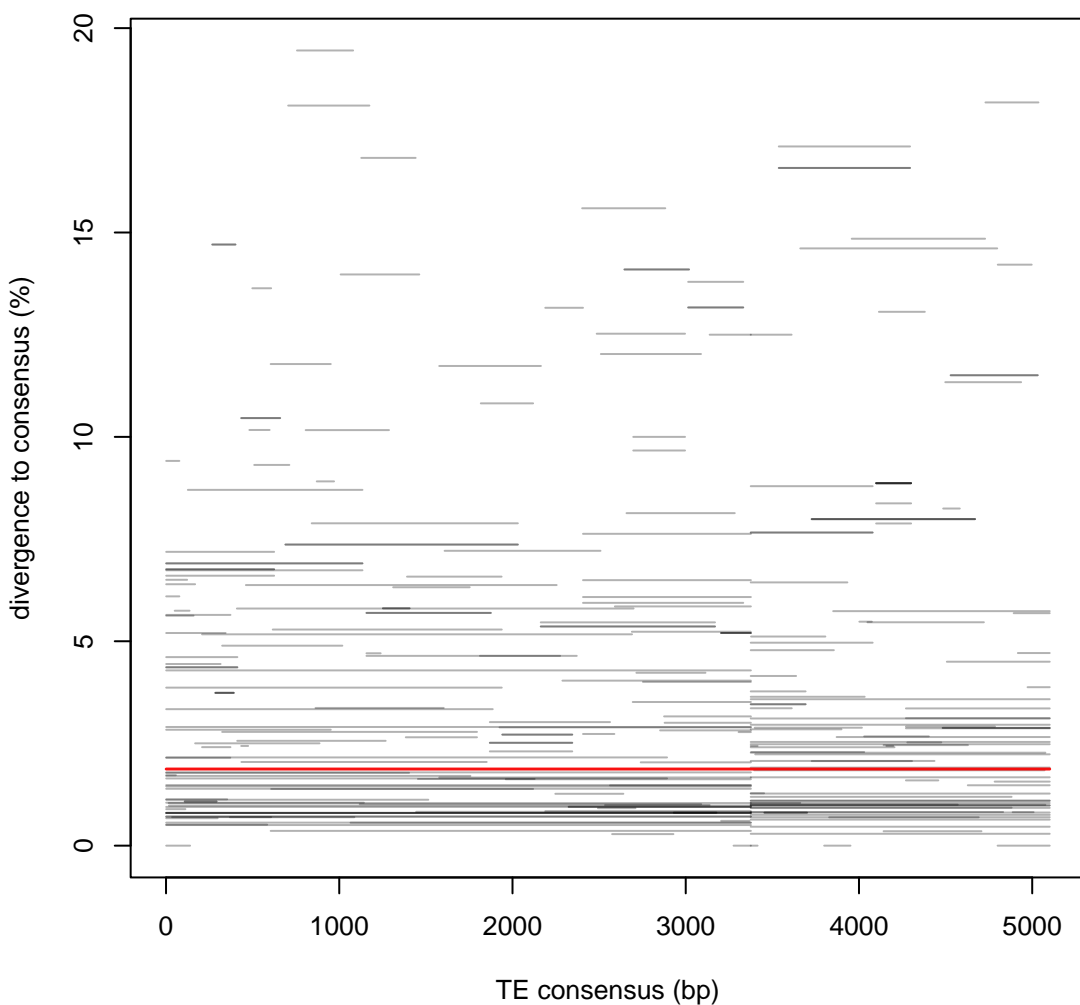
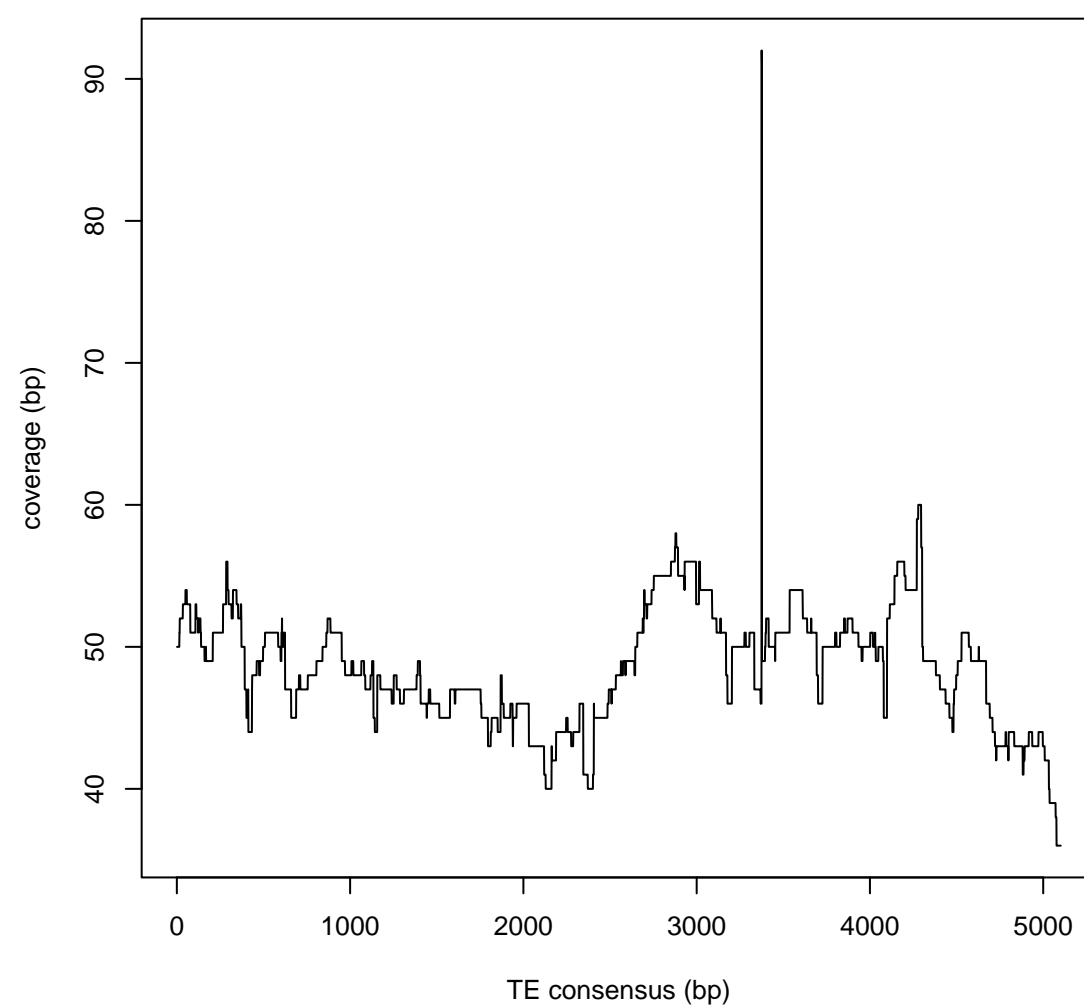


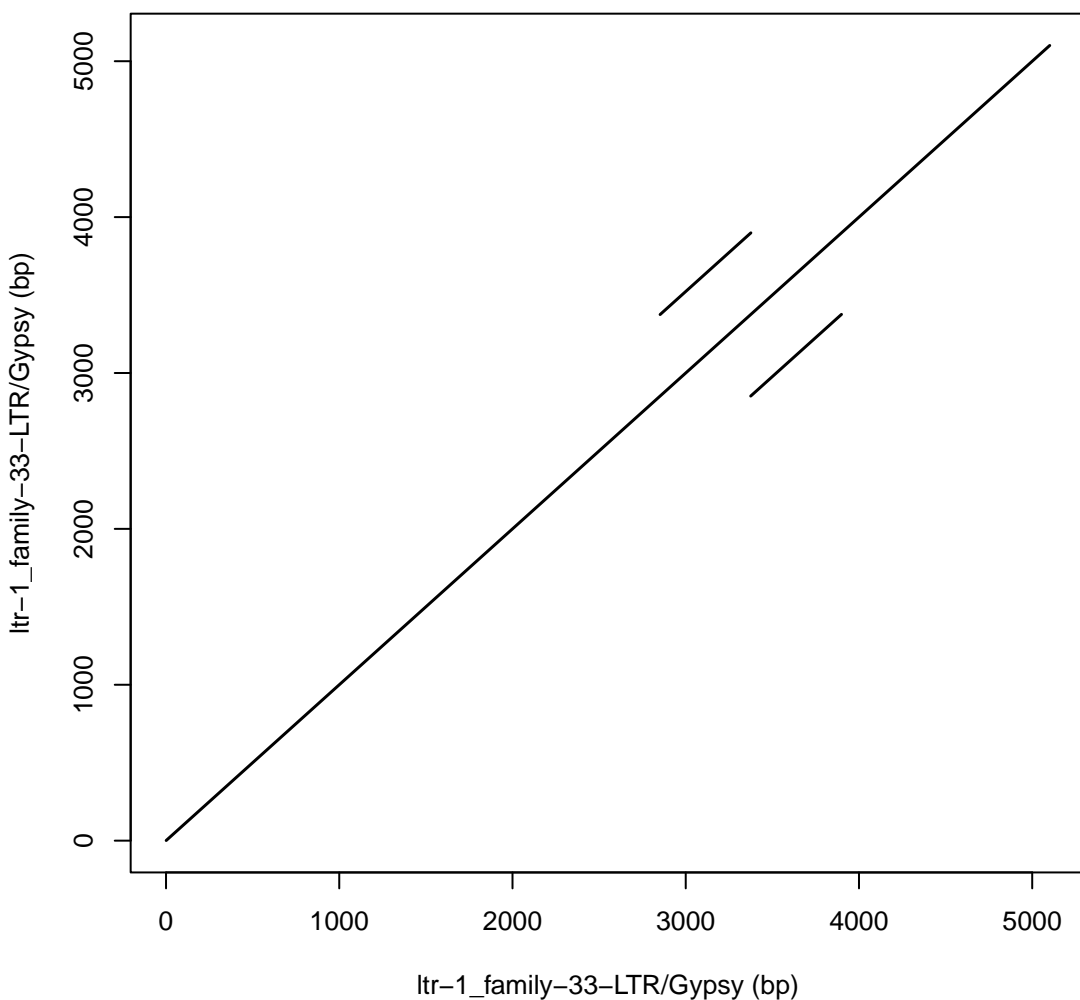
TE: ltr-1\_family-33-LTR/Gypsy  
consensus size: 5101bp; fragments: 267; full length: 1 (>=4590.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

