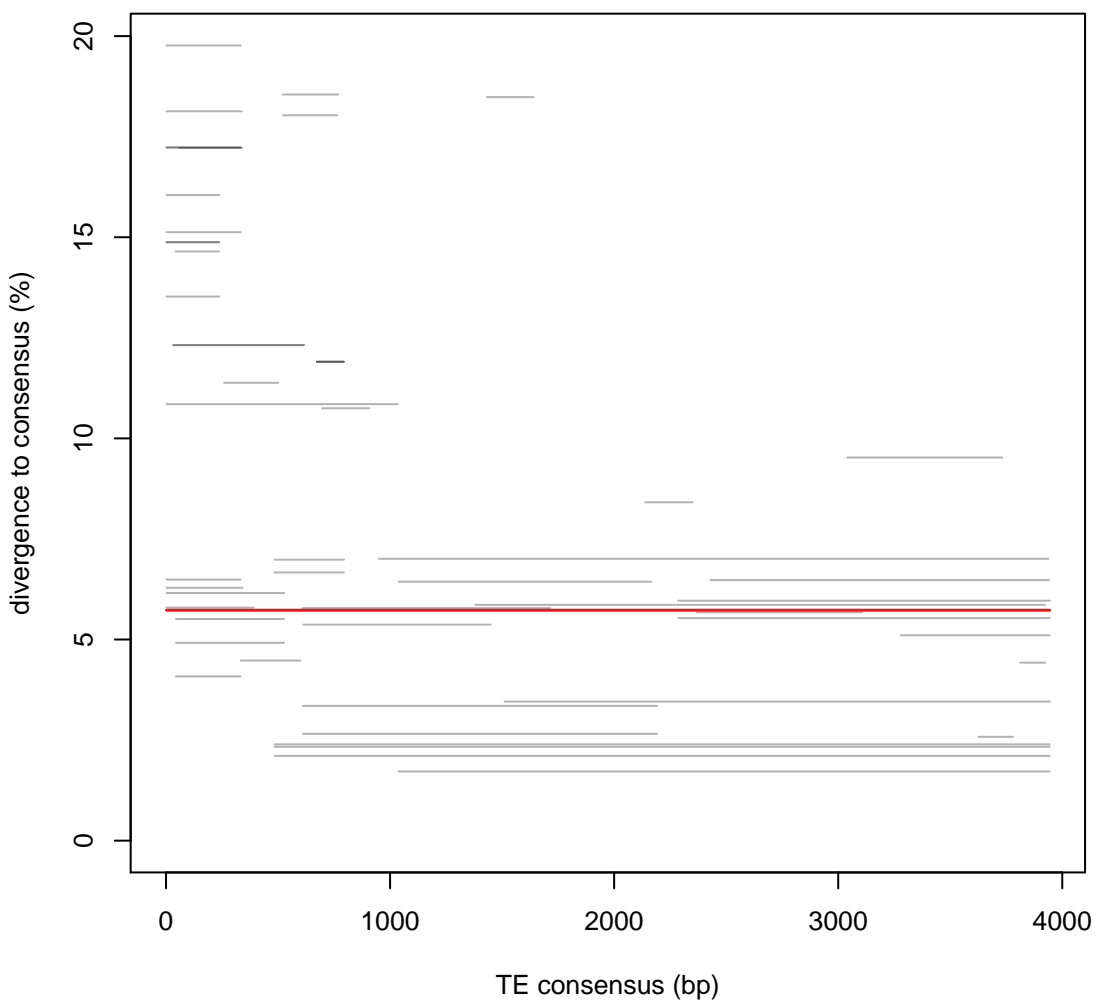
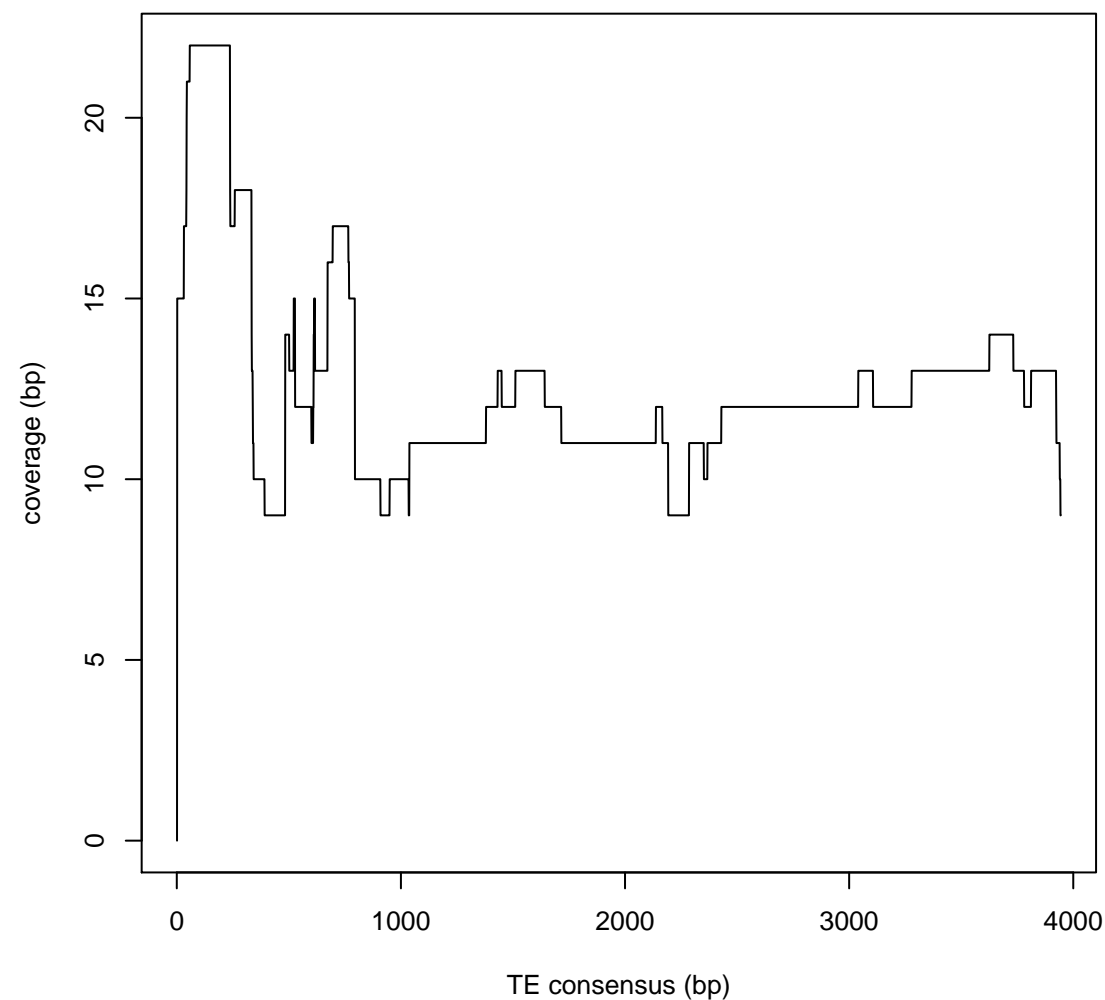


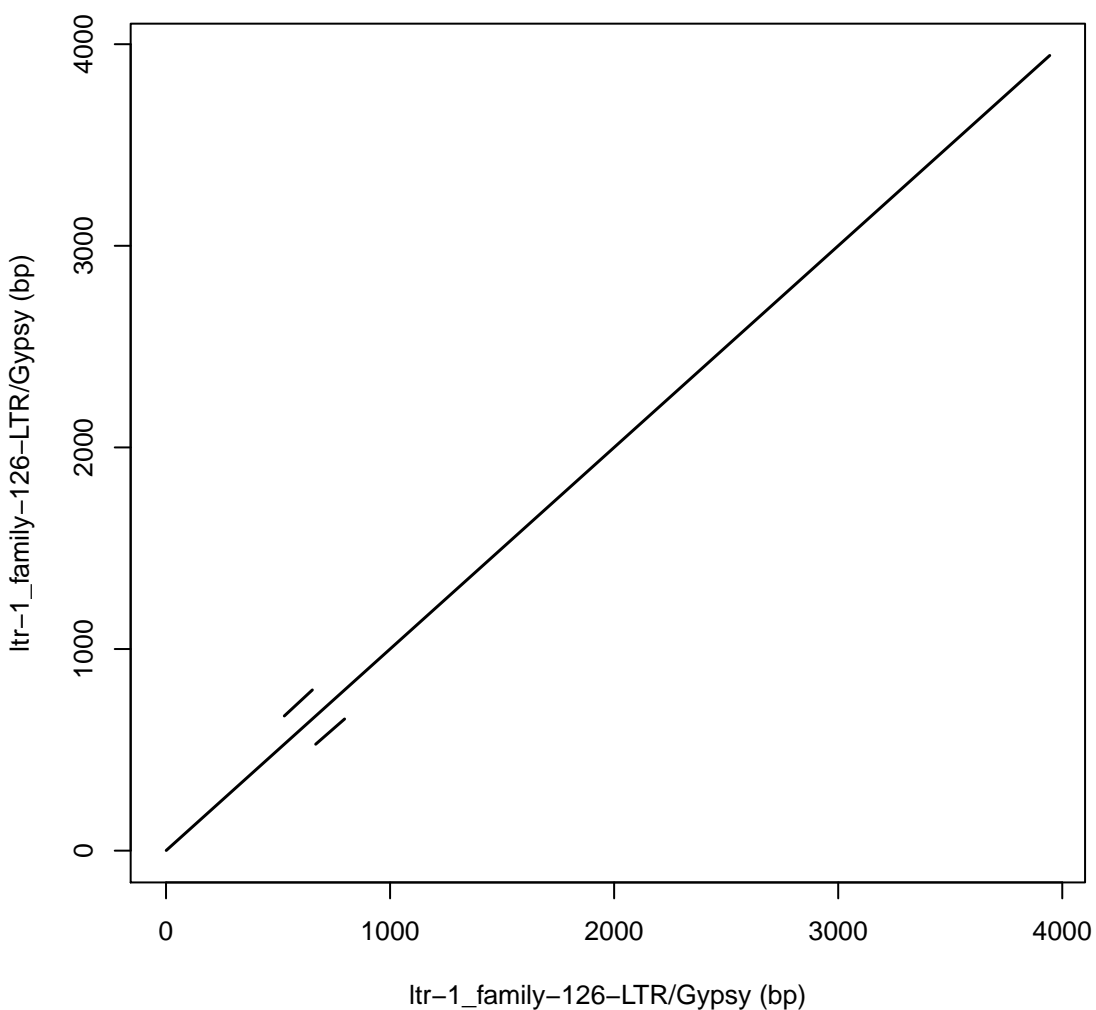
TE: ltr-1_family-126-LTR/Gypsy
consensus size: 3944bp; fragments: 54; full length: 1 (>=3549.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

