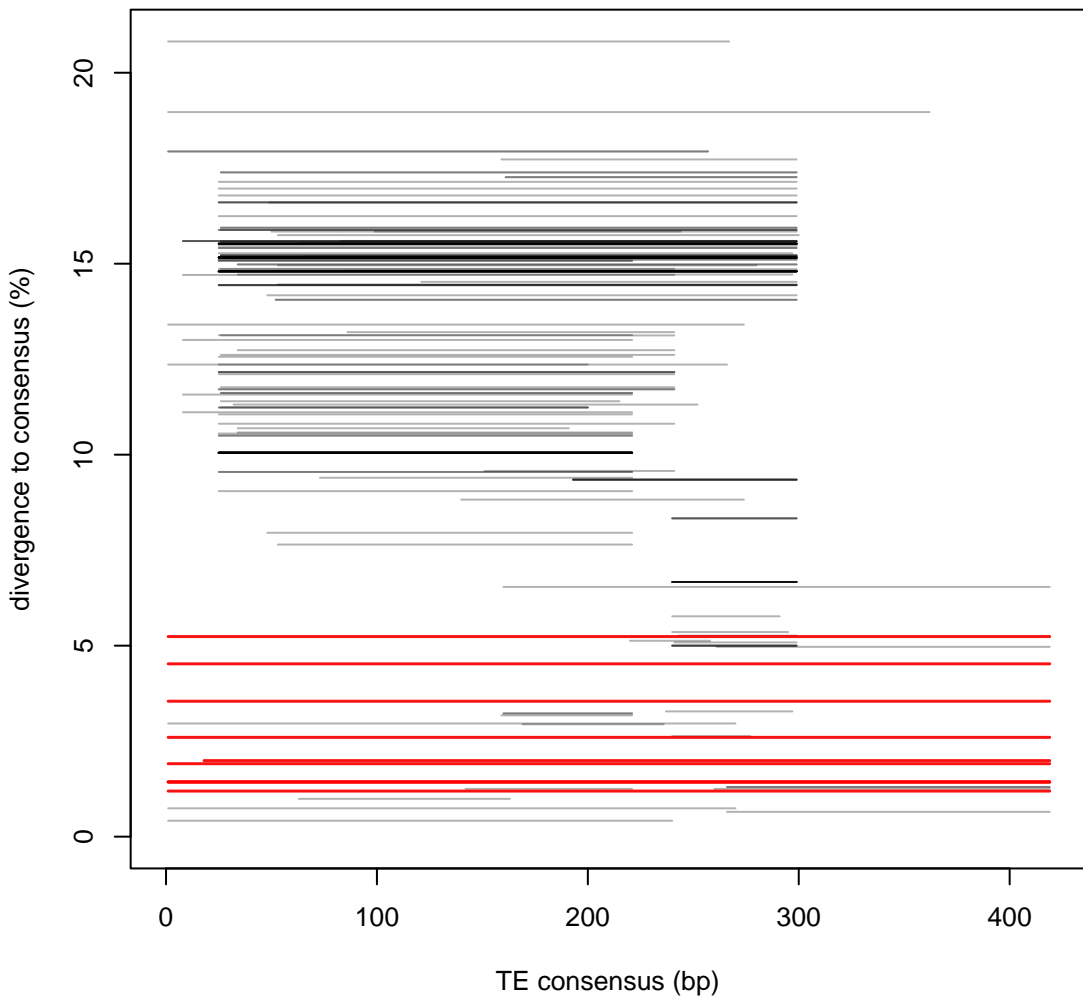
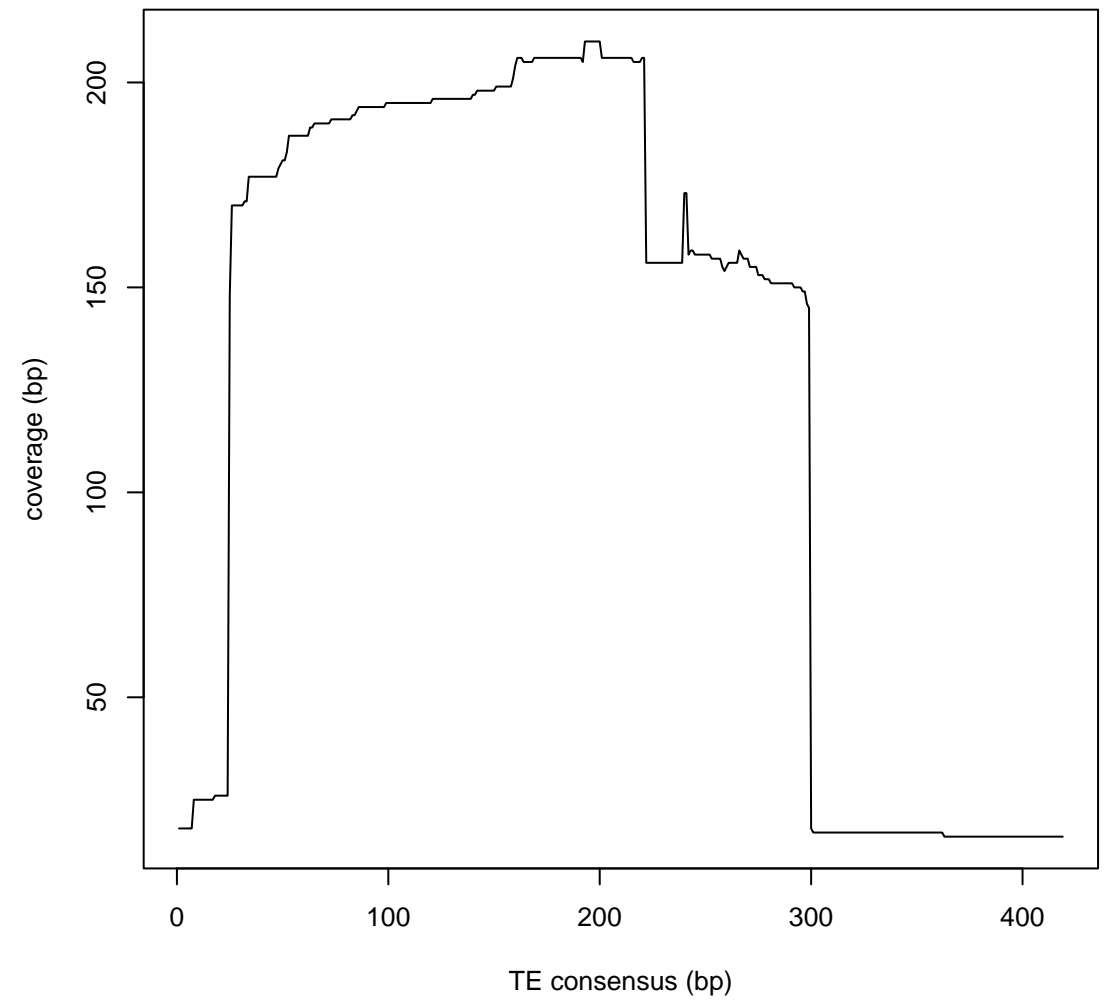


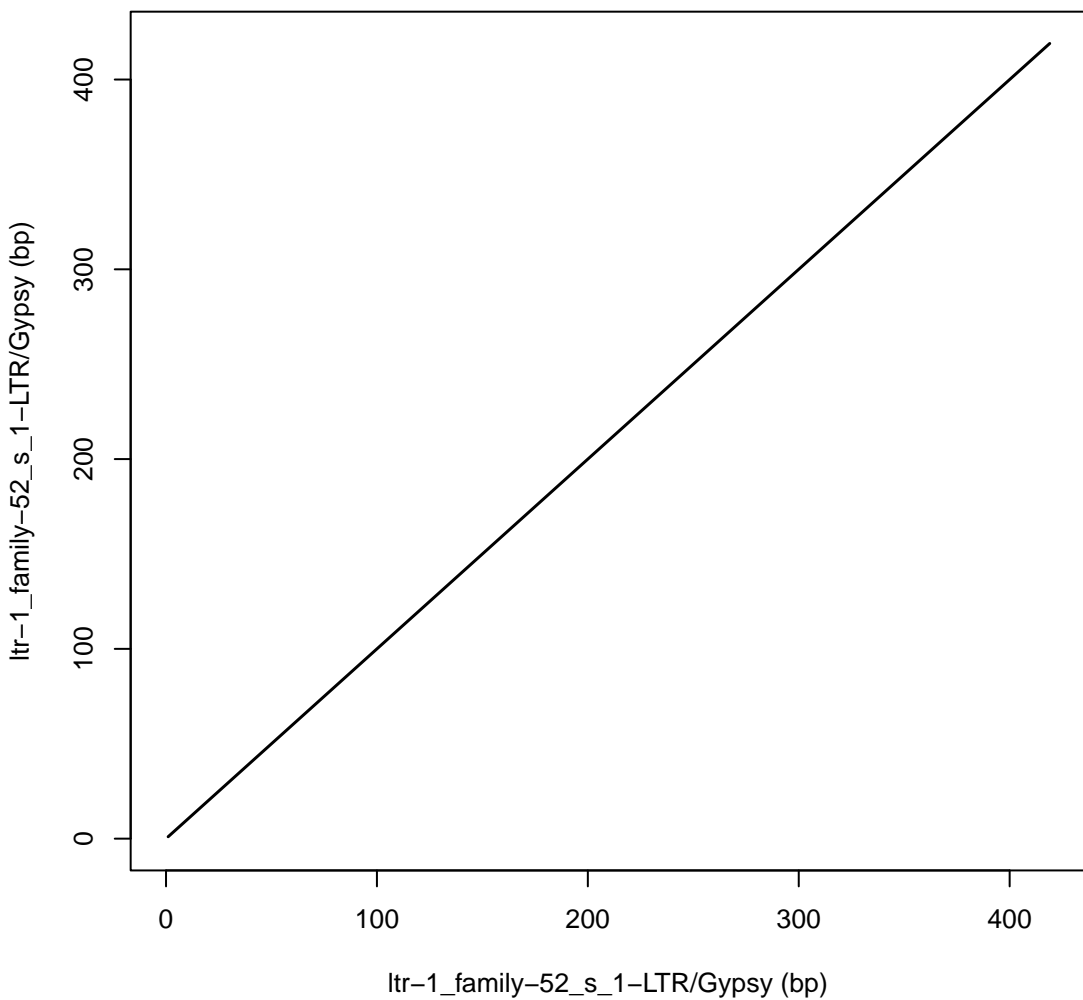
TE: ltr-1\_family-52\_s\_1-LTR/Gypsy  
consensus size: 419bp; fragments: 238; full length: 10 (>=377.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

