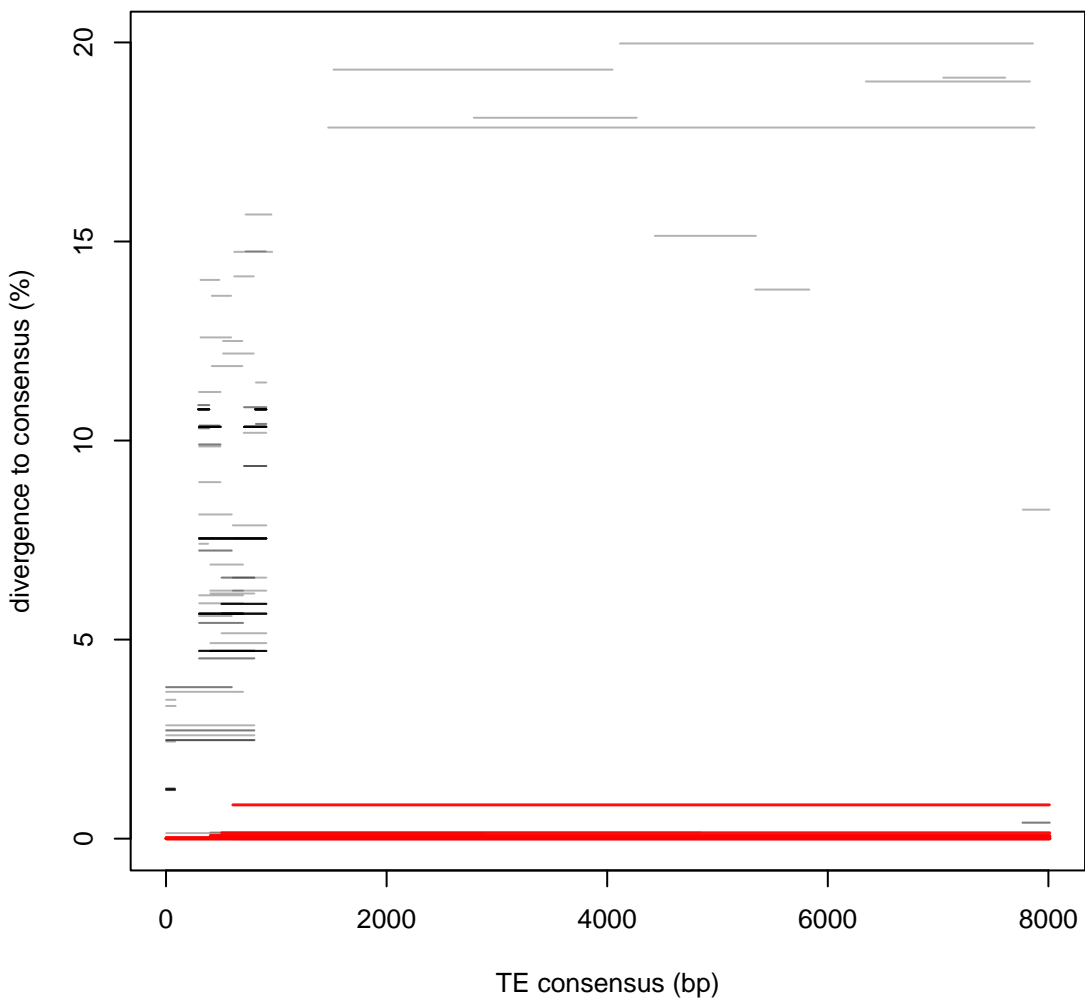
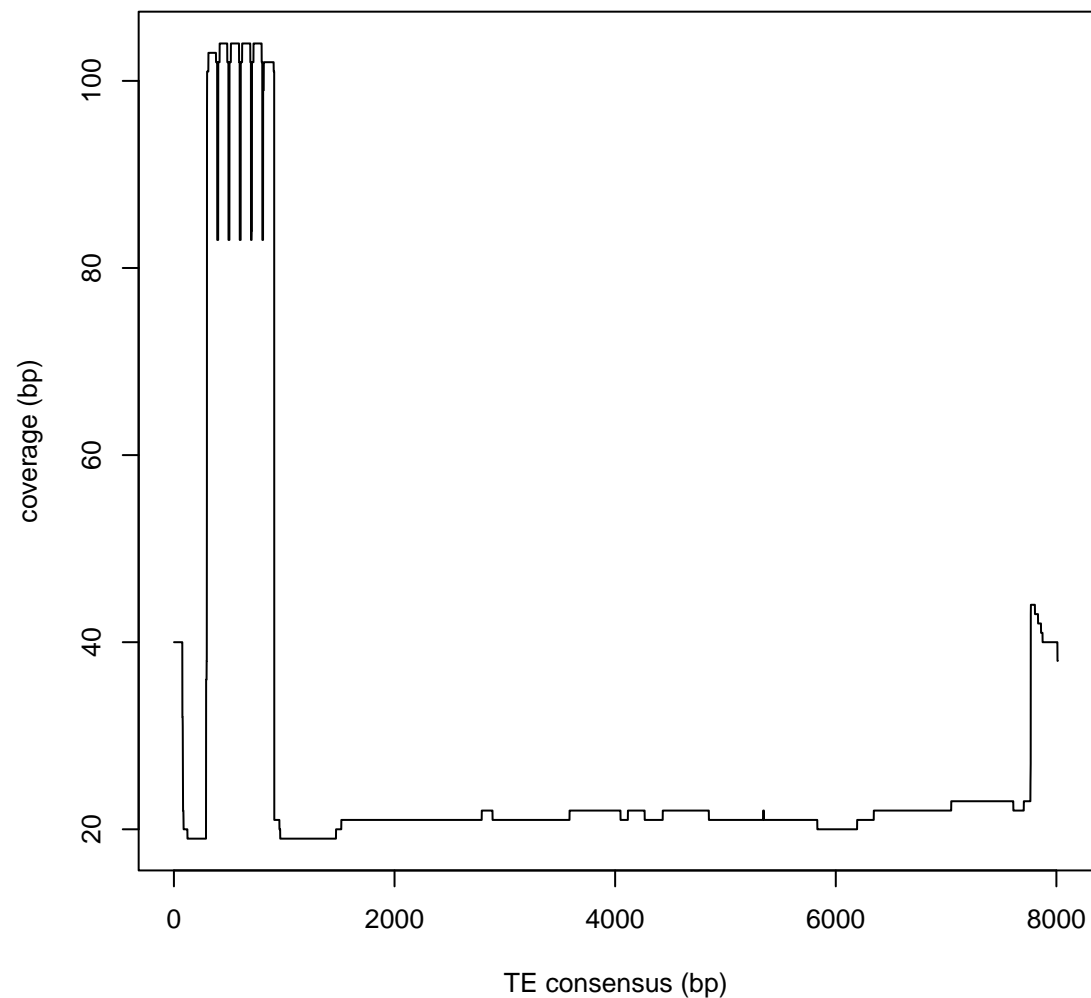


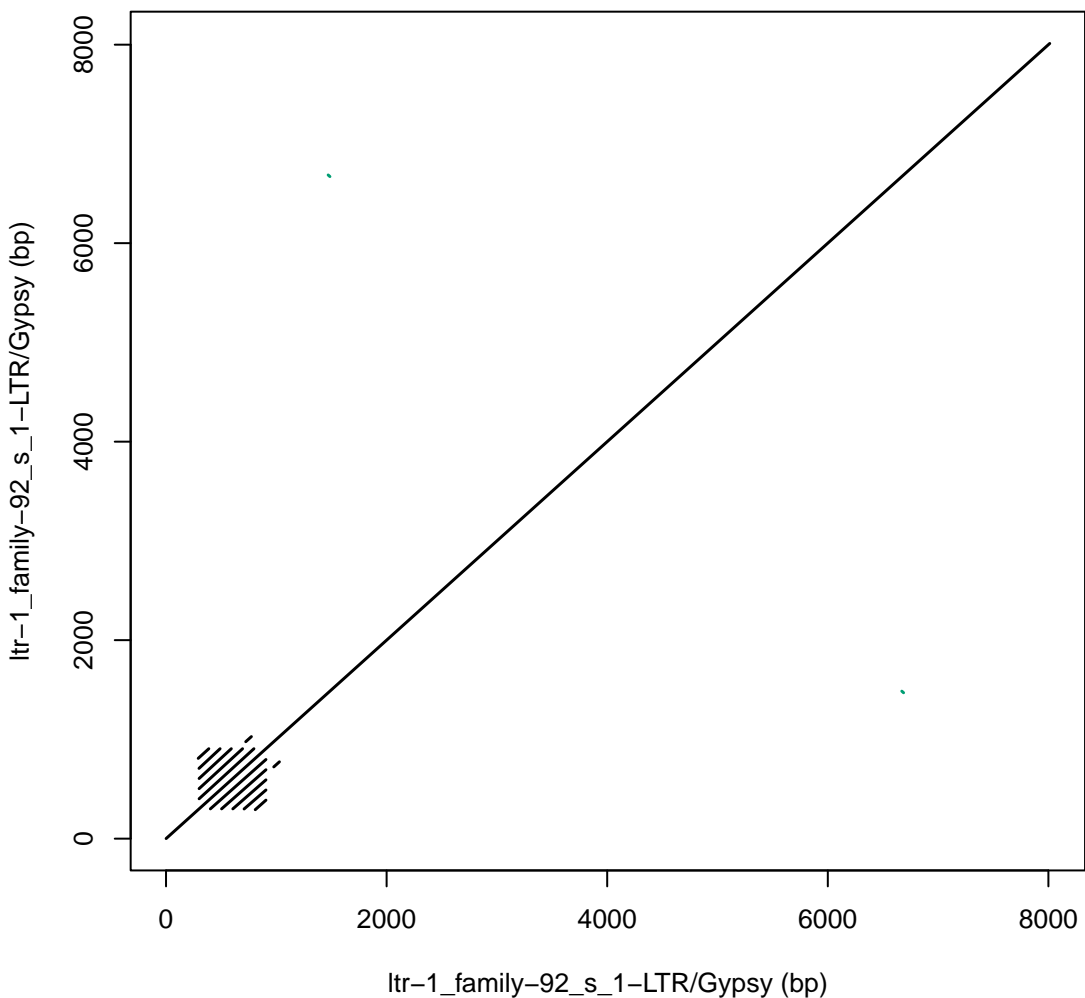
TE: ltr-1_family-92_s_1-LTR/Gypsy
consensus size: 8012bp; fragments: 259; full length: 17 (>=7210.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

