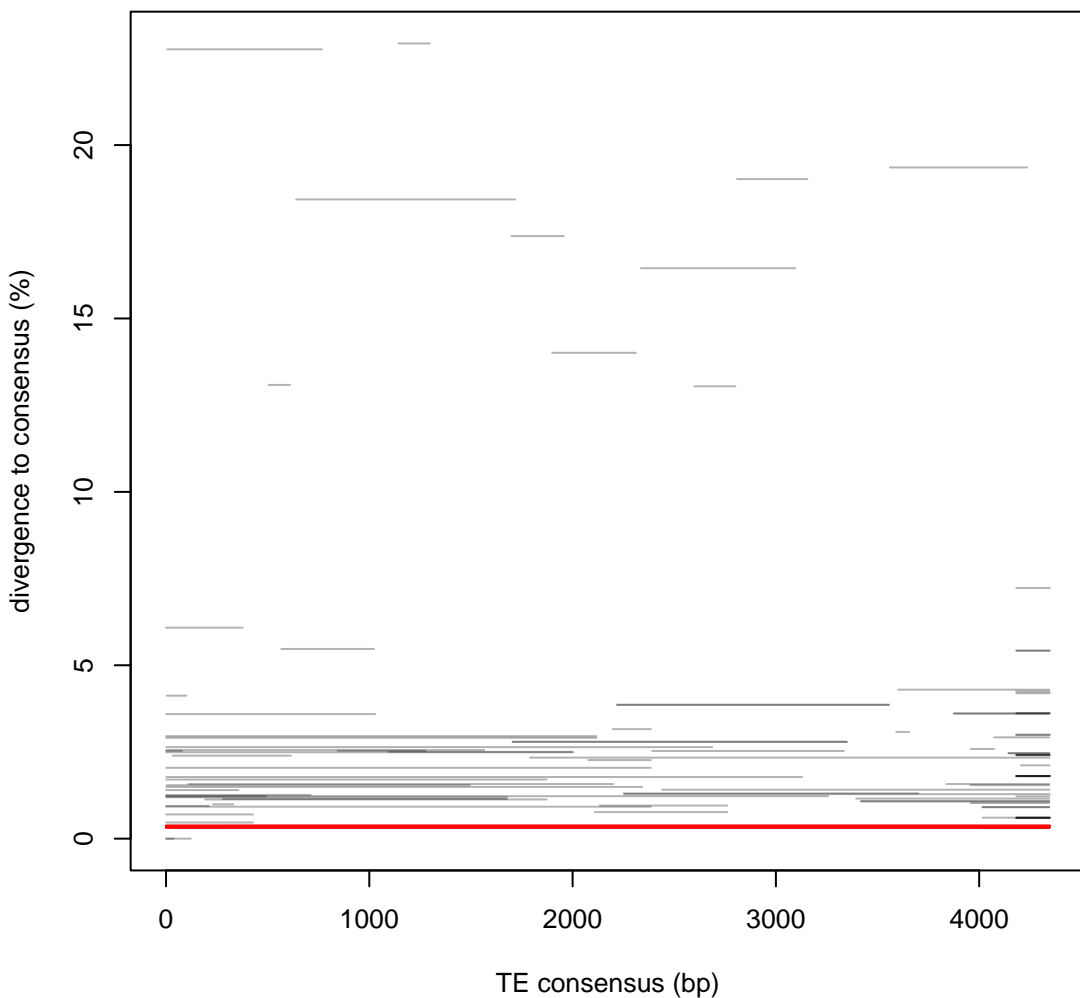
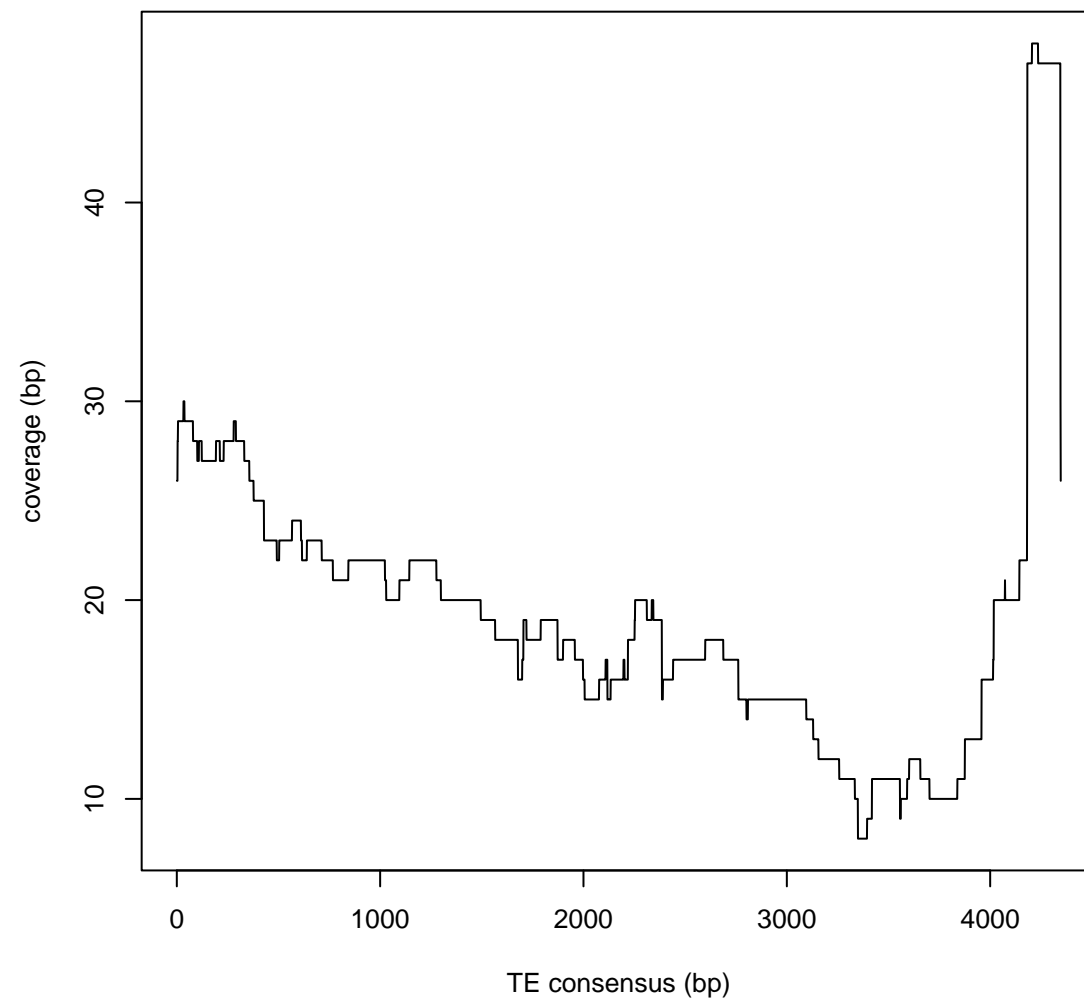


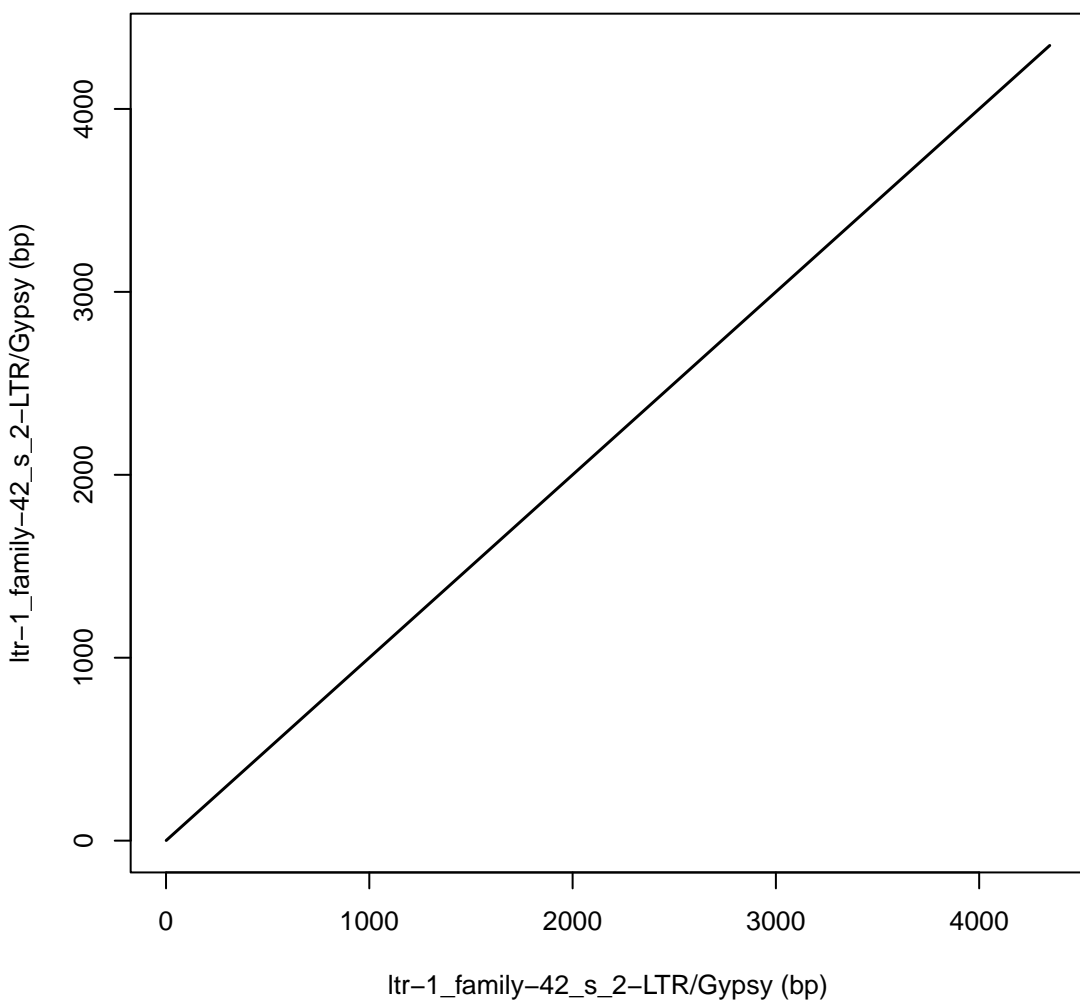
TE: ltr-1_family-42_s_2-LTR/Gypsy
consensus size: 4347bp; fragments: 103; full length: 2 (>=3912.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

