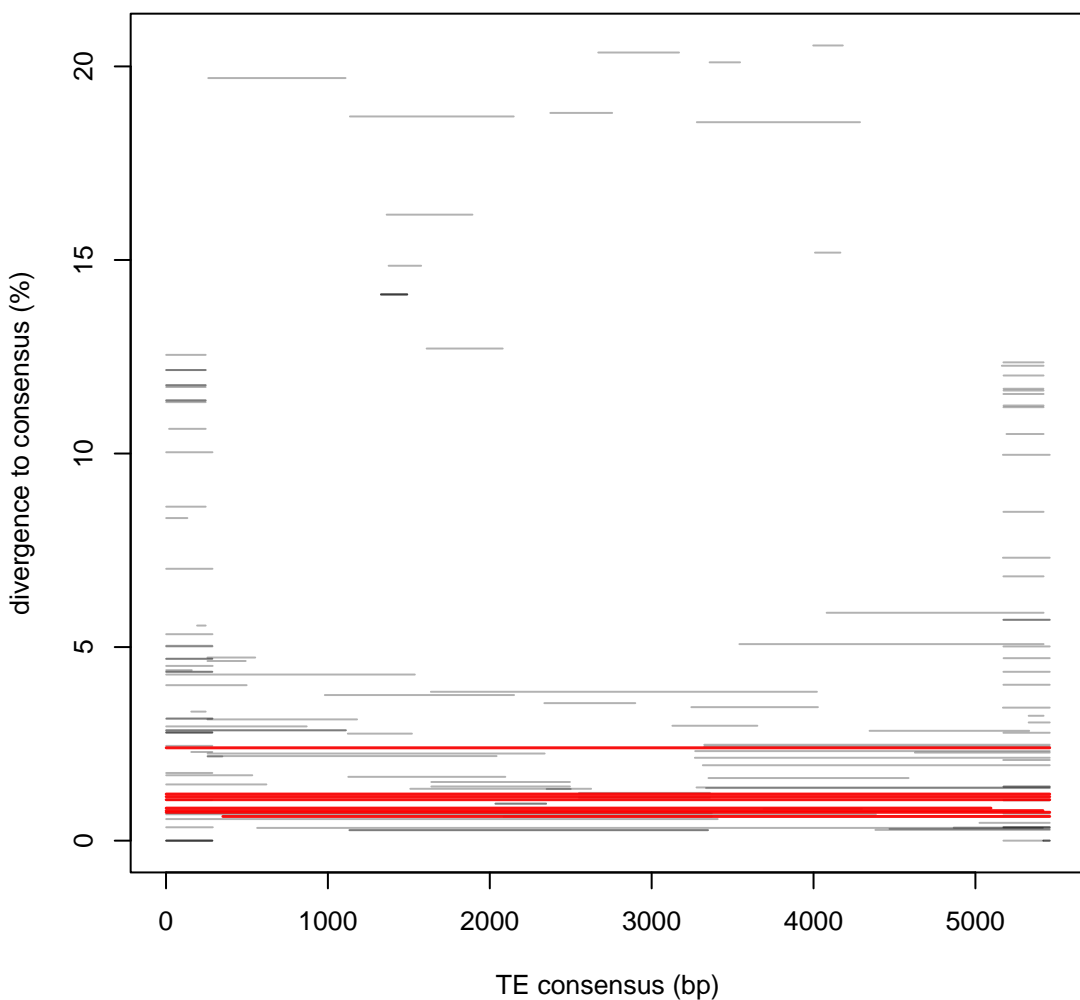
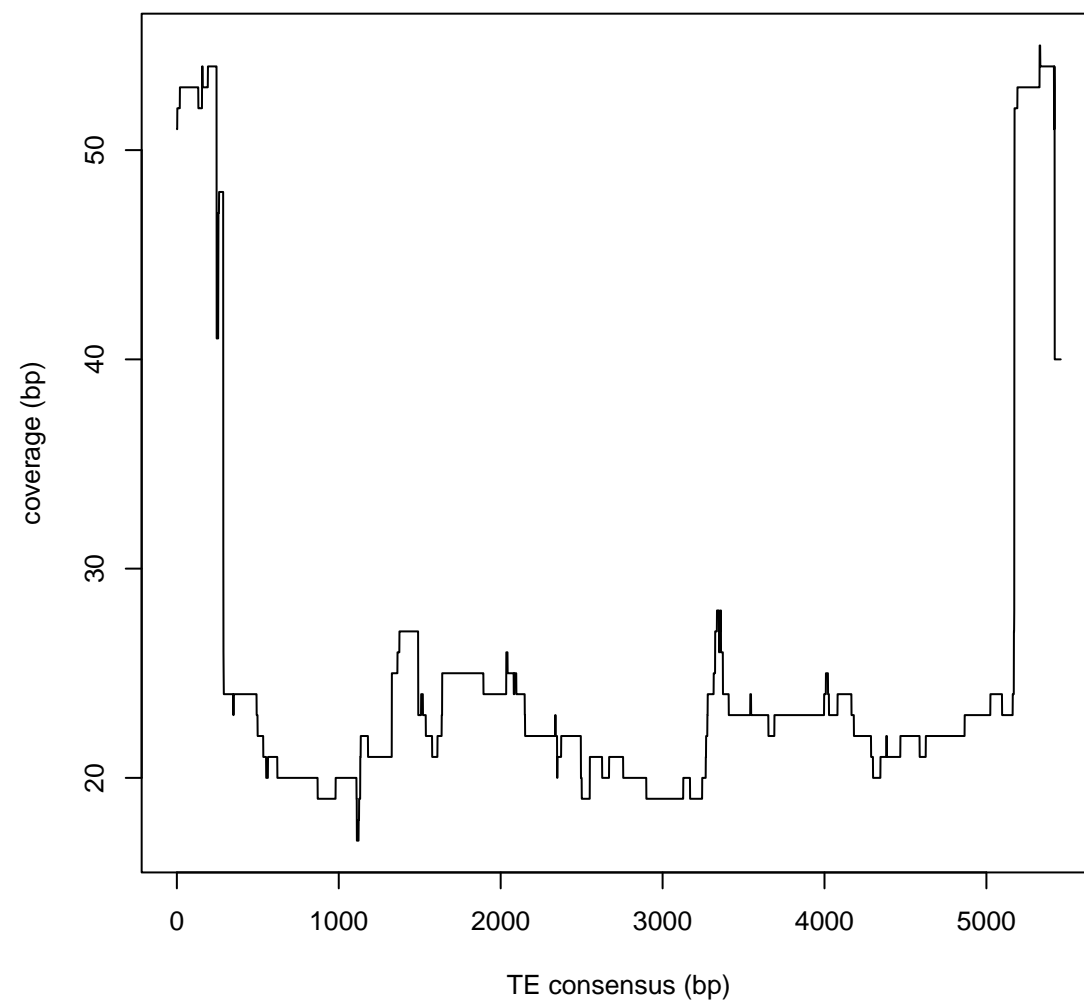


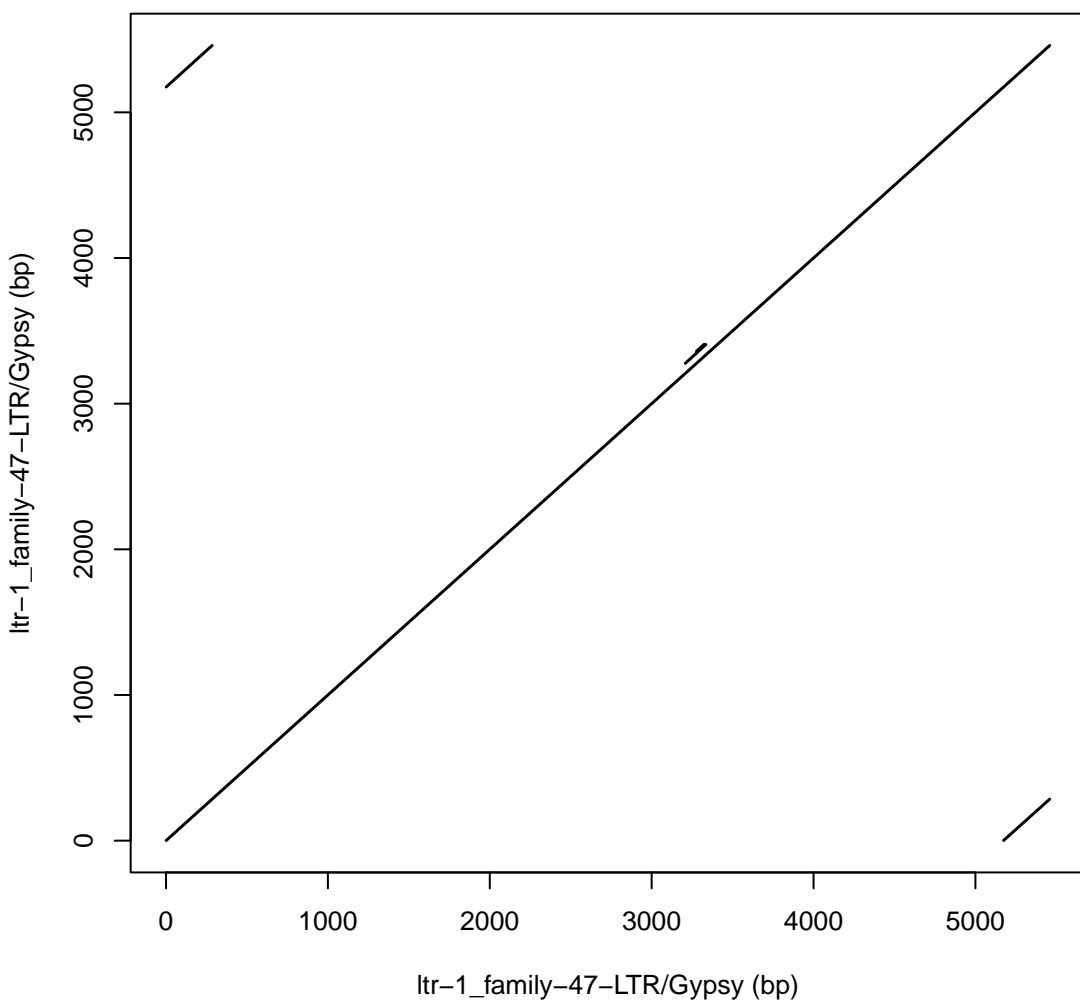
TE: ltr-1\_family-47-LTR/Gypsy  
consensus size: 5459bp; fragments: 149; full length: 8 (>=4913.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

