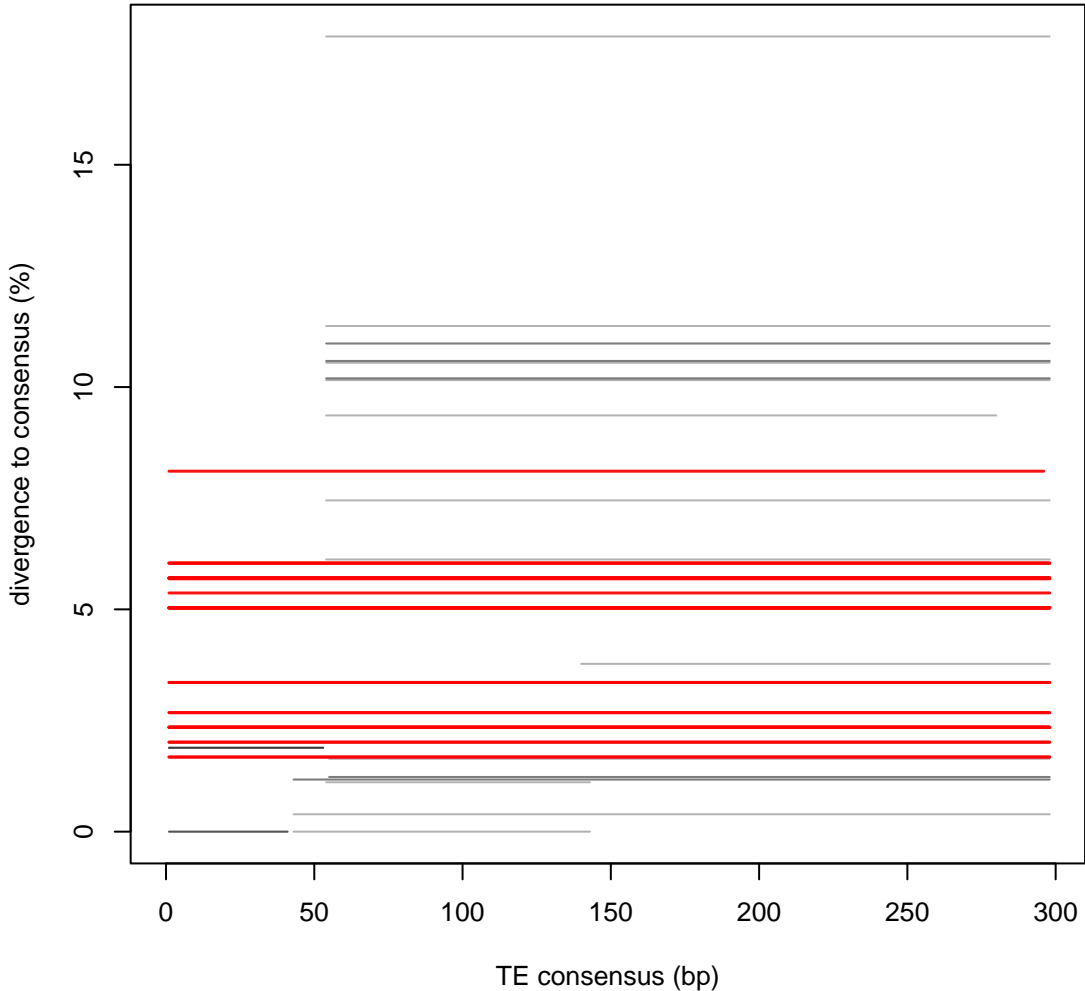
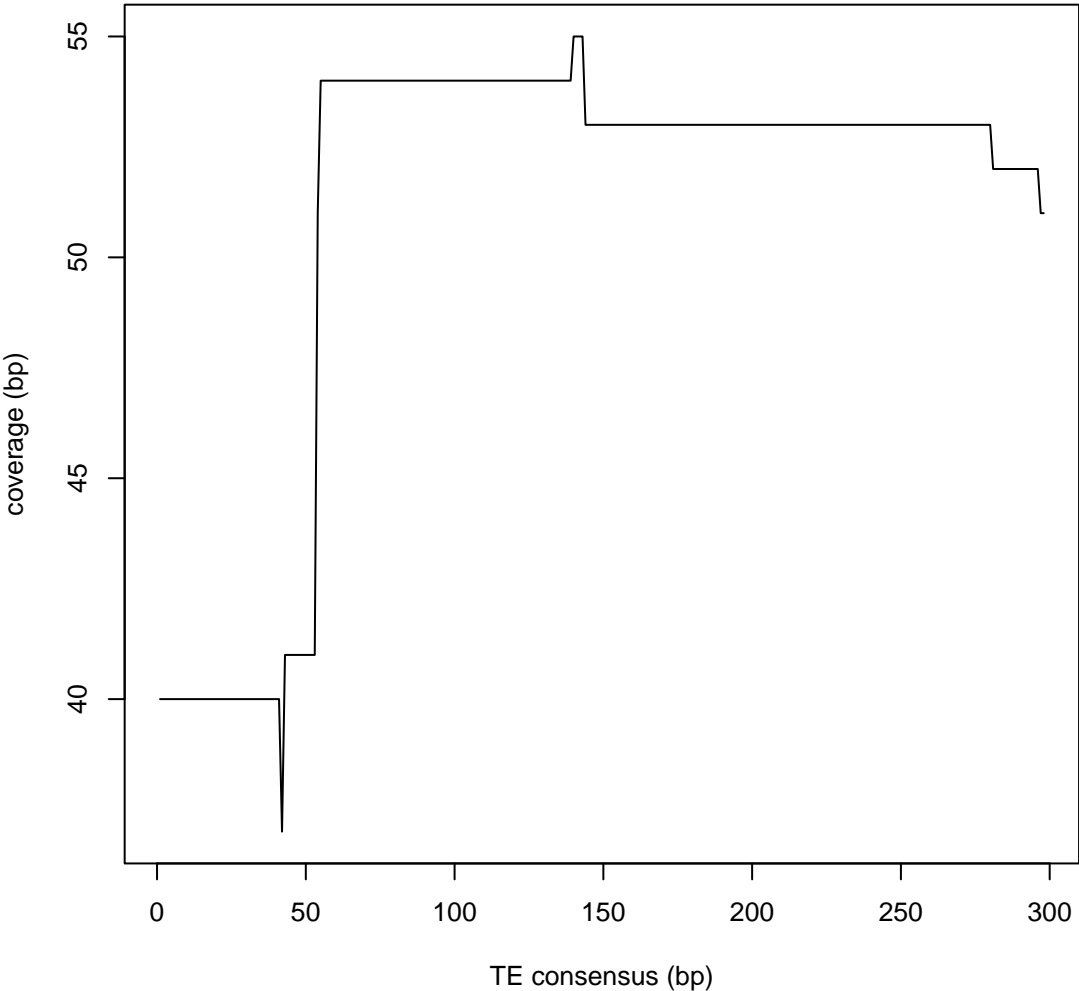


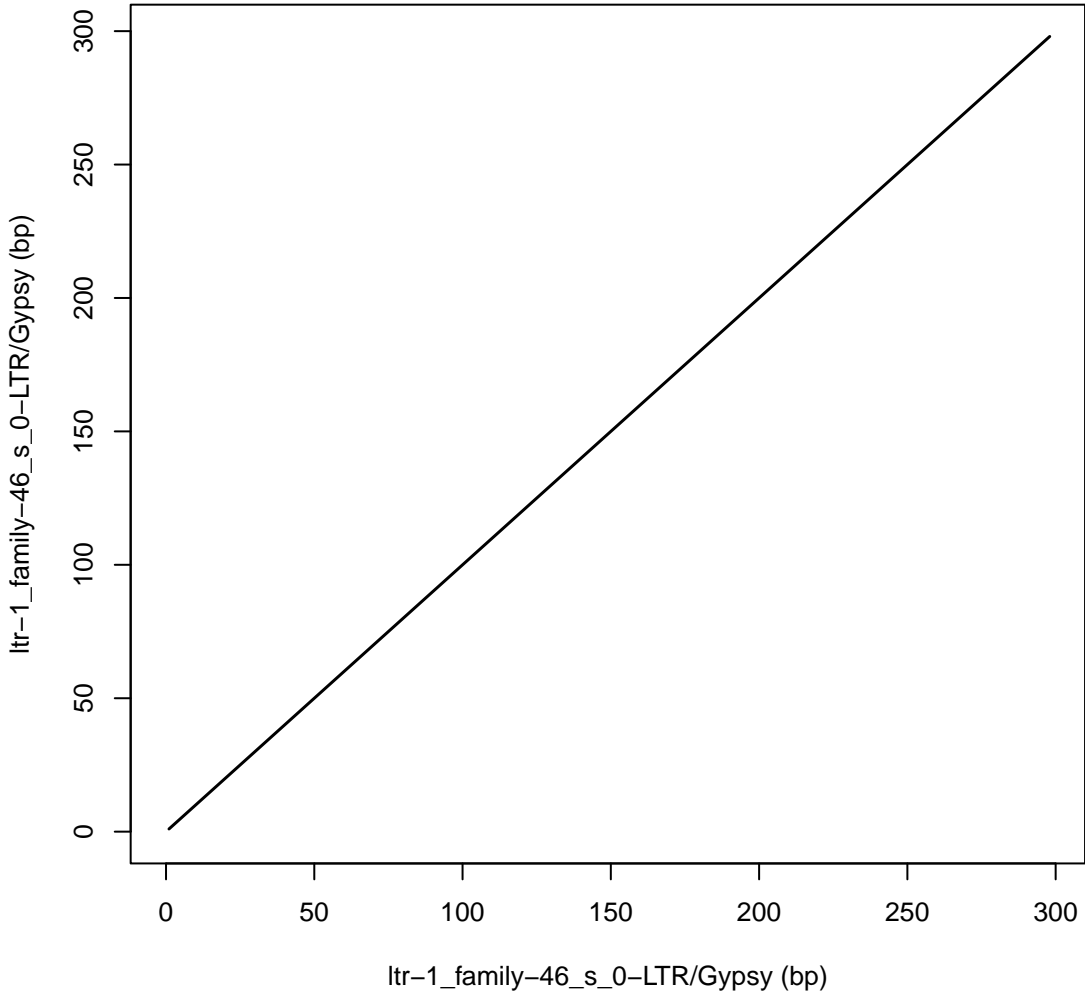
TE: ltr-1_family-46_s_0-LTR/Gypsy
consensus size: 298bp; fragments: 62; full length: 33 (>=268.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

