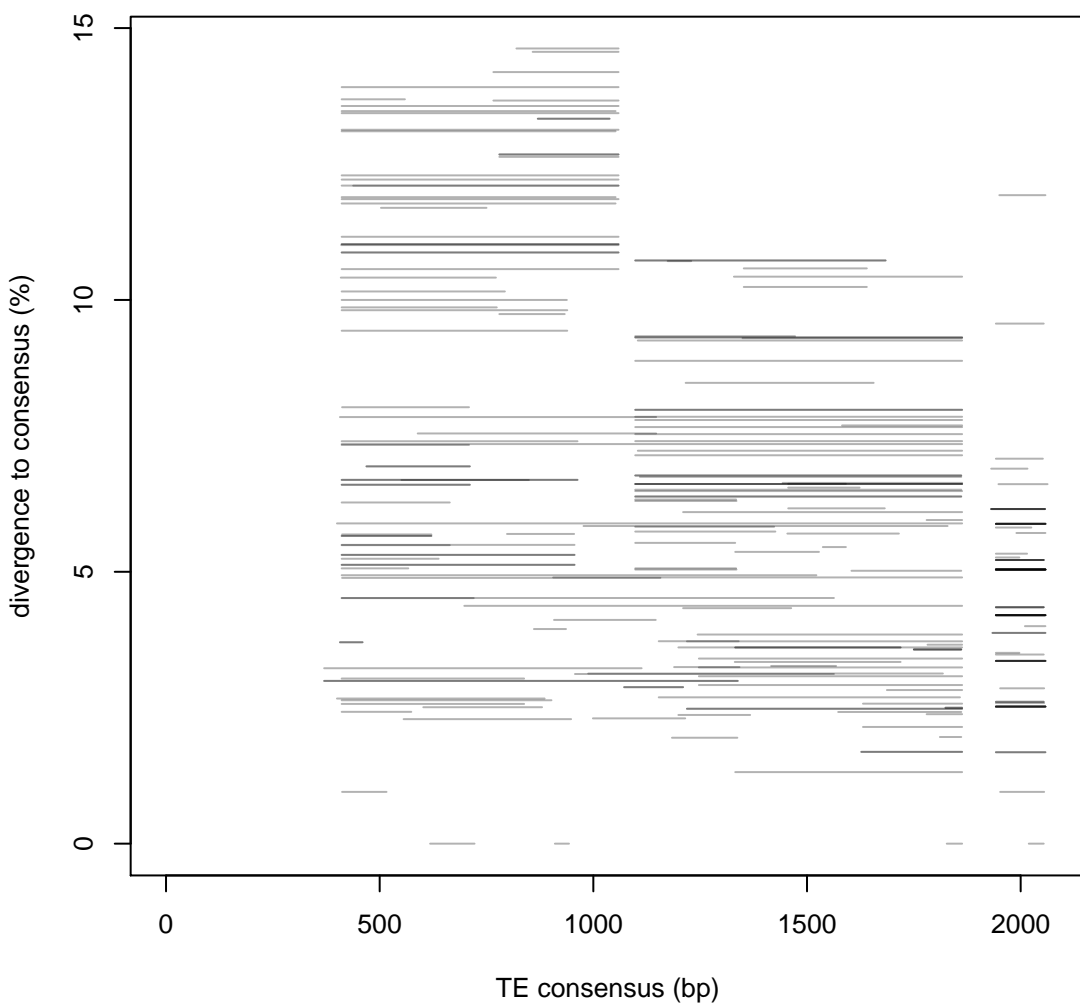
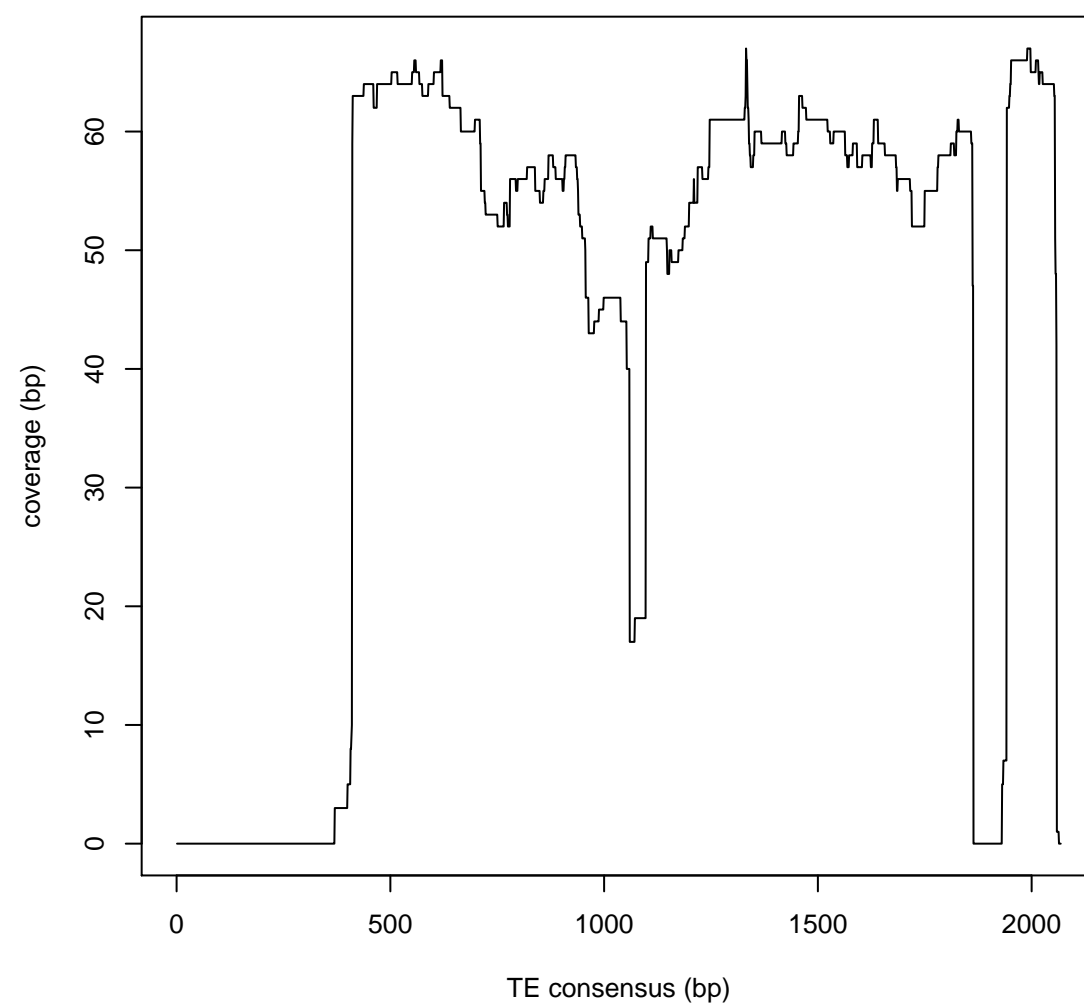


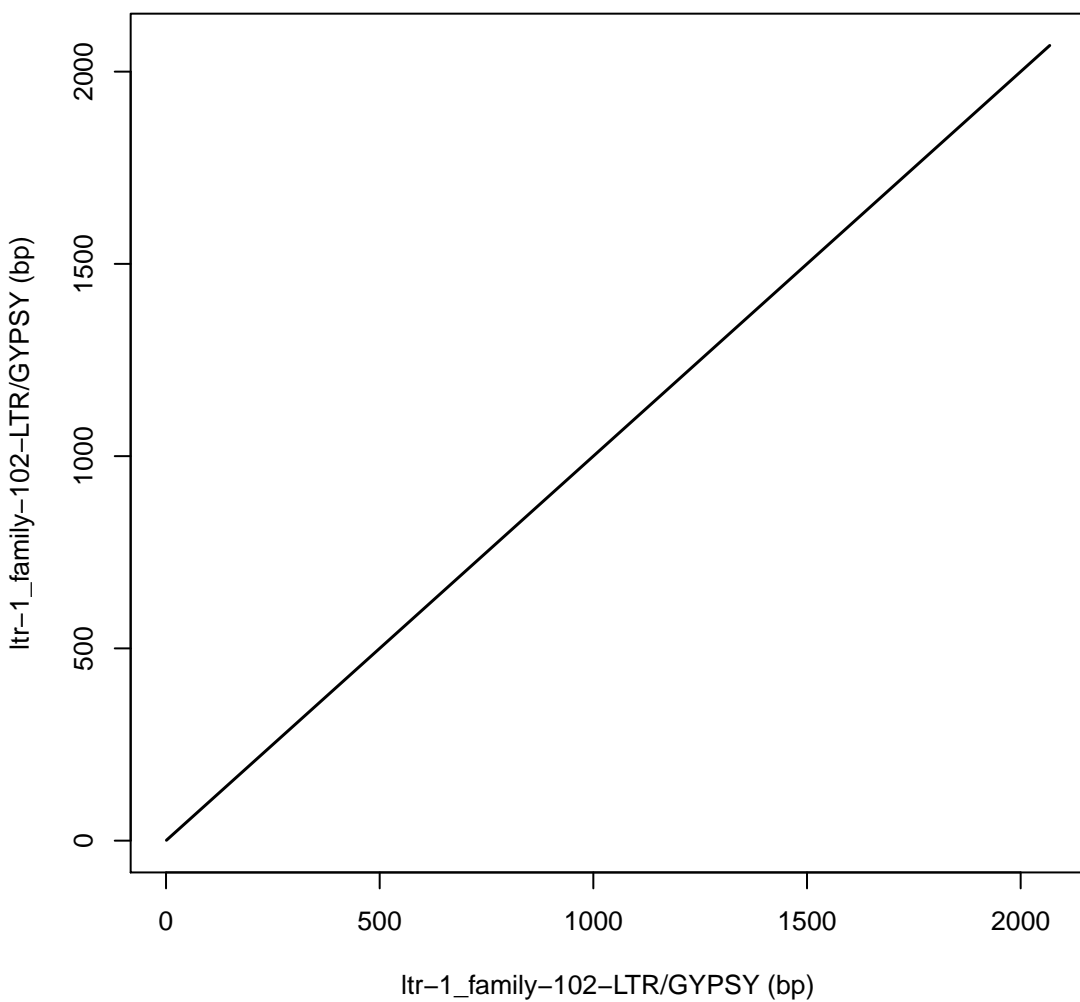
TE: ltr-1\_family-102-LTR/Gypsy  
consensus size: 2068bp; fragments: 248; full length: 0 (>=1861.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

