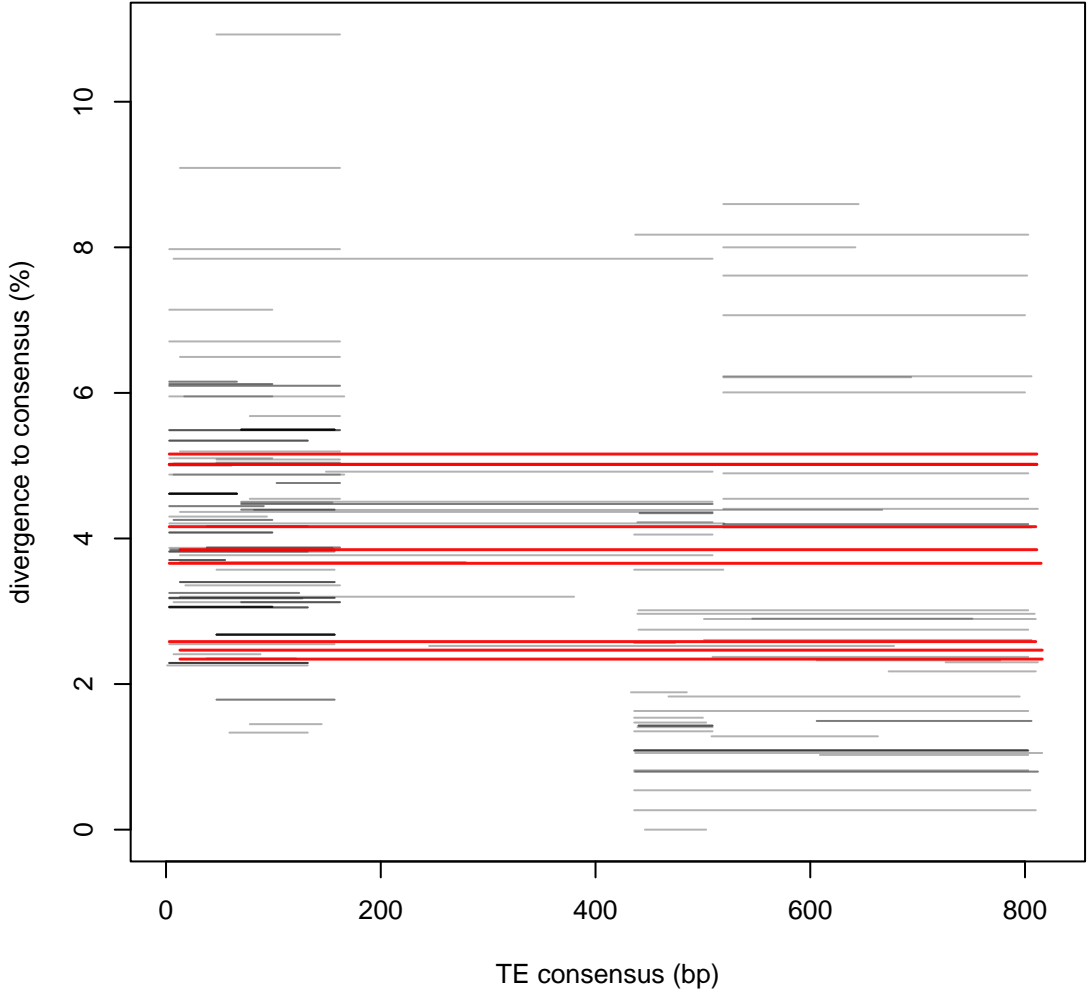
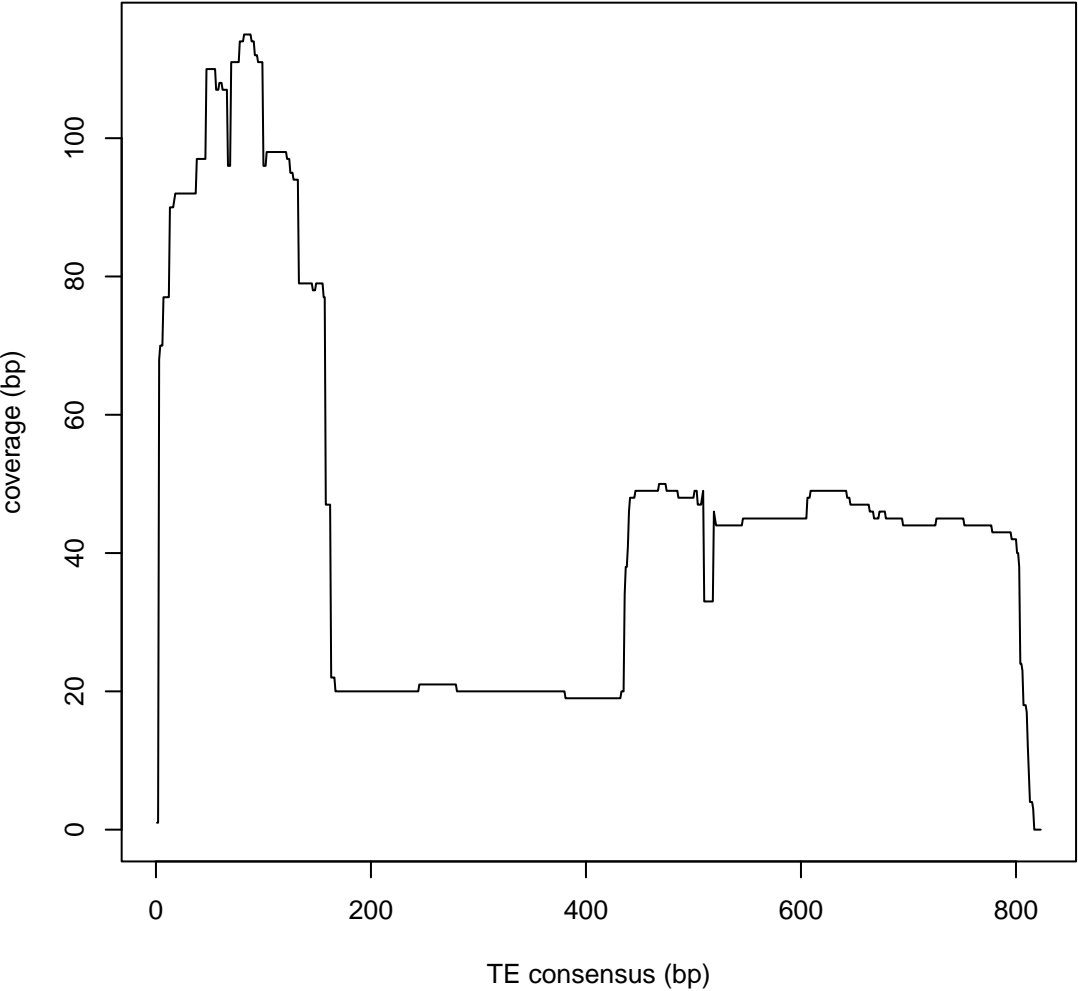


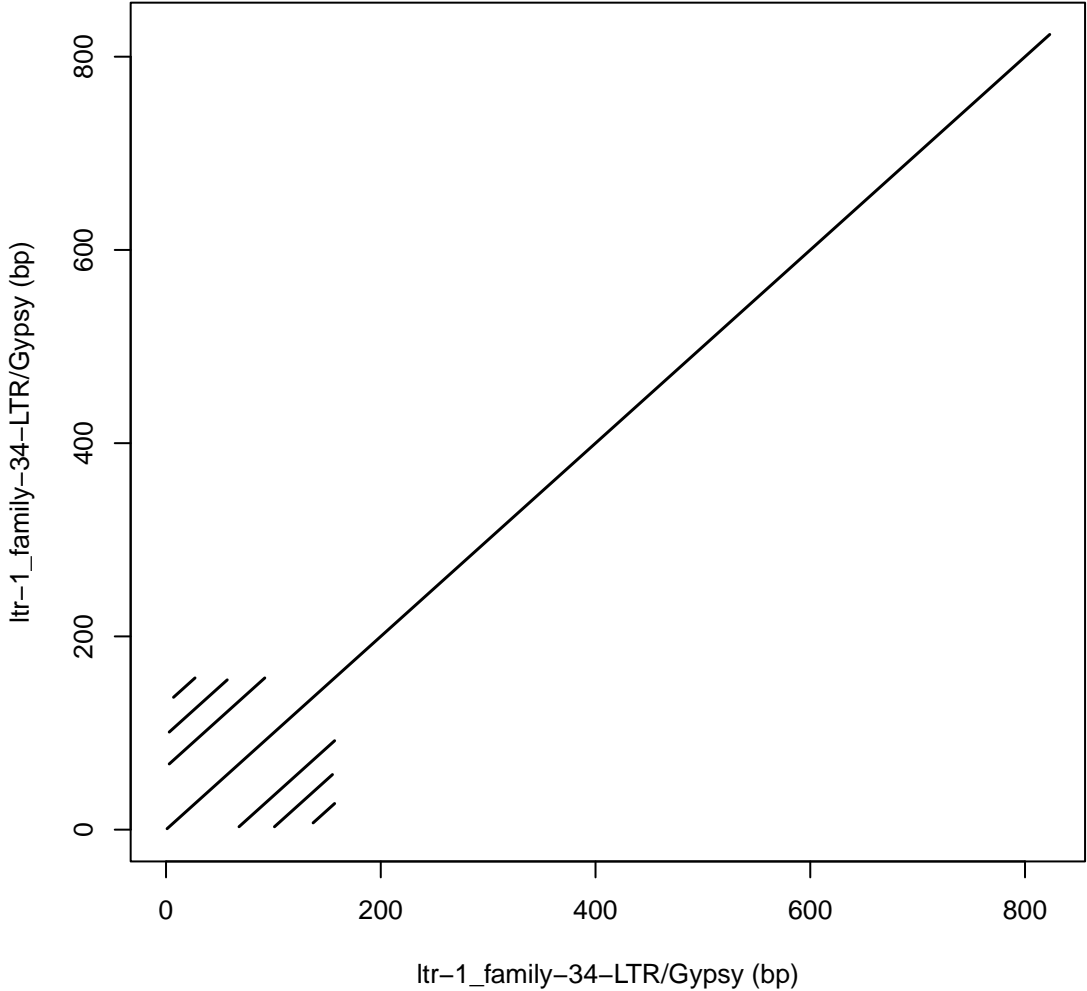
TE: ltr-1\_family-34-LTR/Gypsy  
consensus size: 823bp; fragments: 189; full length: 9 (>=740.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

