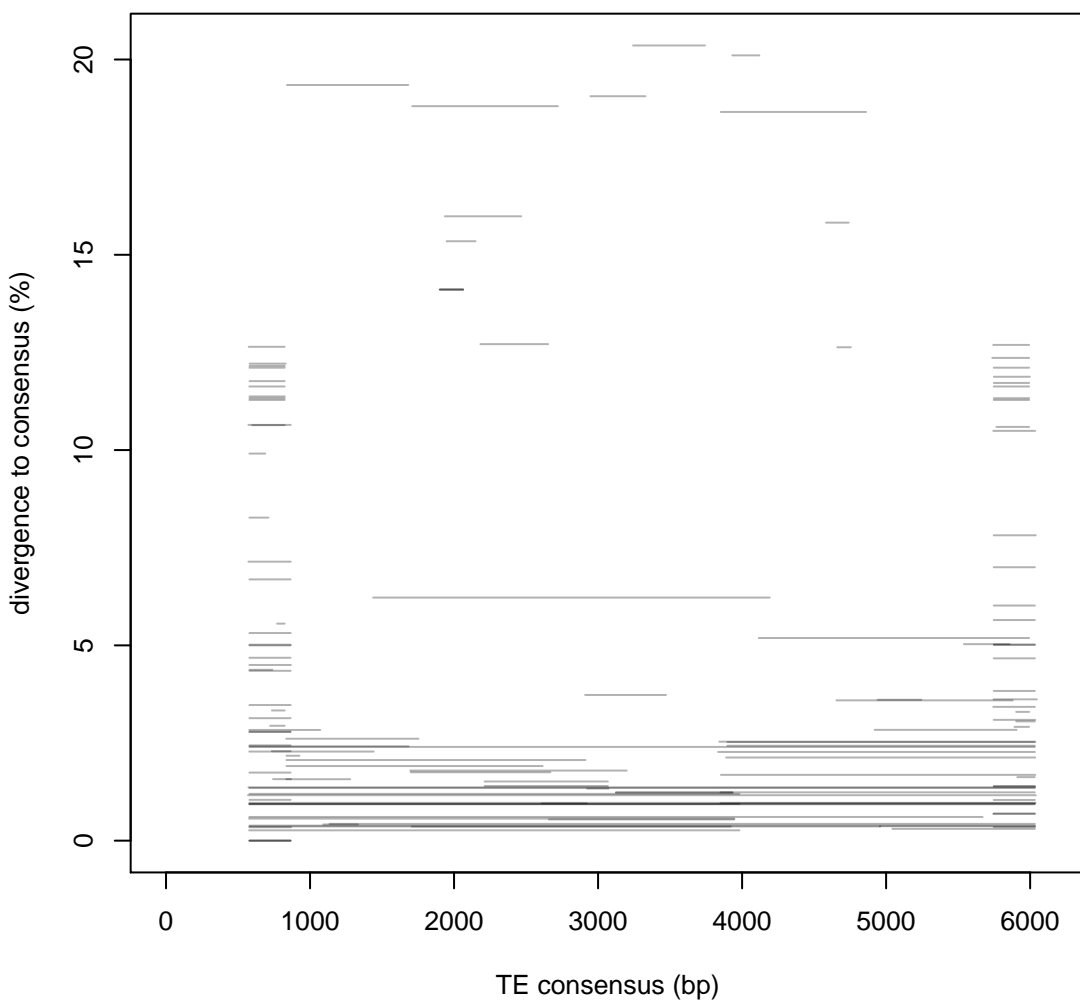
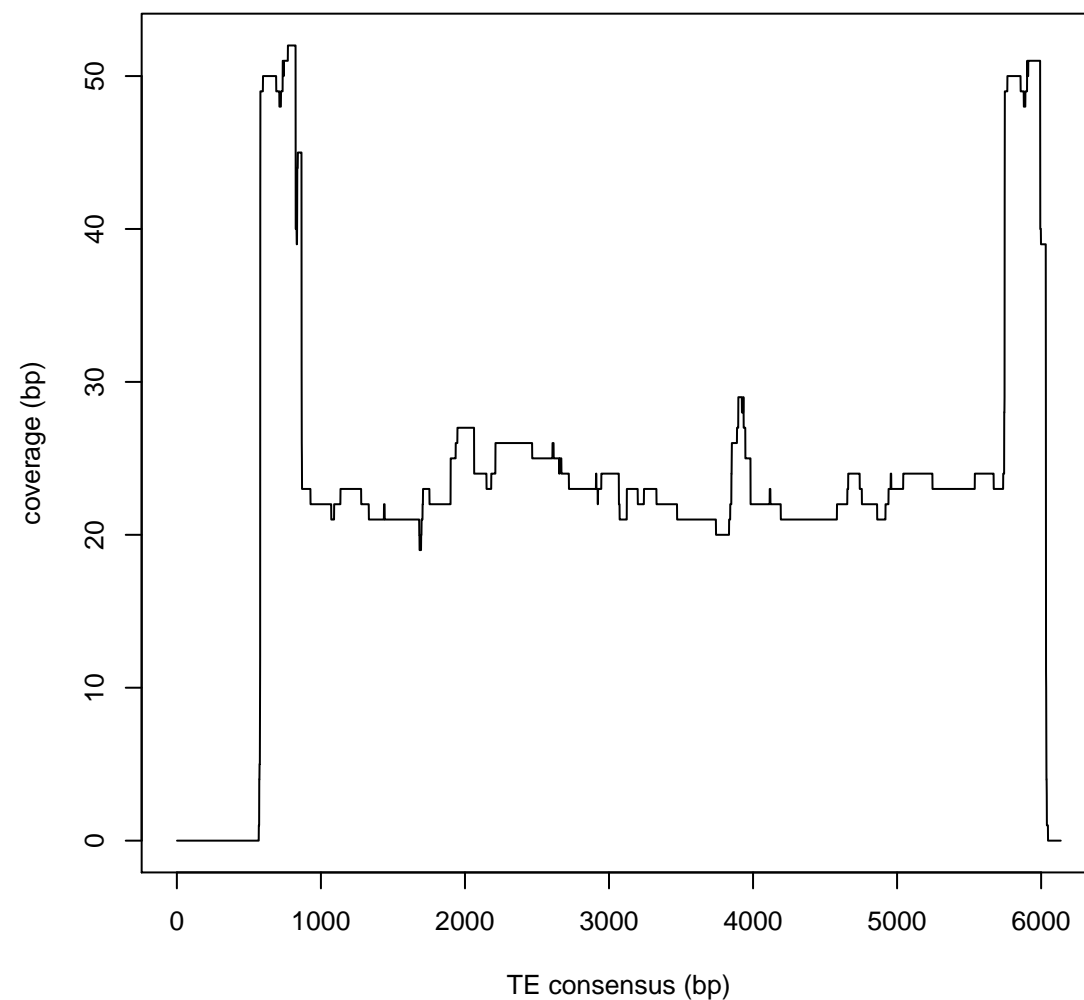


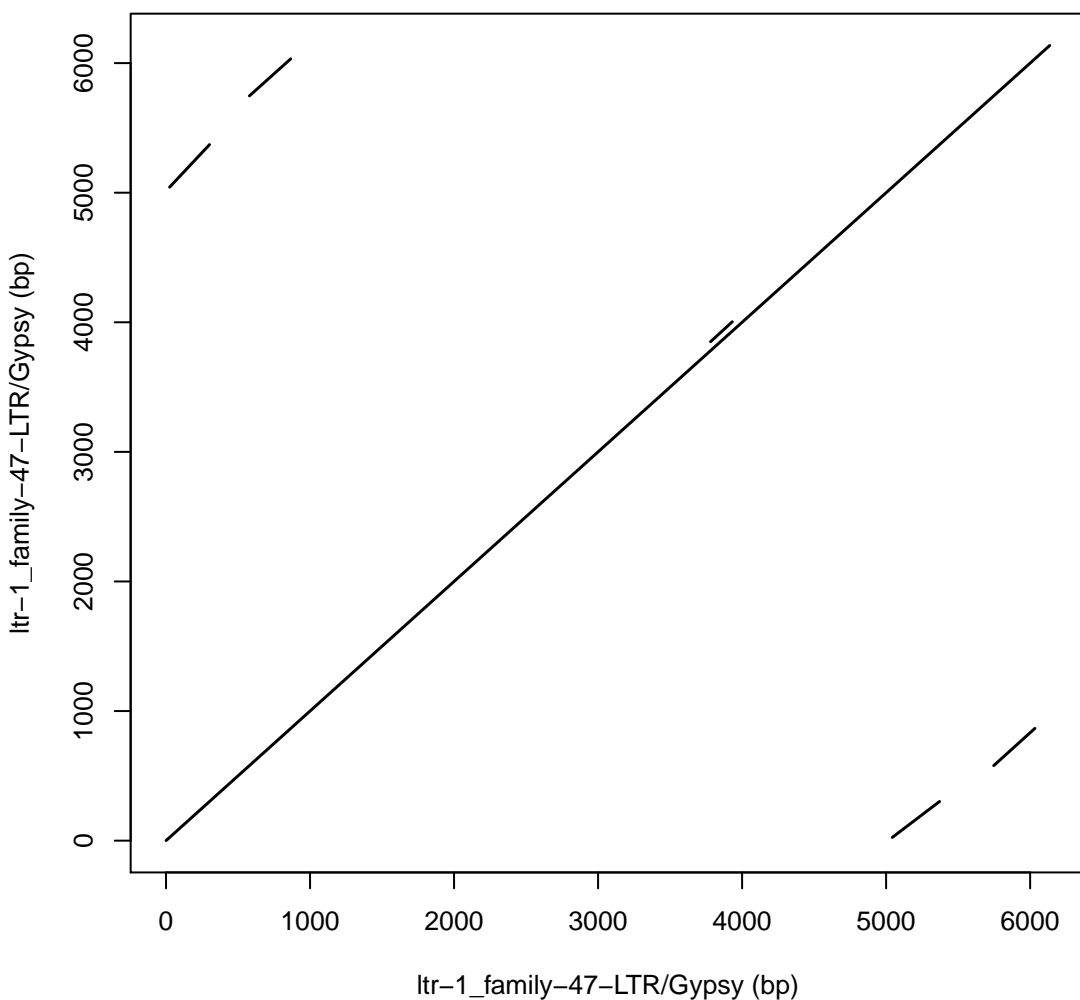
TE: ltr-1\_family-47-LTR/Gypsy  
consensus size: 6136bp; fragments: 134; full length: 0 (>=5522.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

