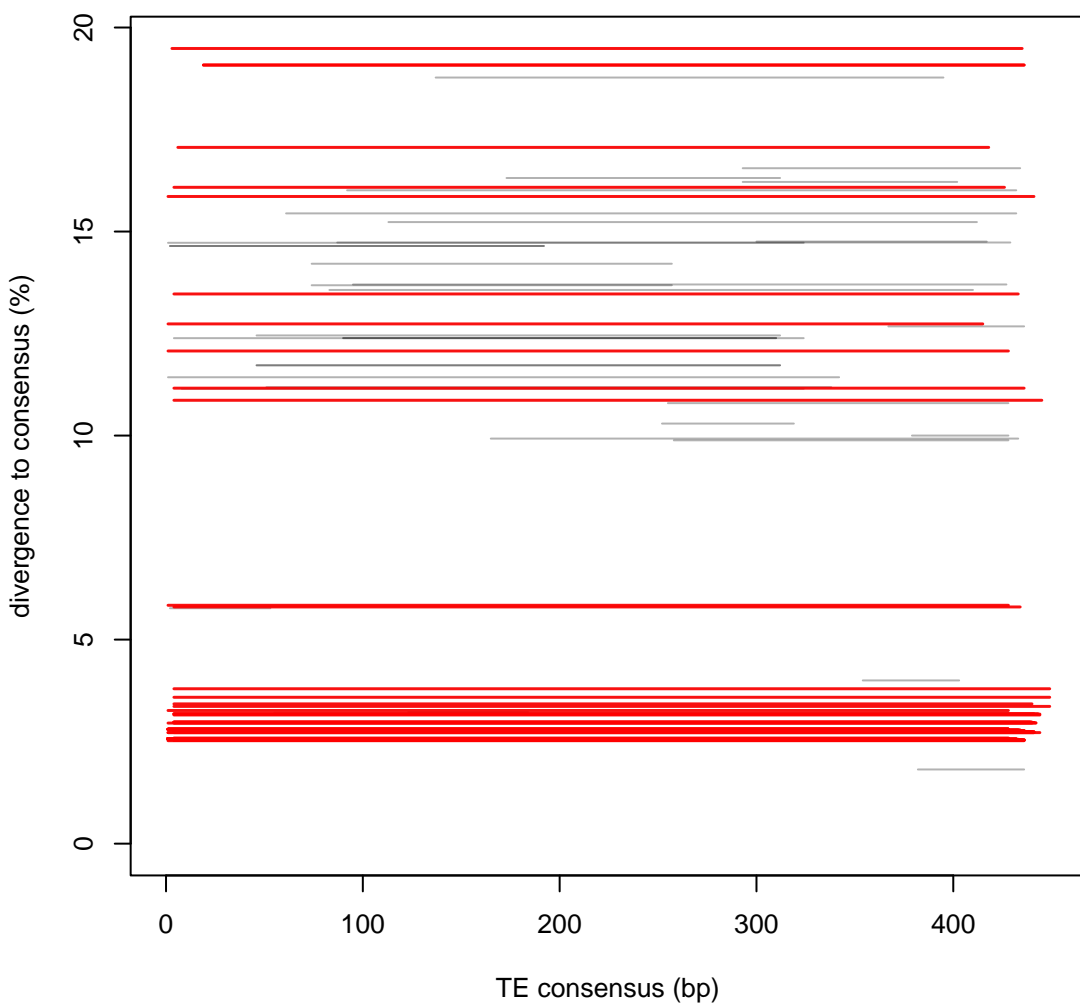
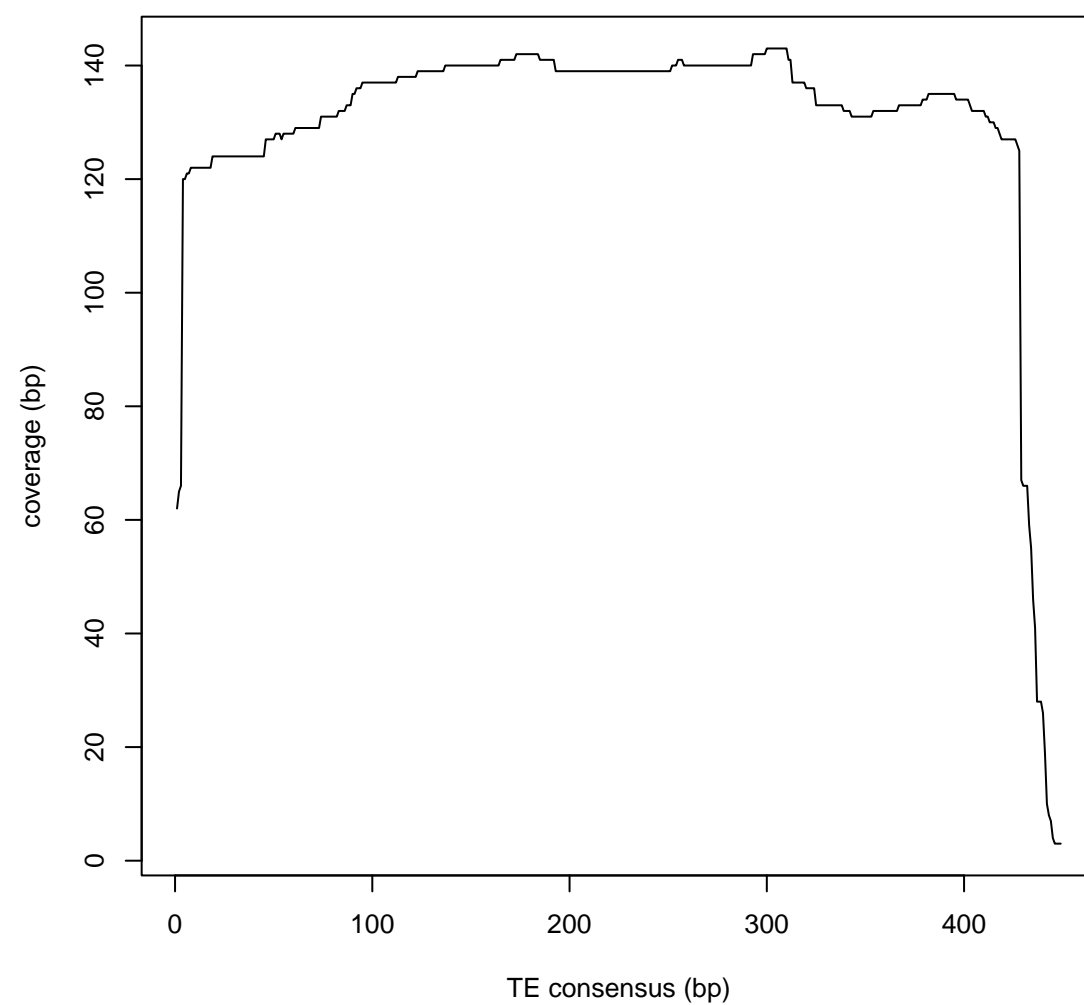


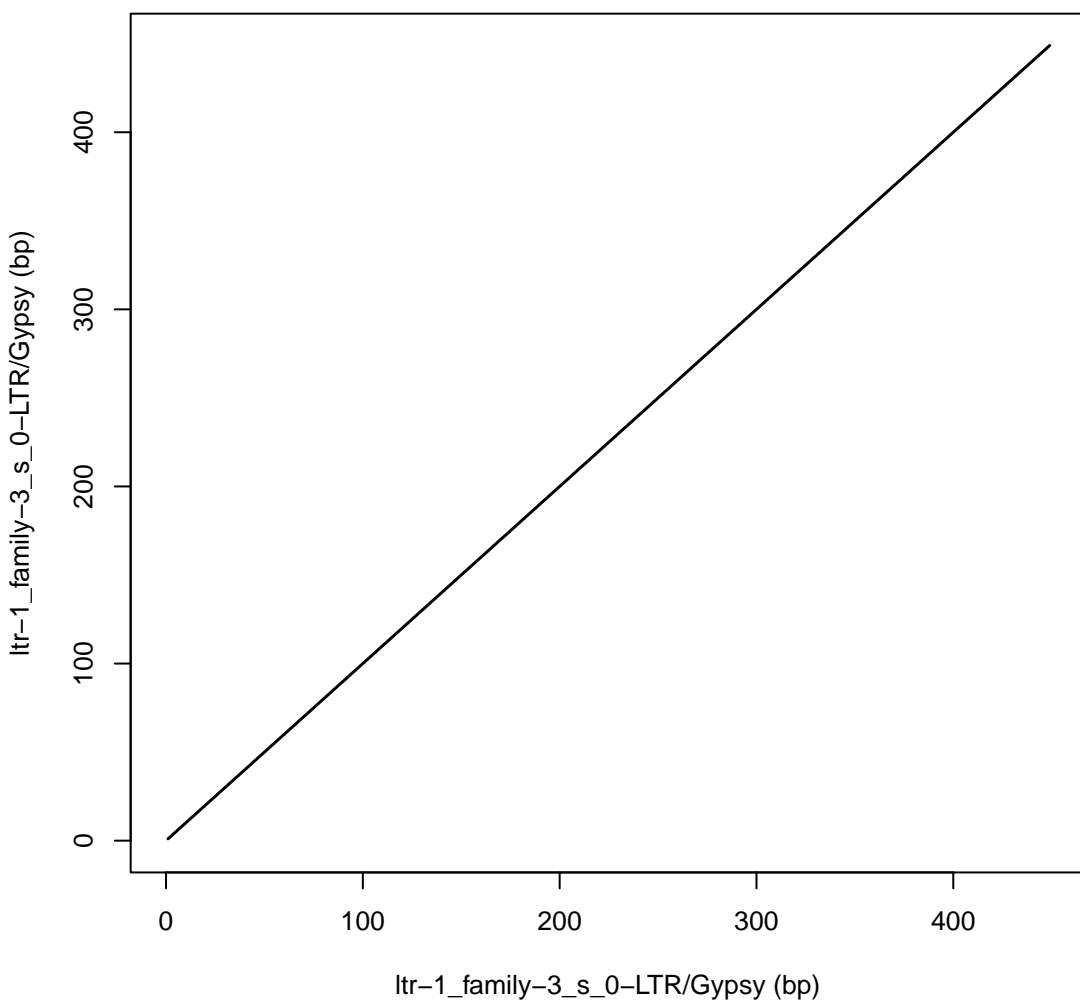
TE: ltr-1_family-3_s_0-LTR/Gypsy
consensus size: 449bp; fragments: 153; full length: 117 (>=404.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

