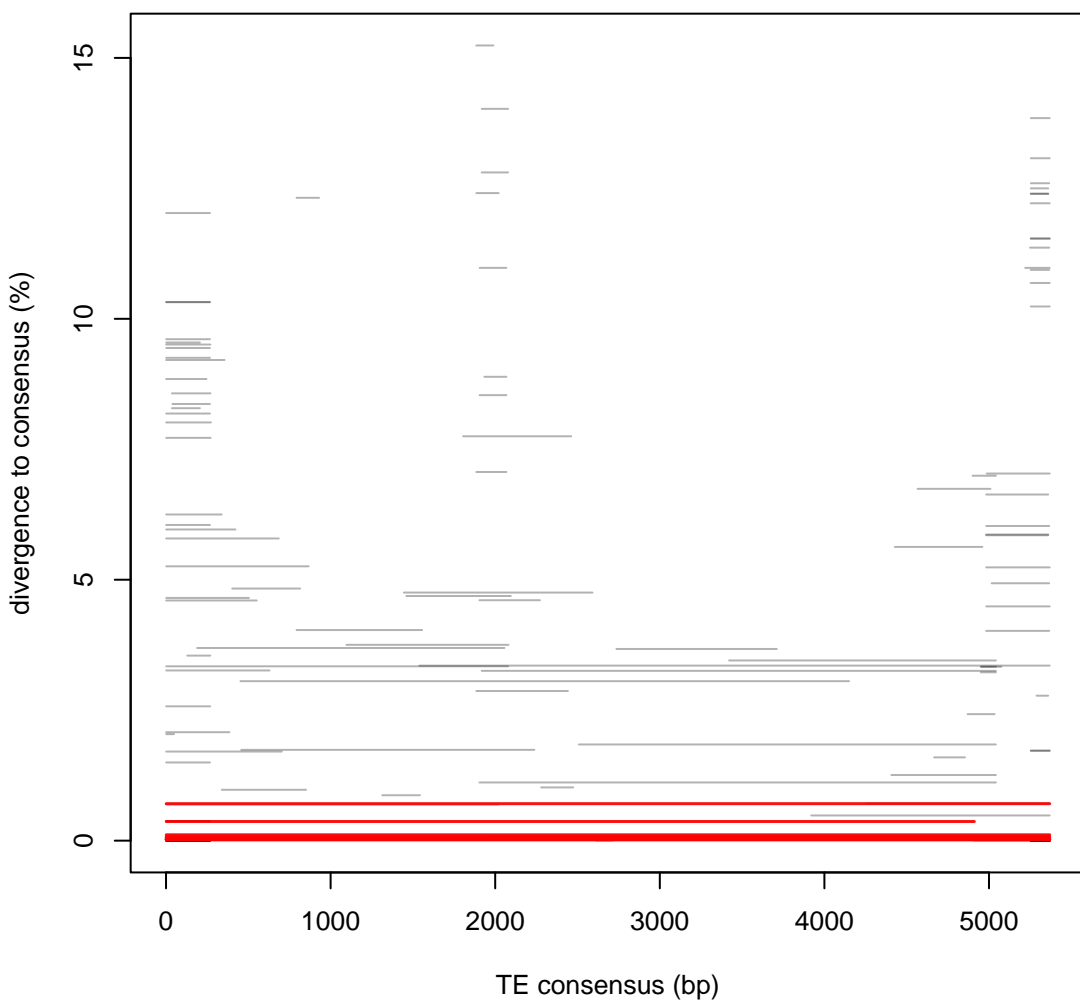
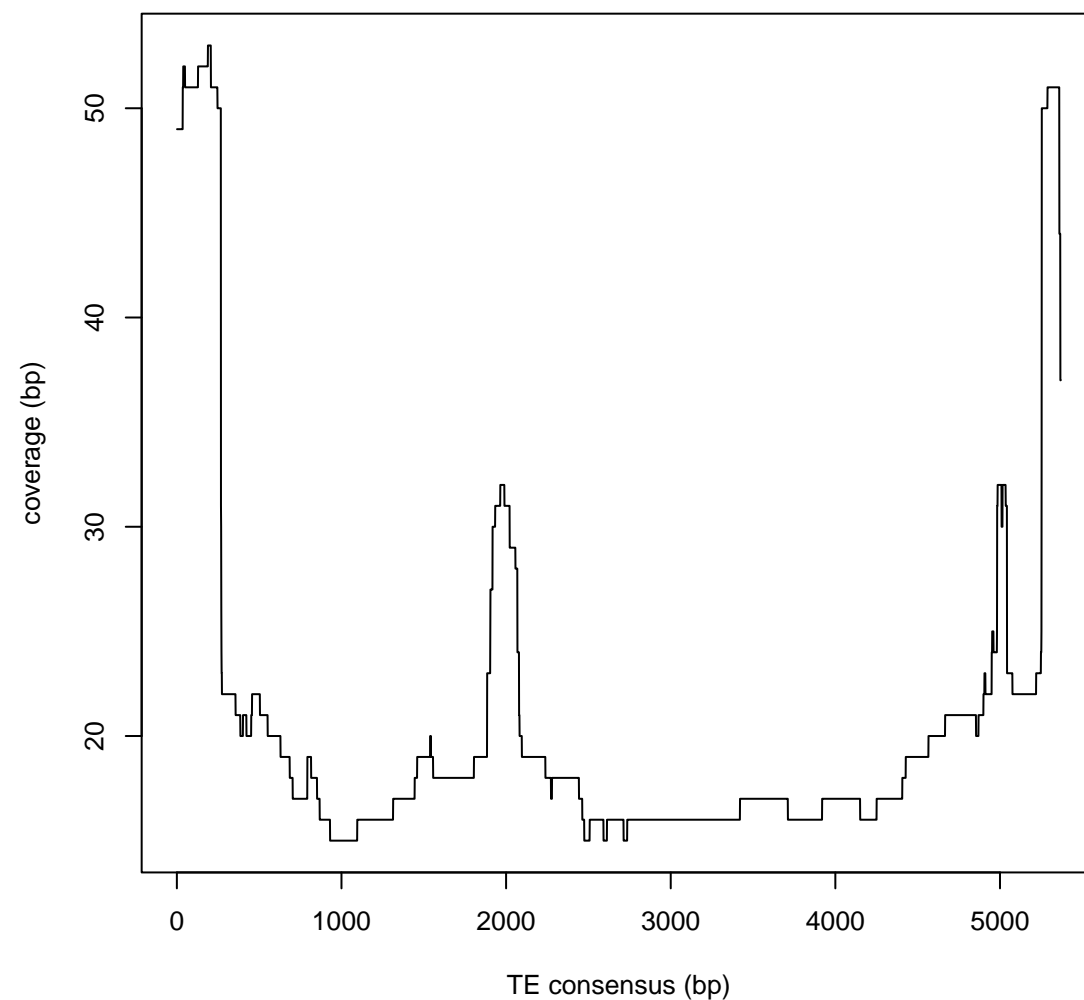


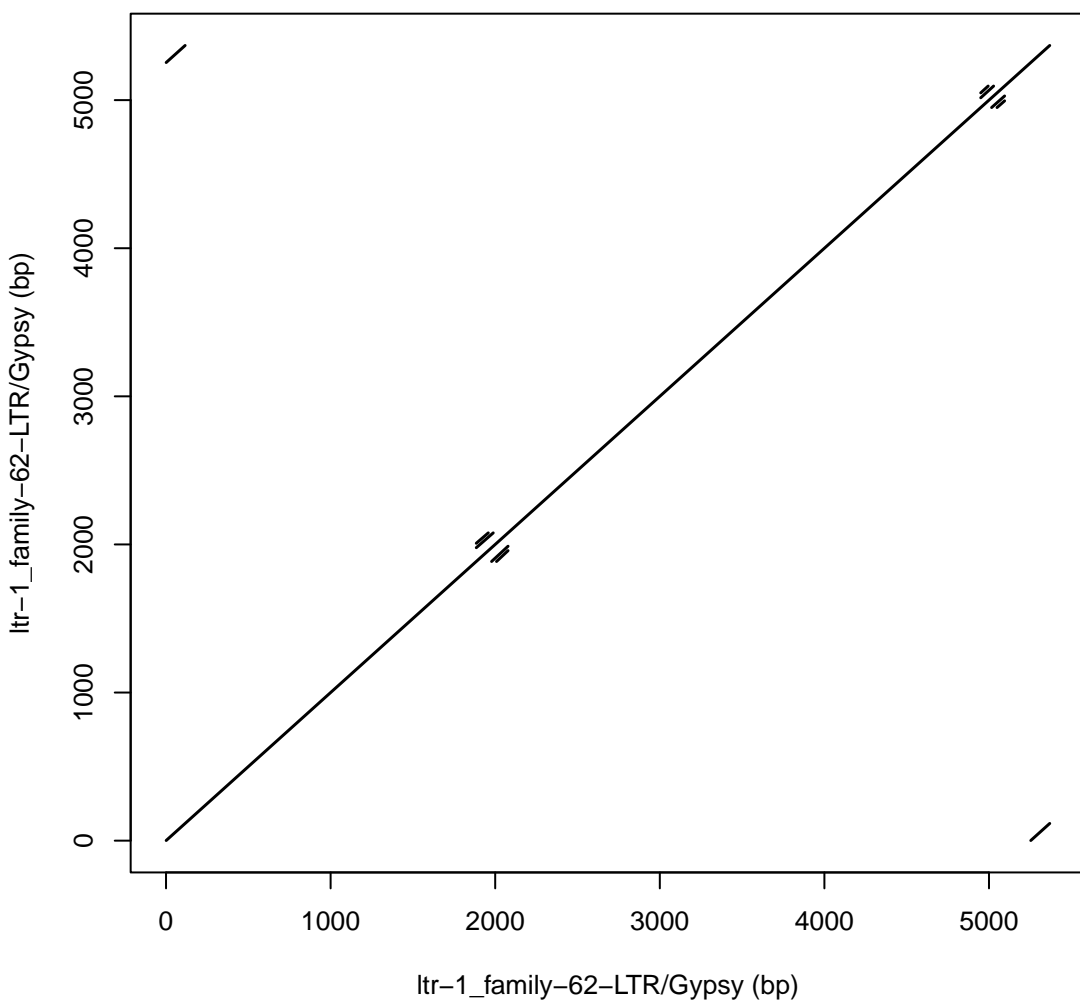
TE: ltr-1\_family-62-LTR/Gypsy  
consensus size: 5369bp; fragments: 135; full length: 9 (>=4832.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

