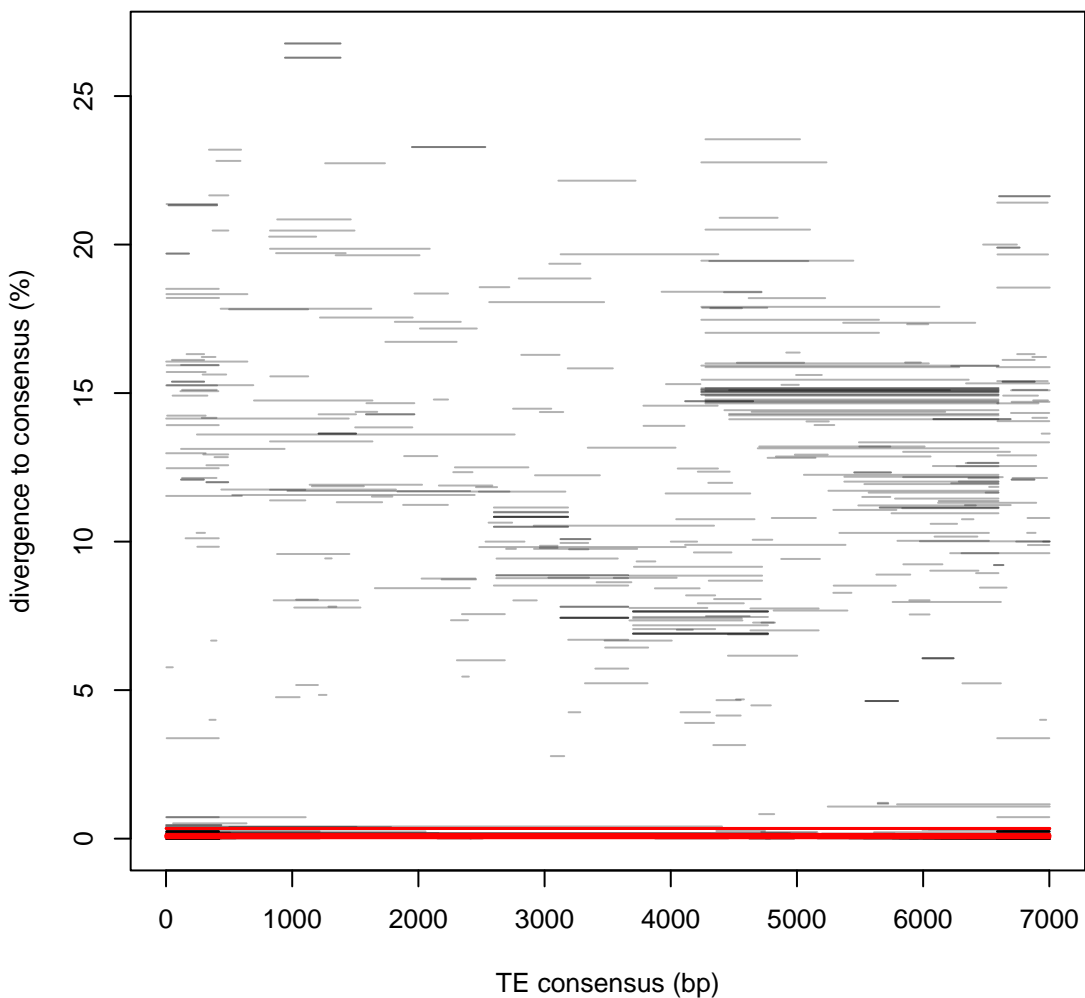
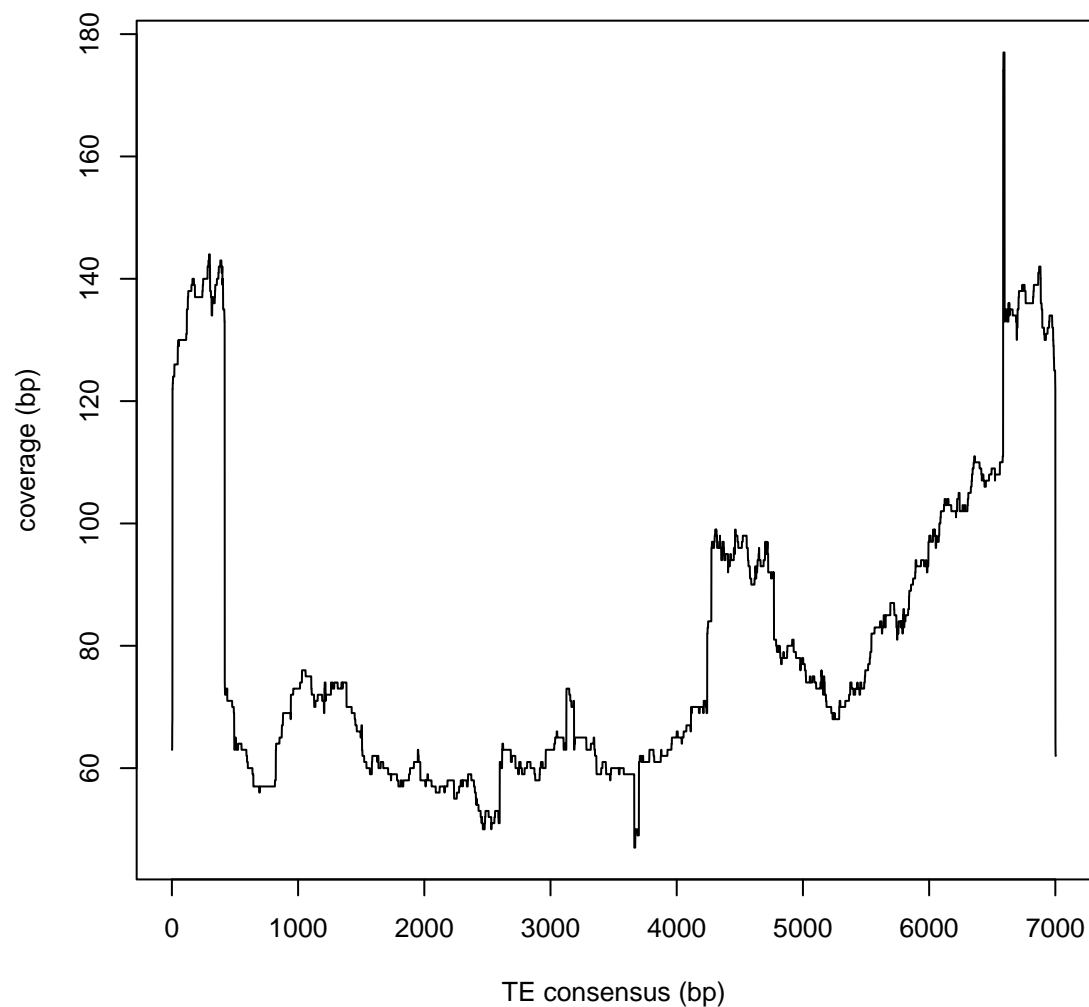


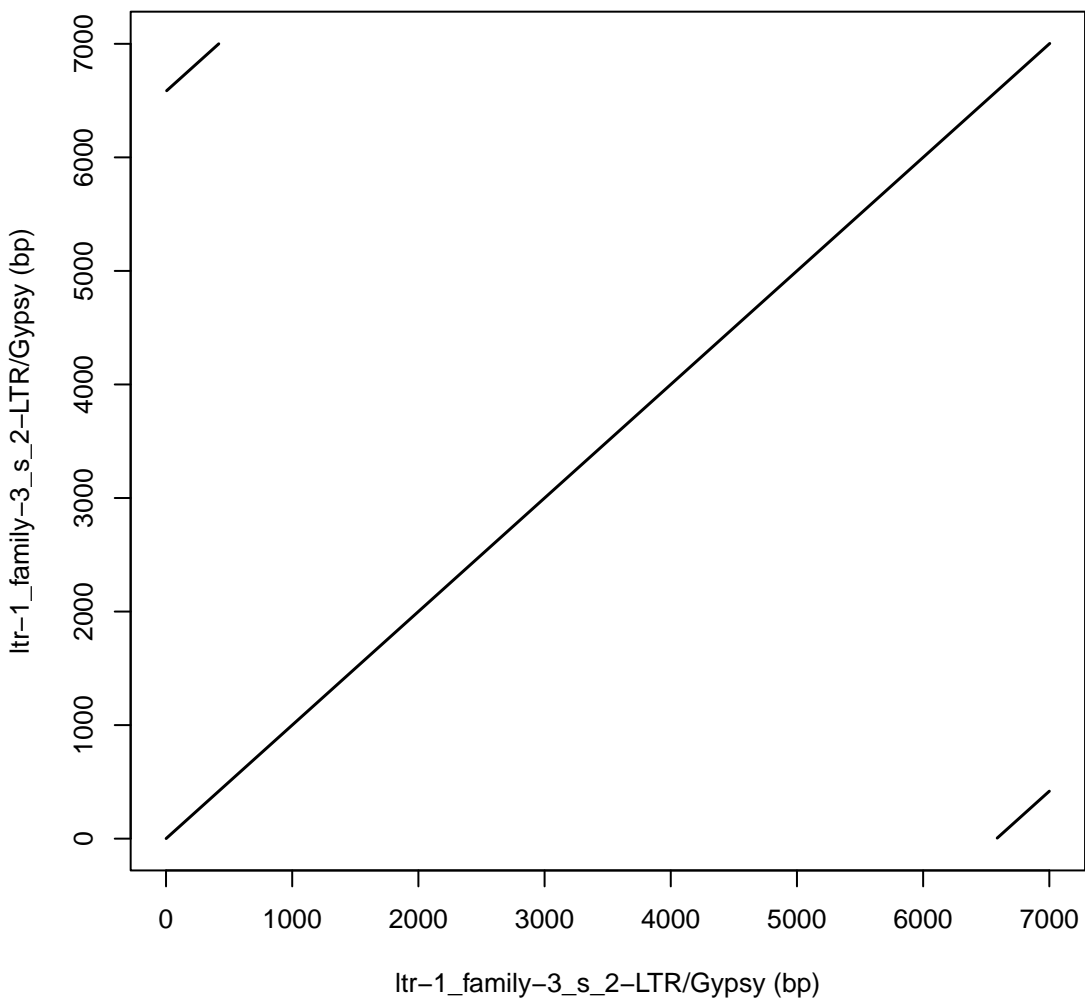
TE: ltr-1_family-3_s_2-LTR/Gypsy
consensus size: 7003bp; fragments: 593; full length: 18 (>=6302.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

