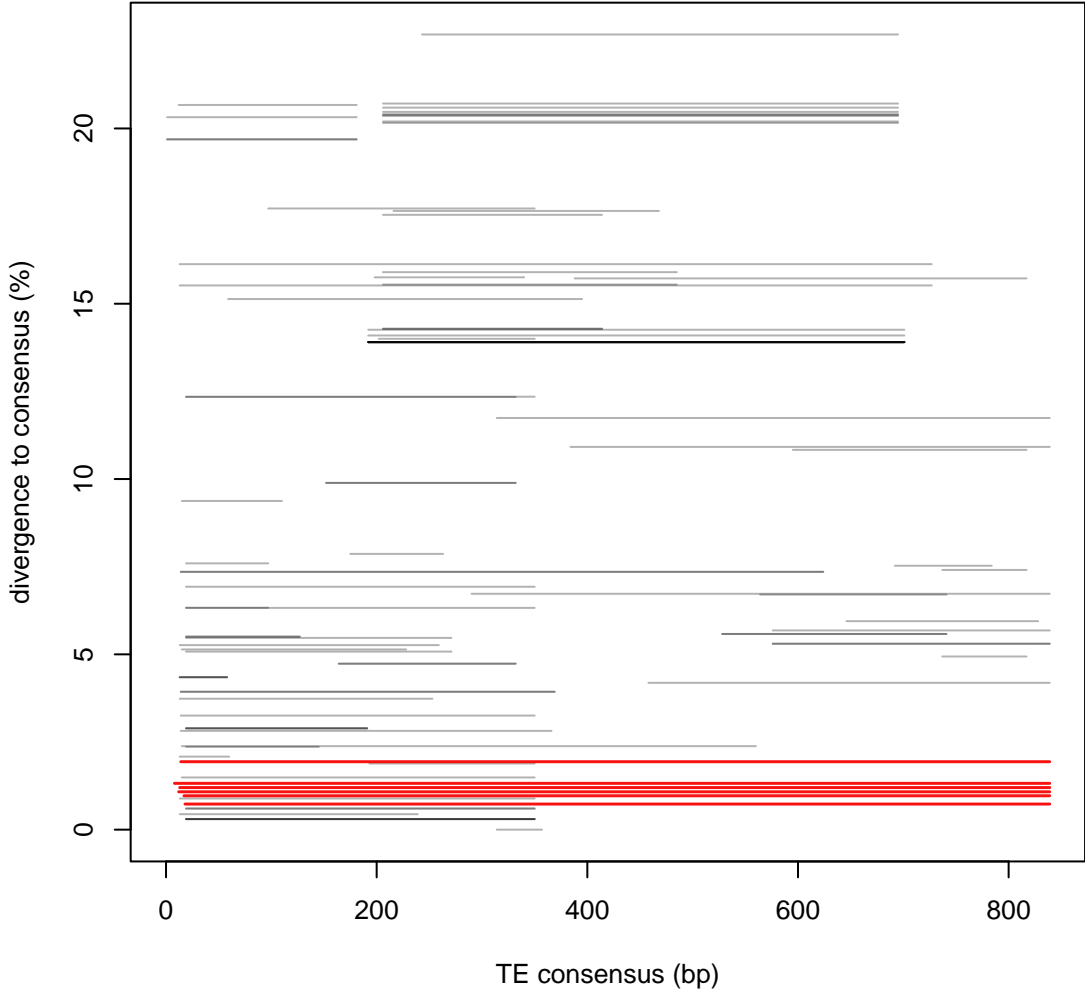
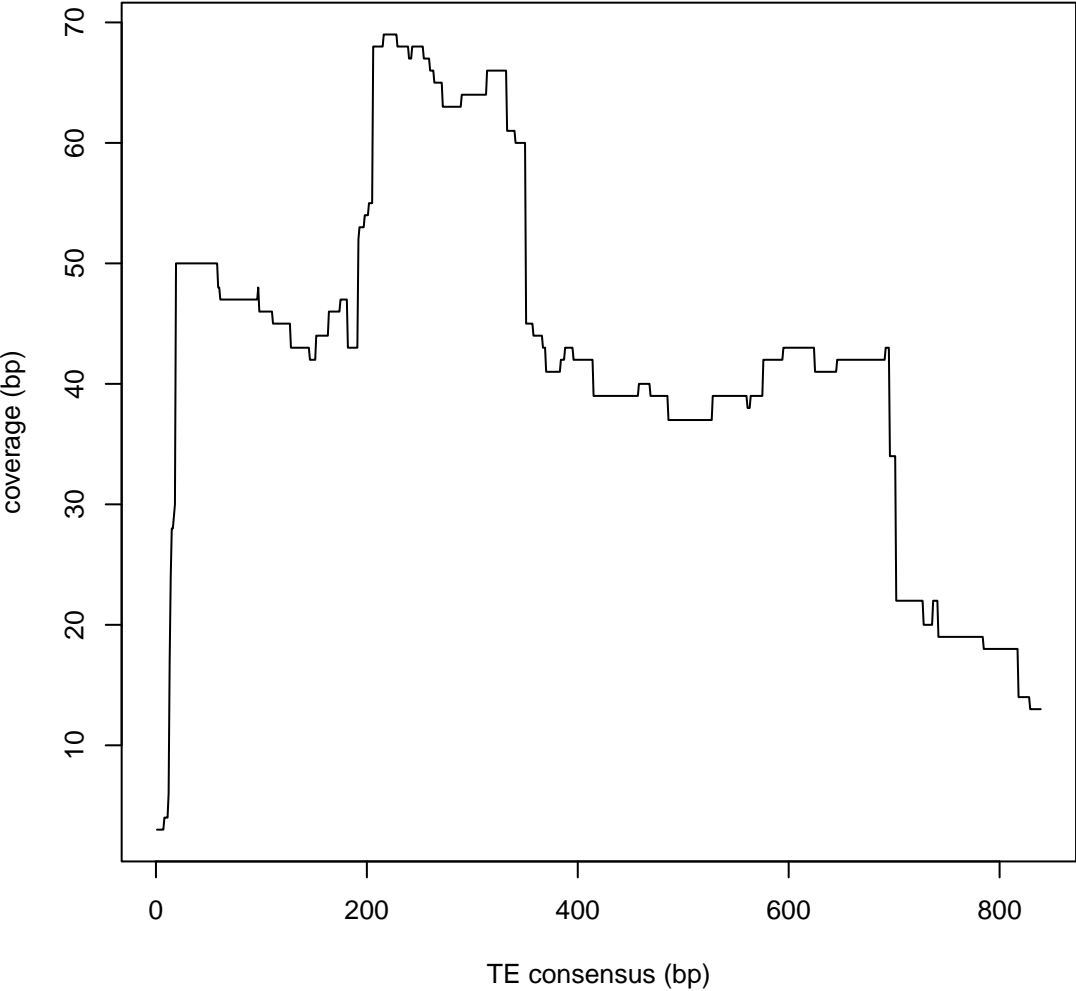


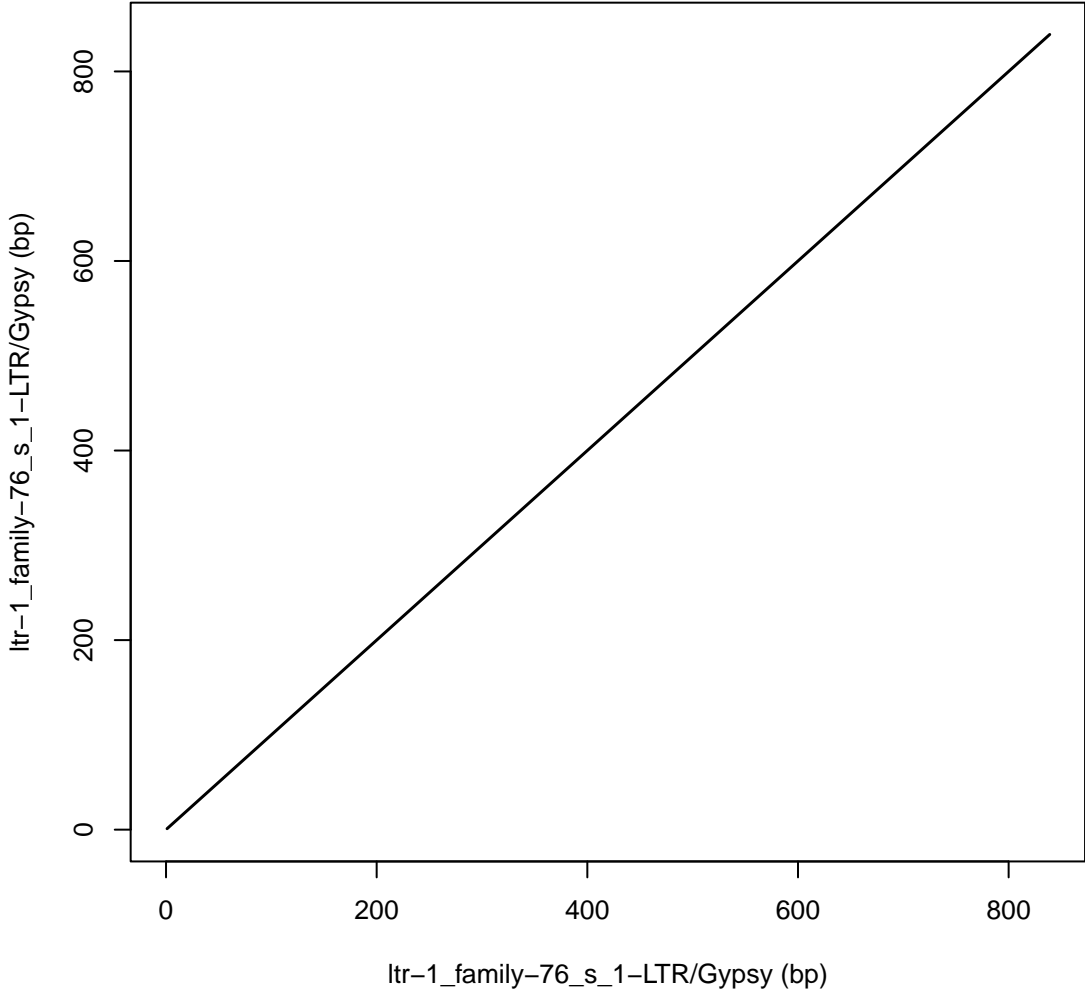
TE: ltr-1_family-76_s_1-LTR/Gypsy
consensus size: 839bp; fragments: 104; full length: 6 (>=755.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

