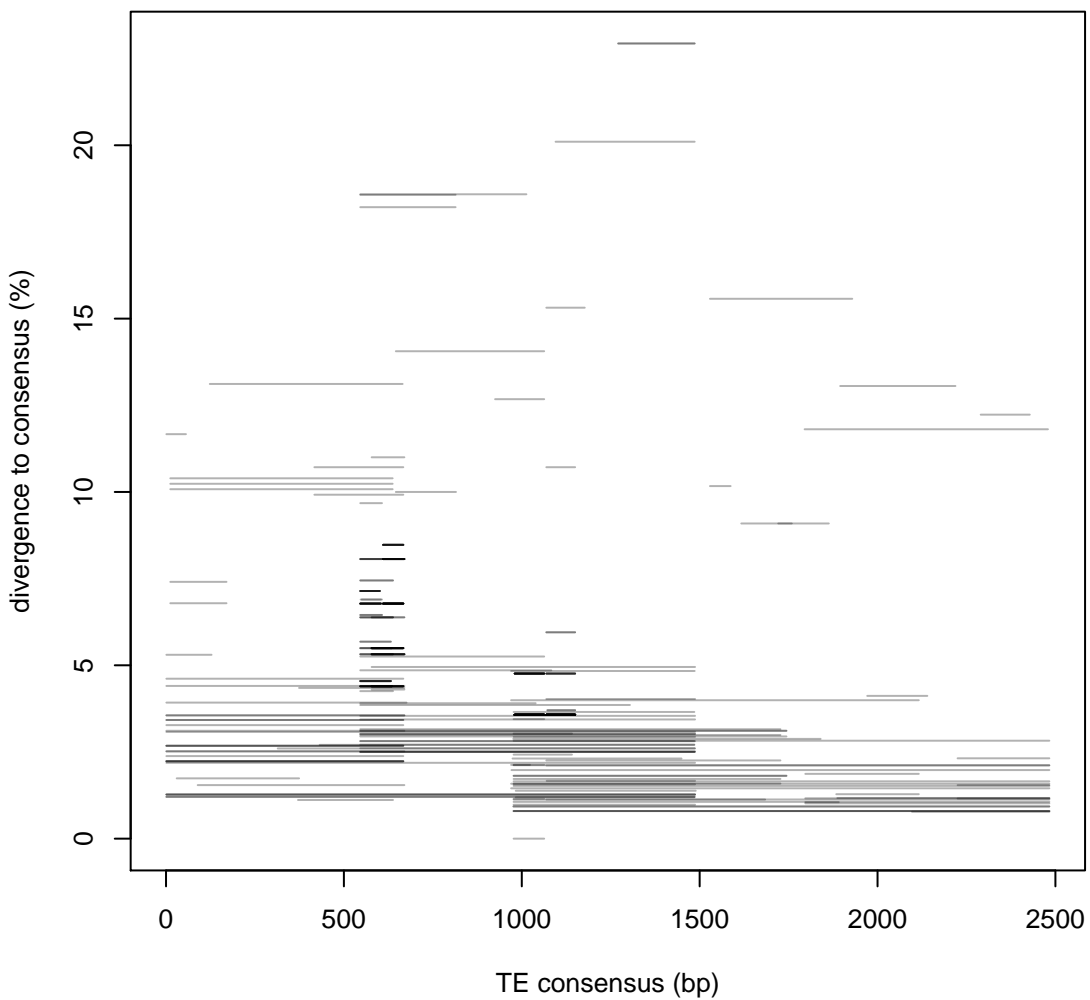
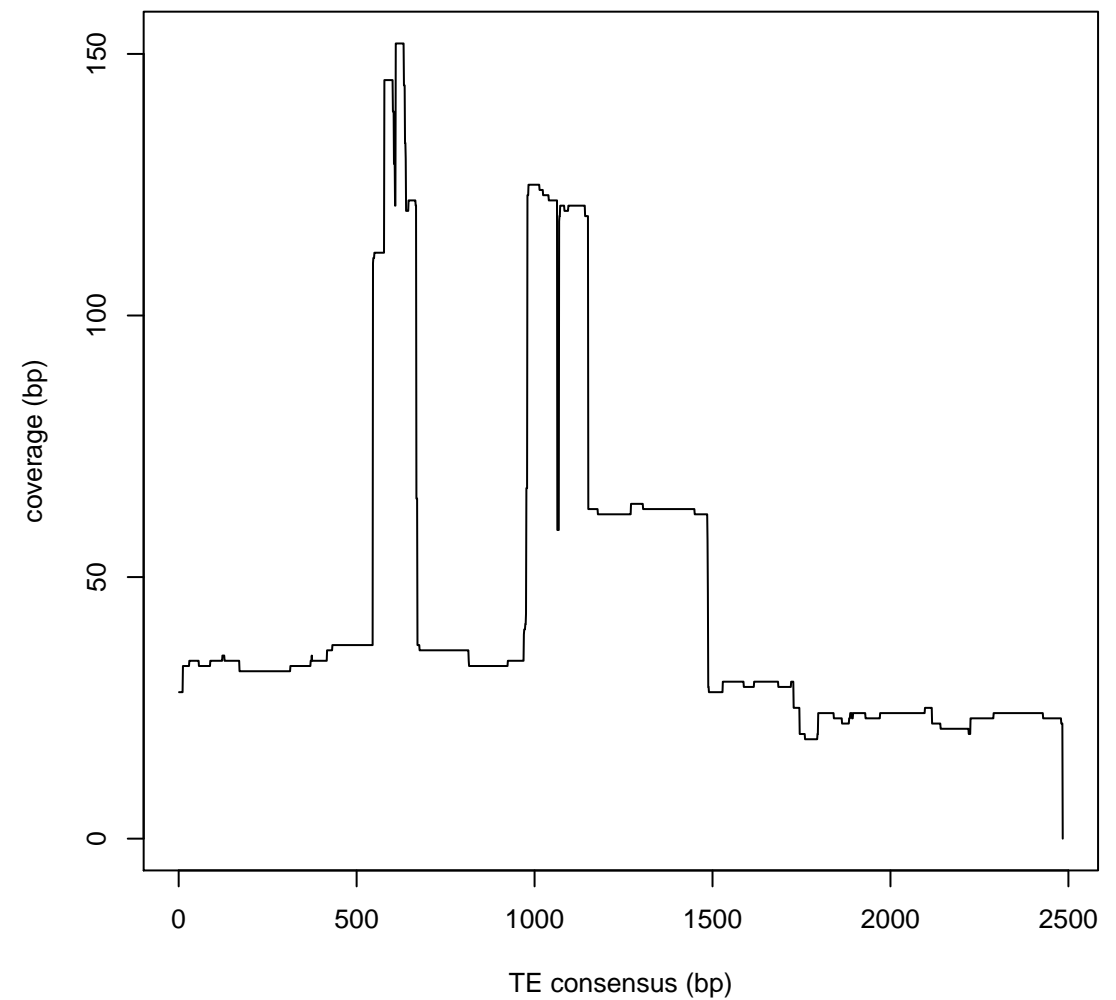


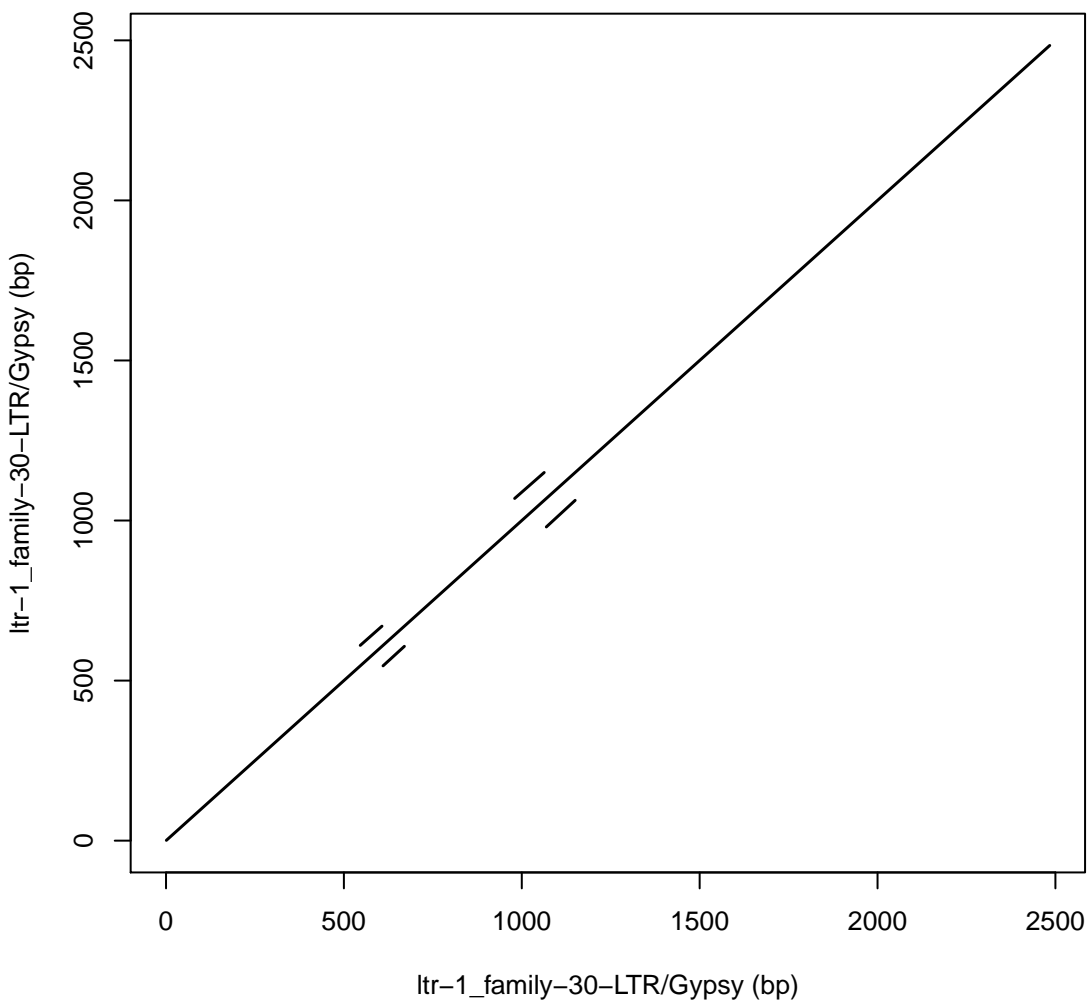
TE: ltr-1\_family-30-LTR/Gypsy  
consensus size: 2484bp; fragments: 358; full length: 0 (>=2235.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

