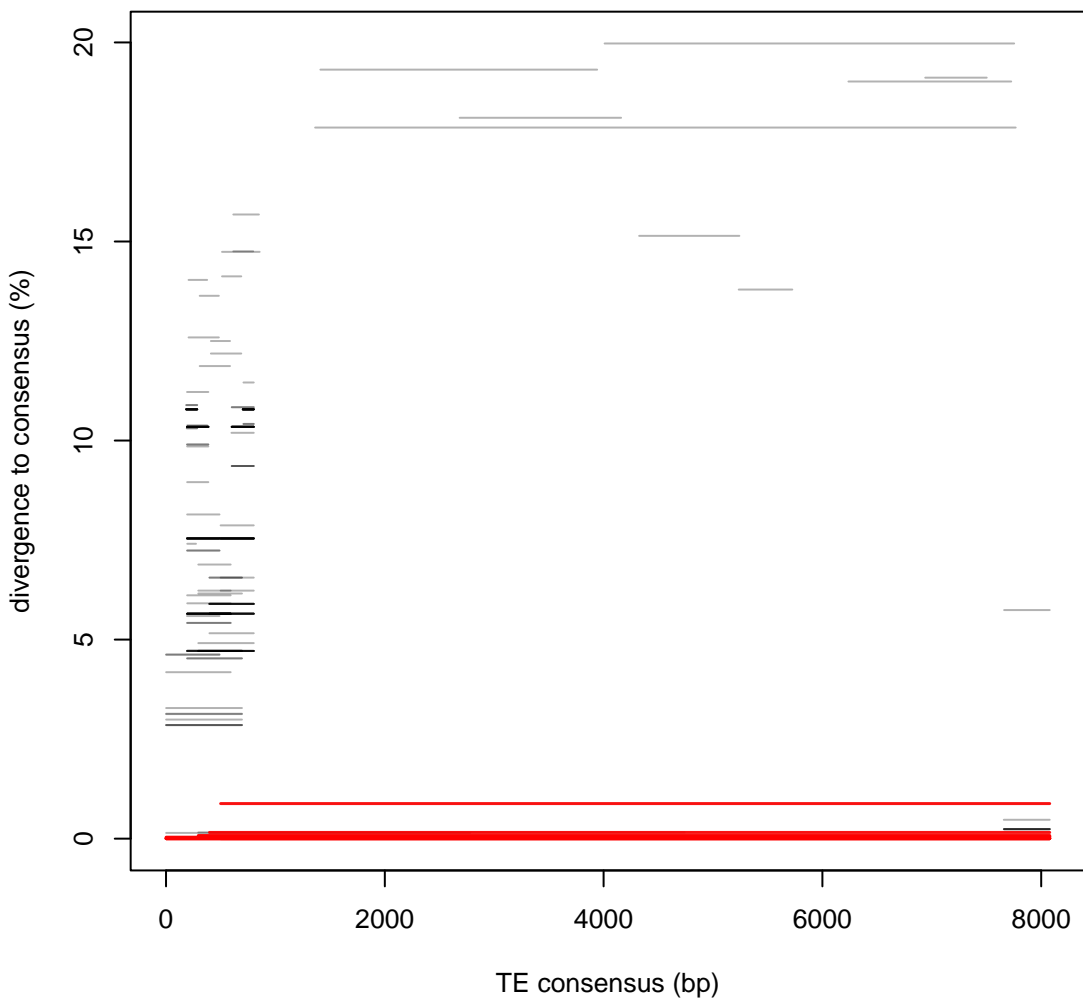
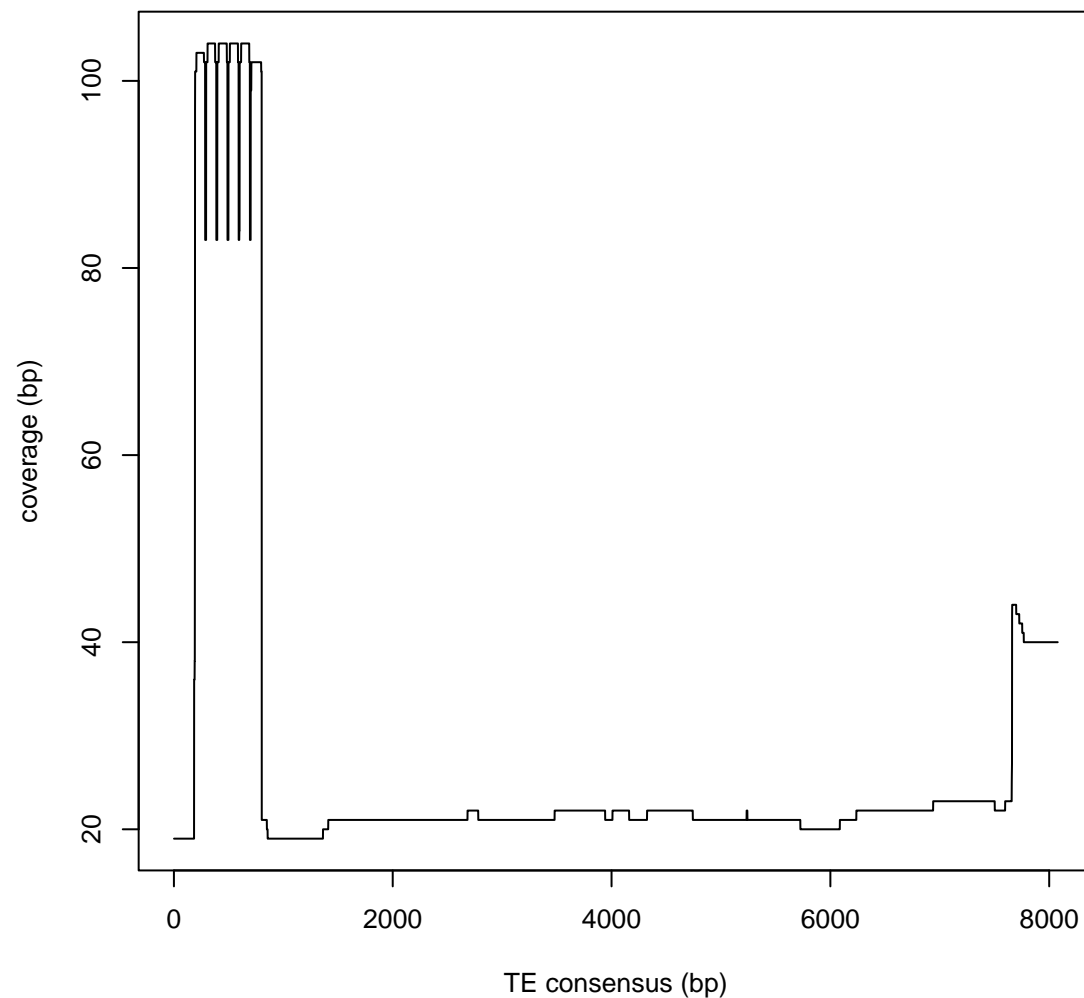


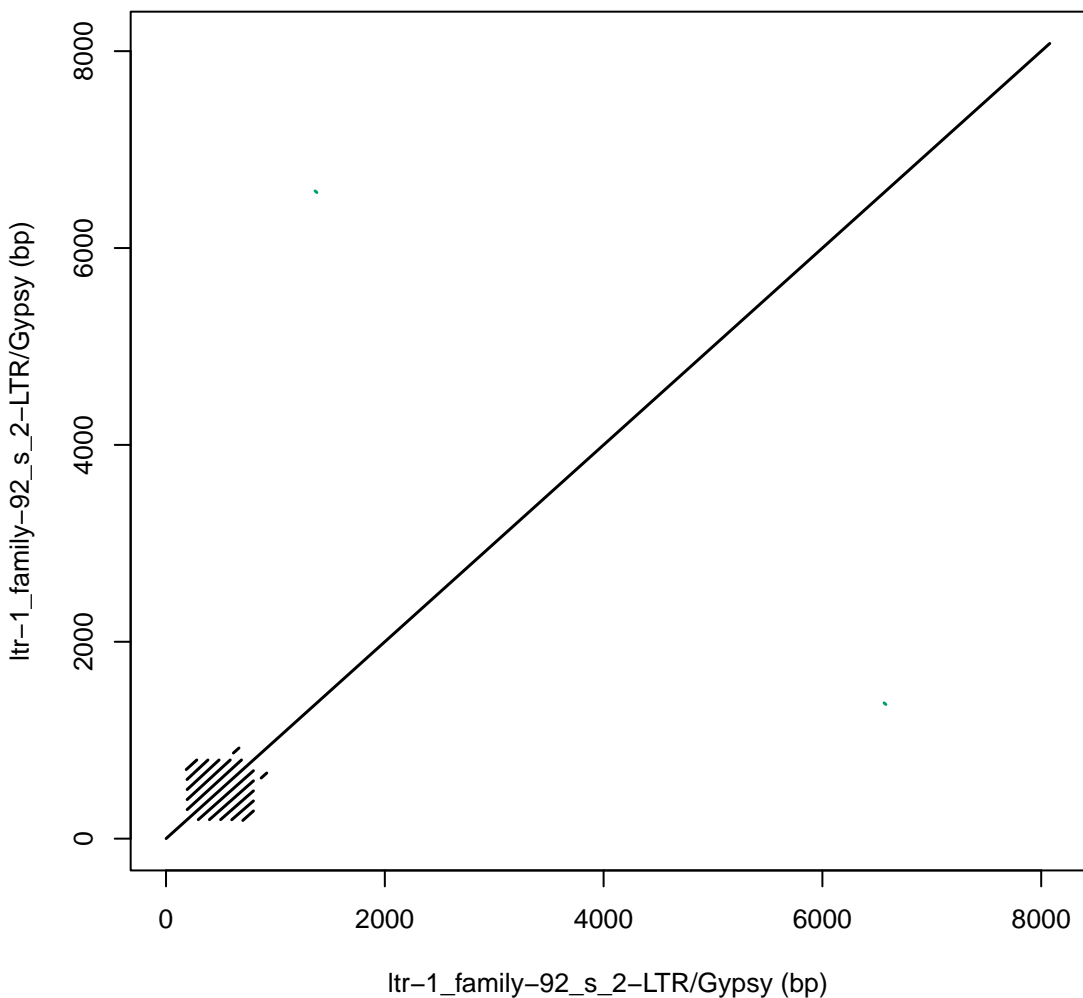
TE: ltr-1_family-92_s_2-LTR/Gypsy
consensus size: 8078bp; fragments: 238; full length: 17 (>=7270.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

