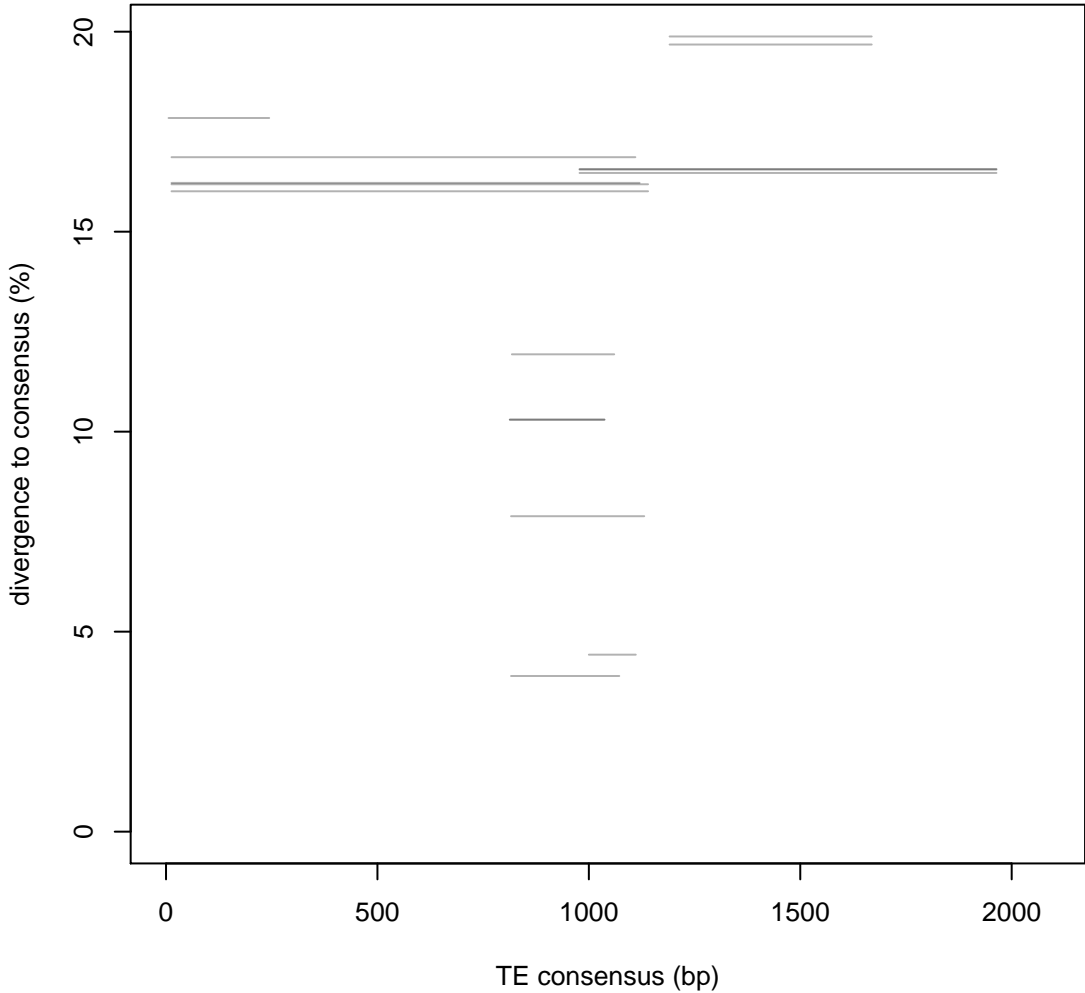
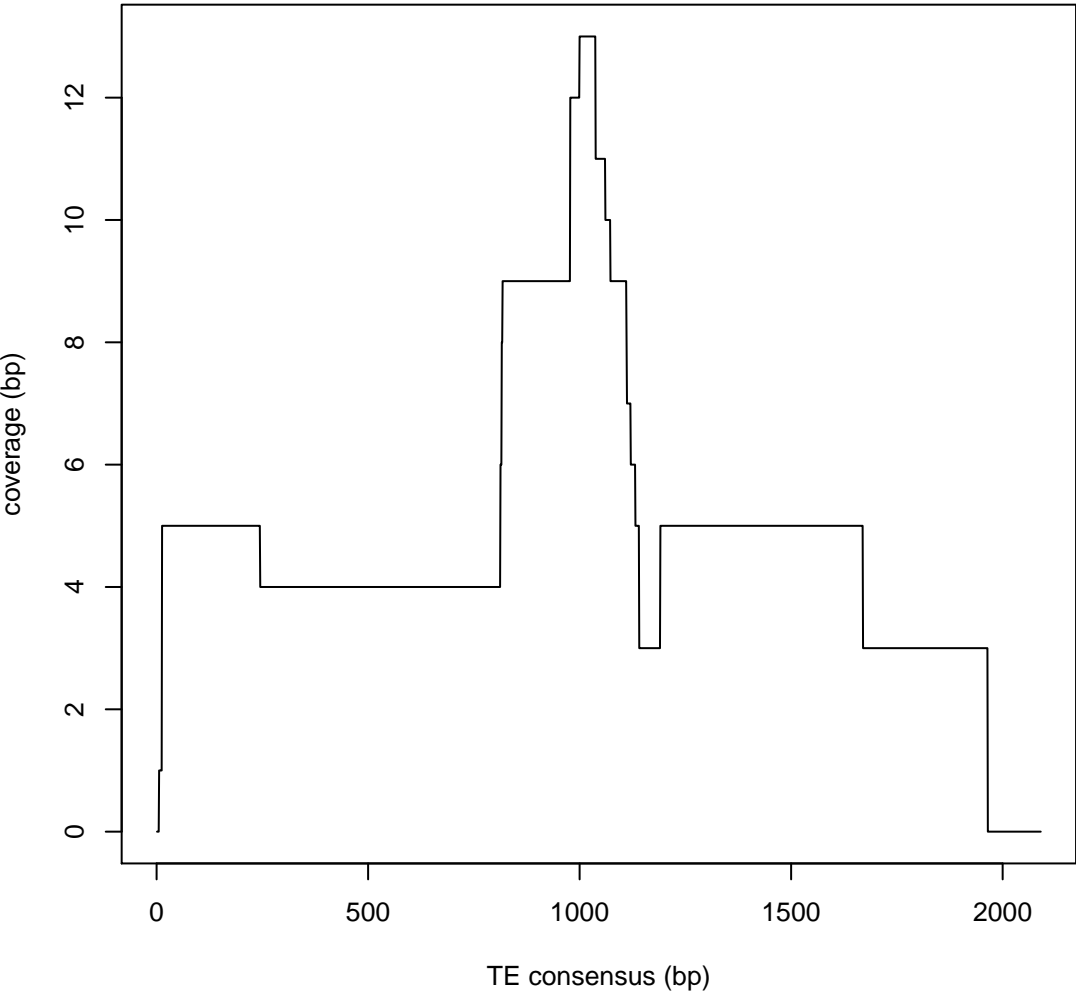


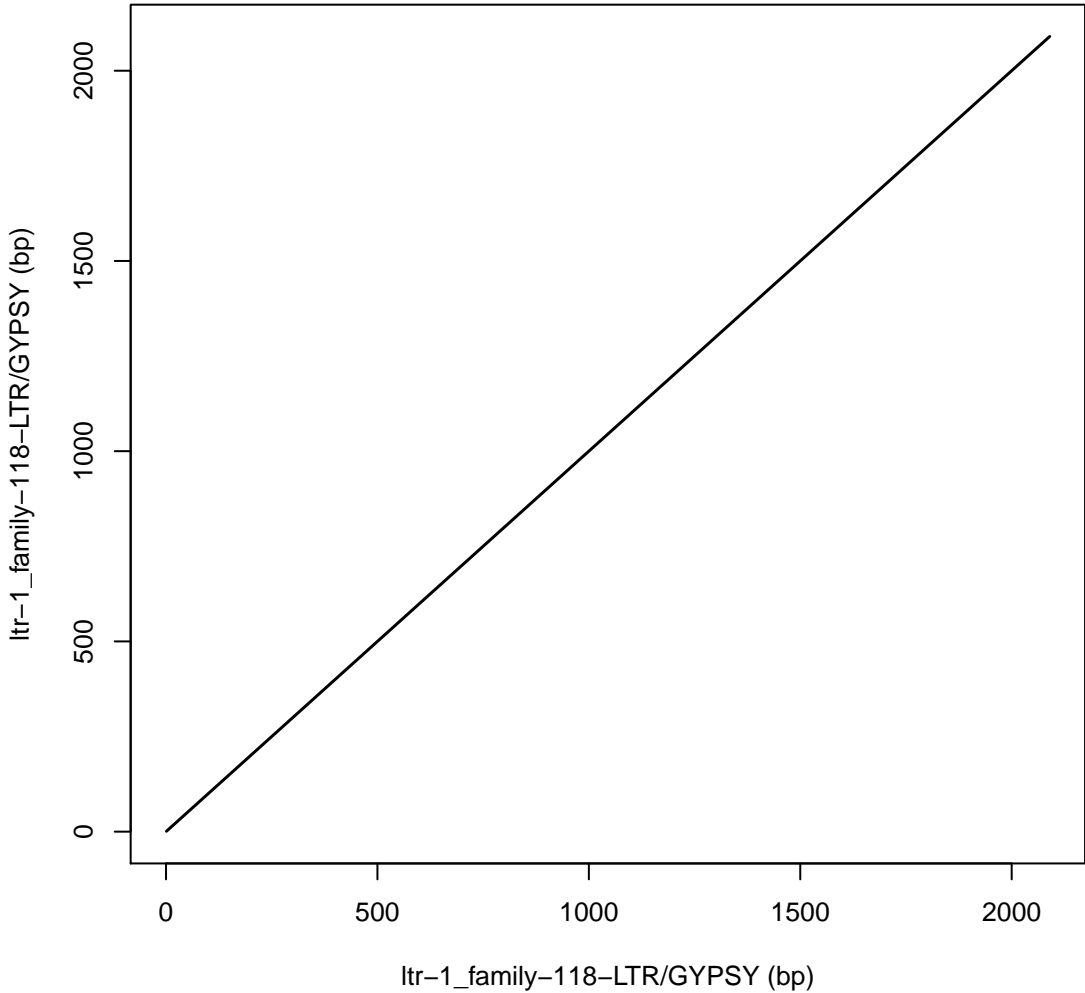
TE: ltr-1\_family-118-LTR/GYPSY  
consensus size: 2090bp; fragments: 16; full length: 0 (>=1881bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

