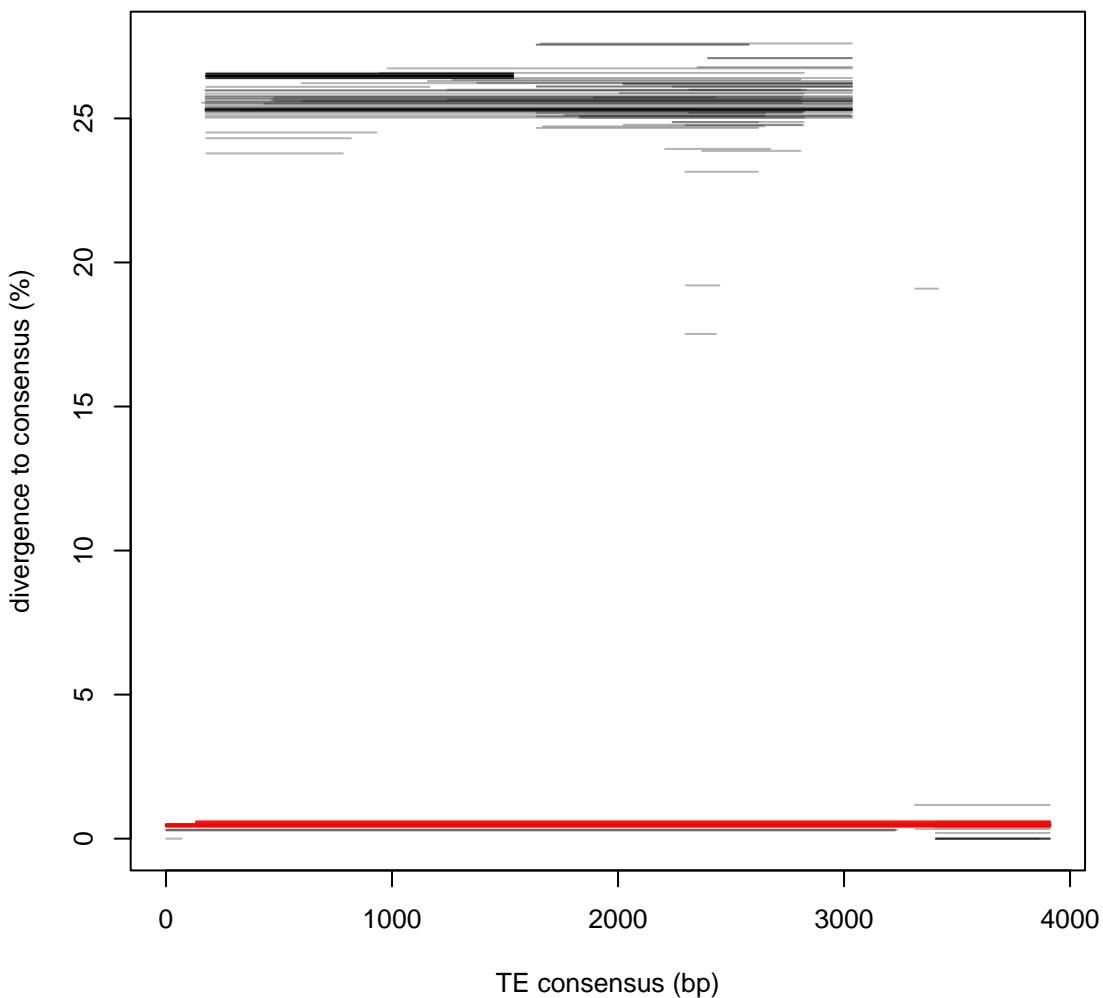
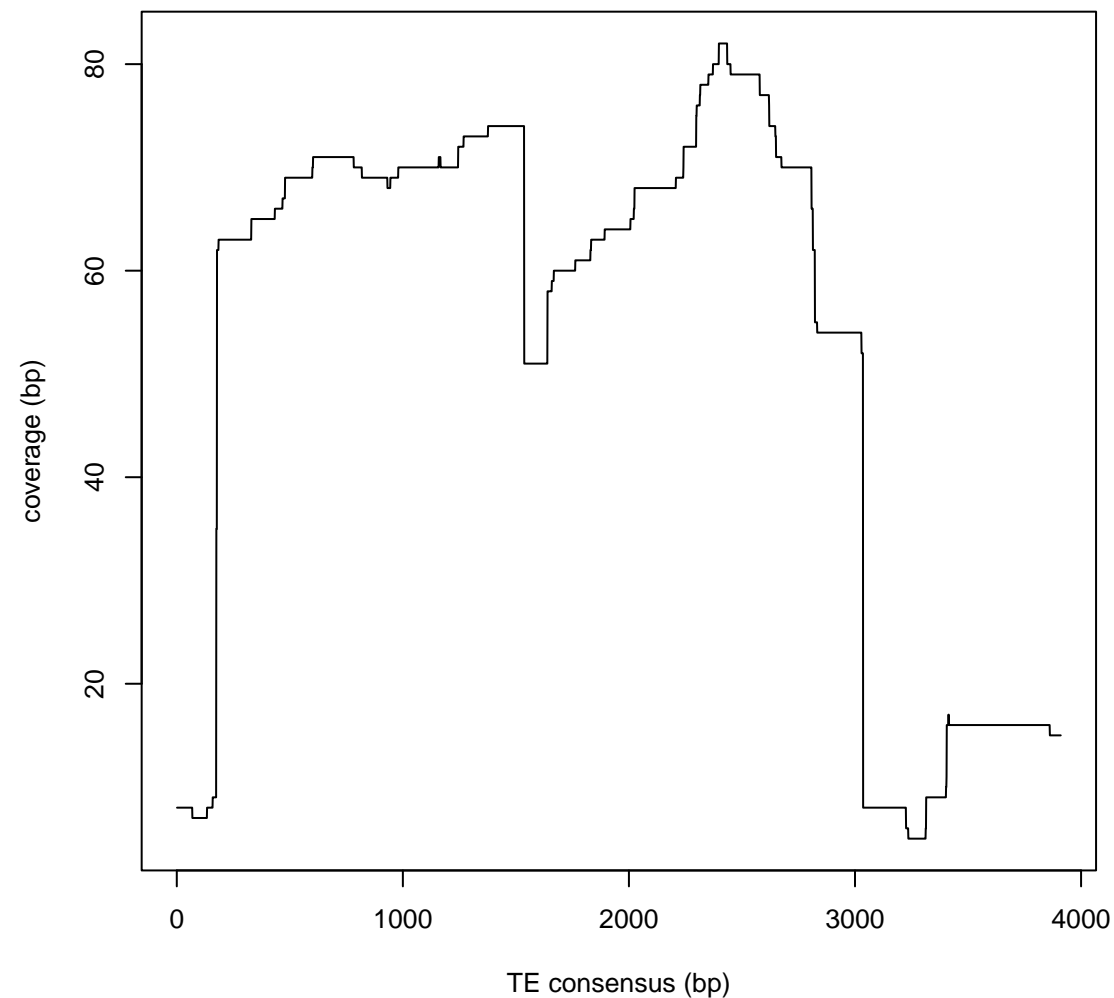


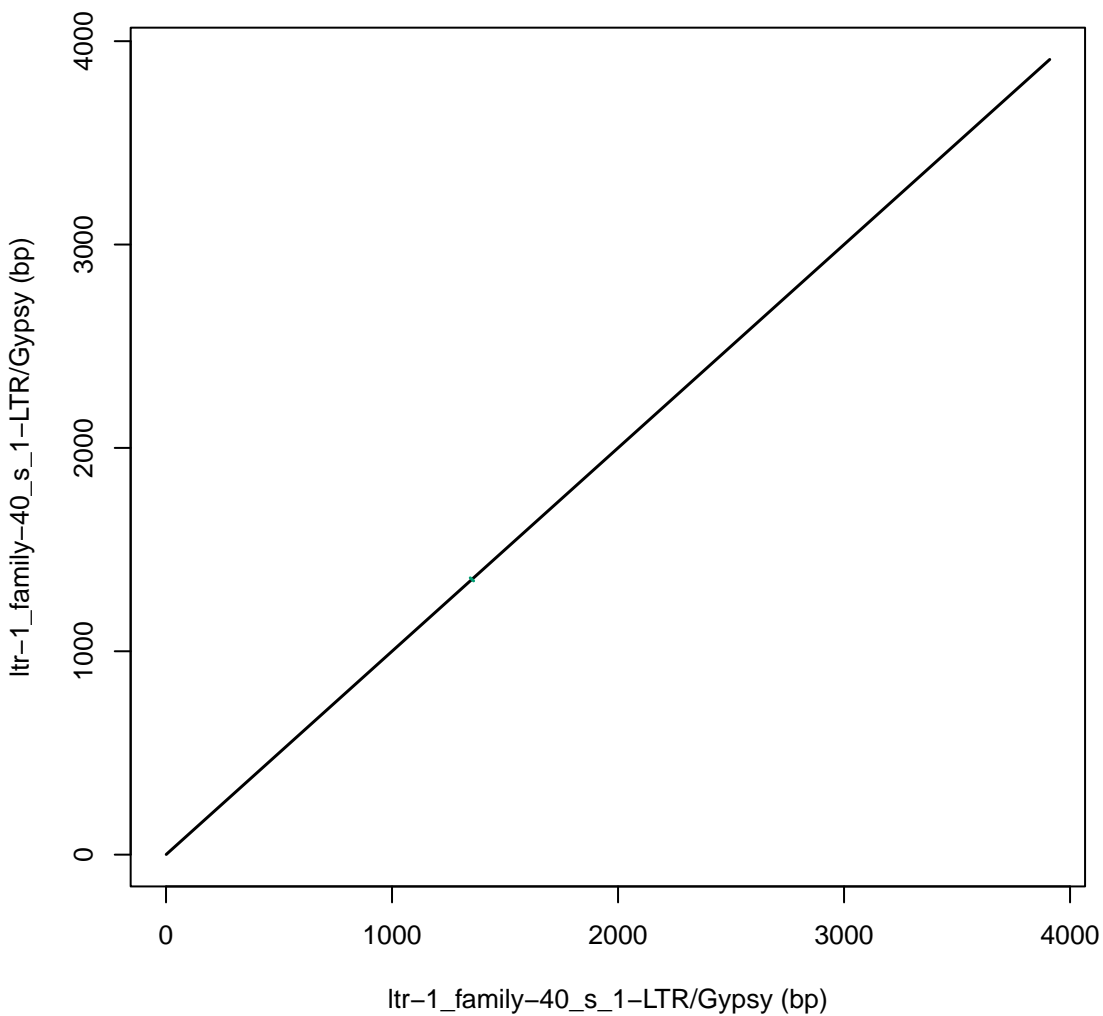
TE: ltr-1_family-40_s_1-LTR/Gypsy
consensus size: 3910bp; fragments: 122; full length: 5 (>=3519bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

