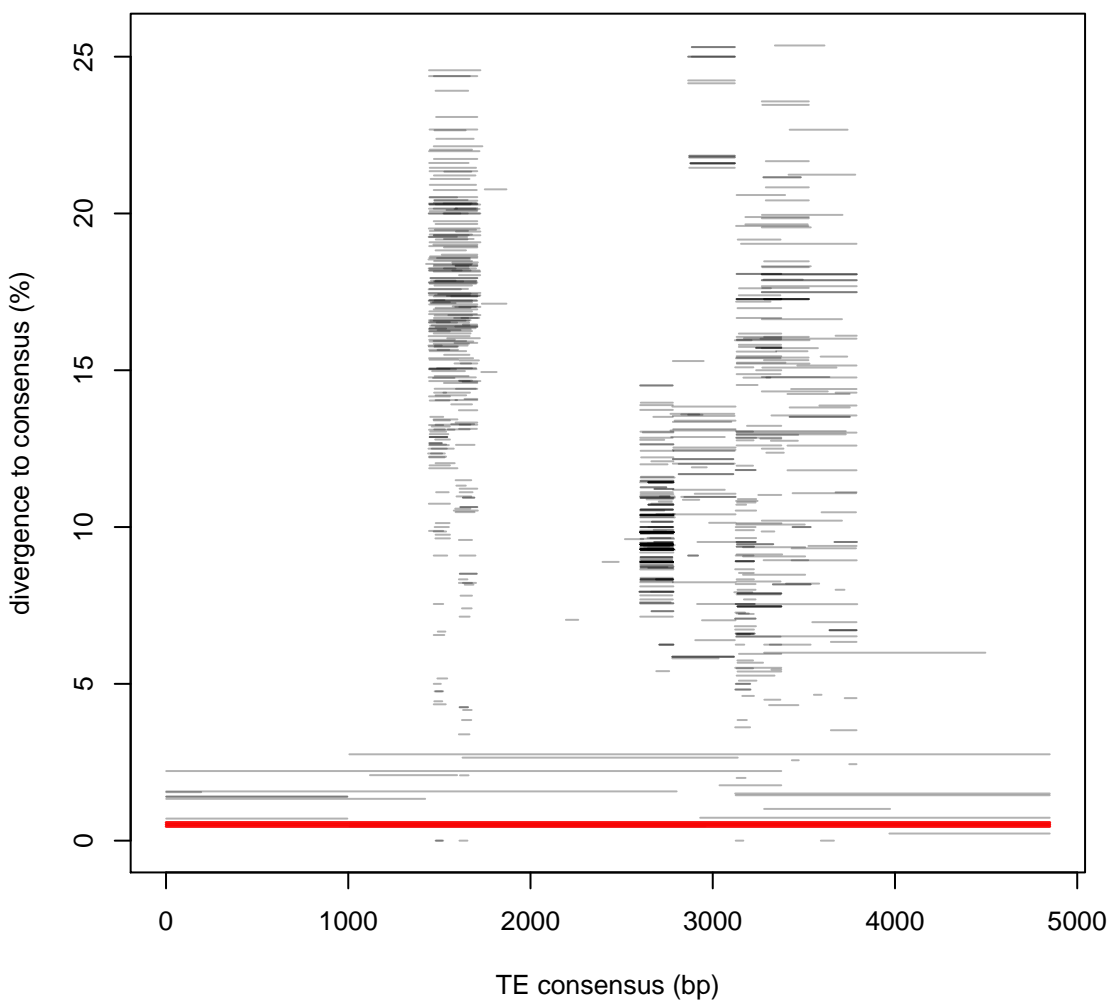
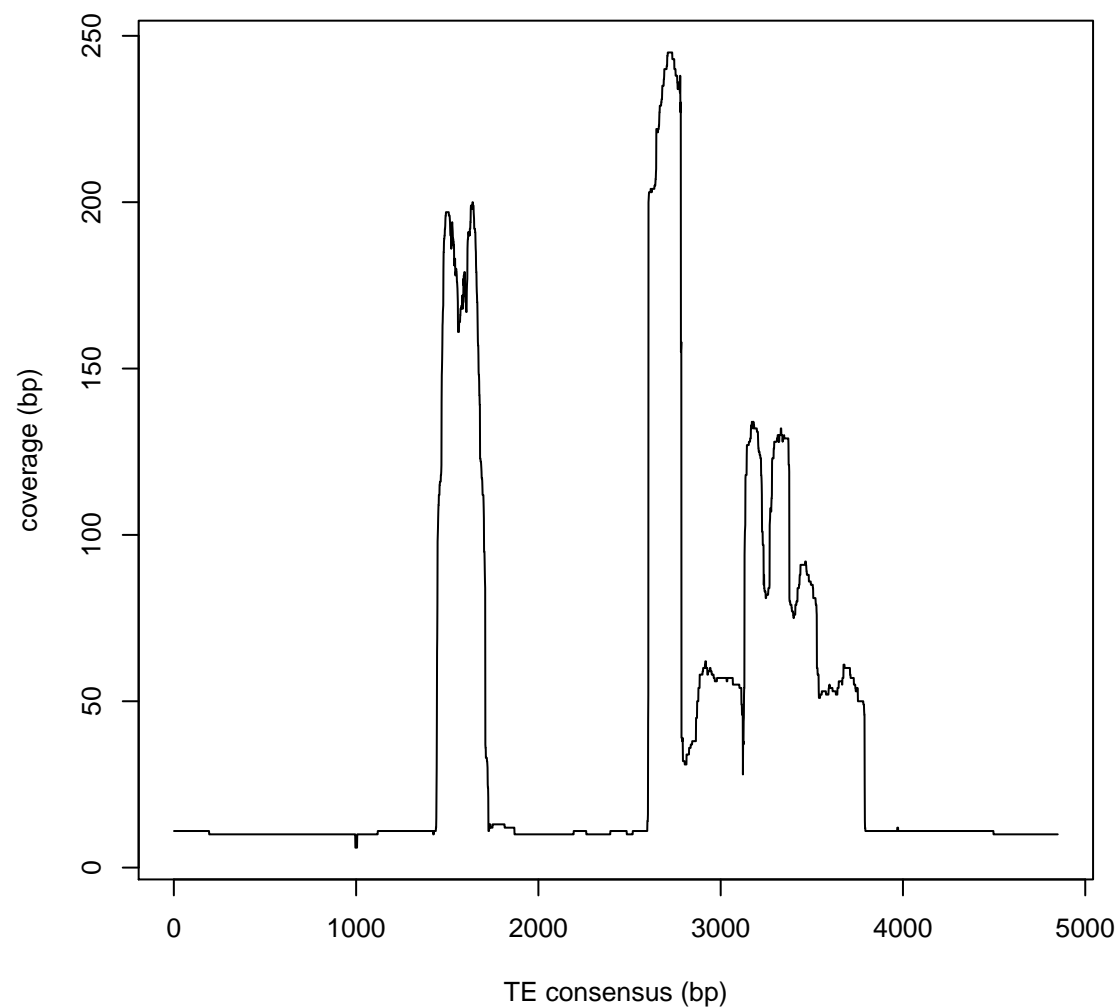


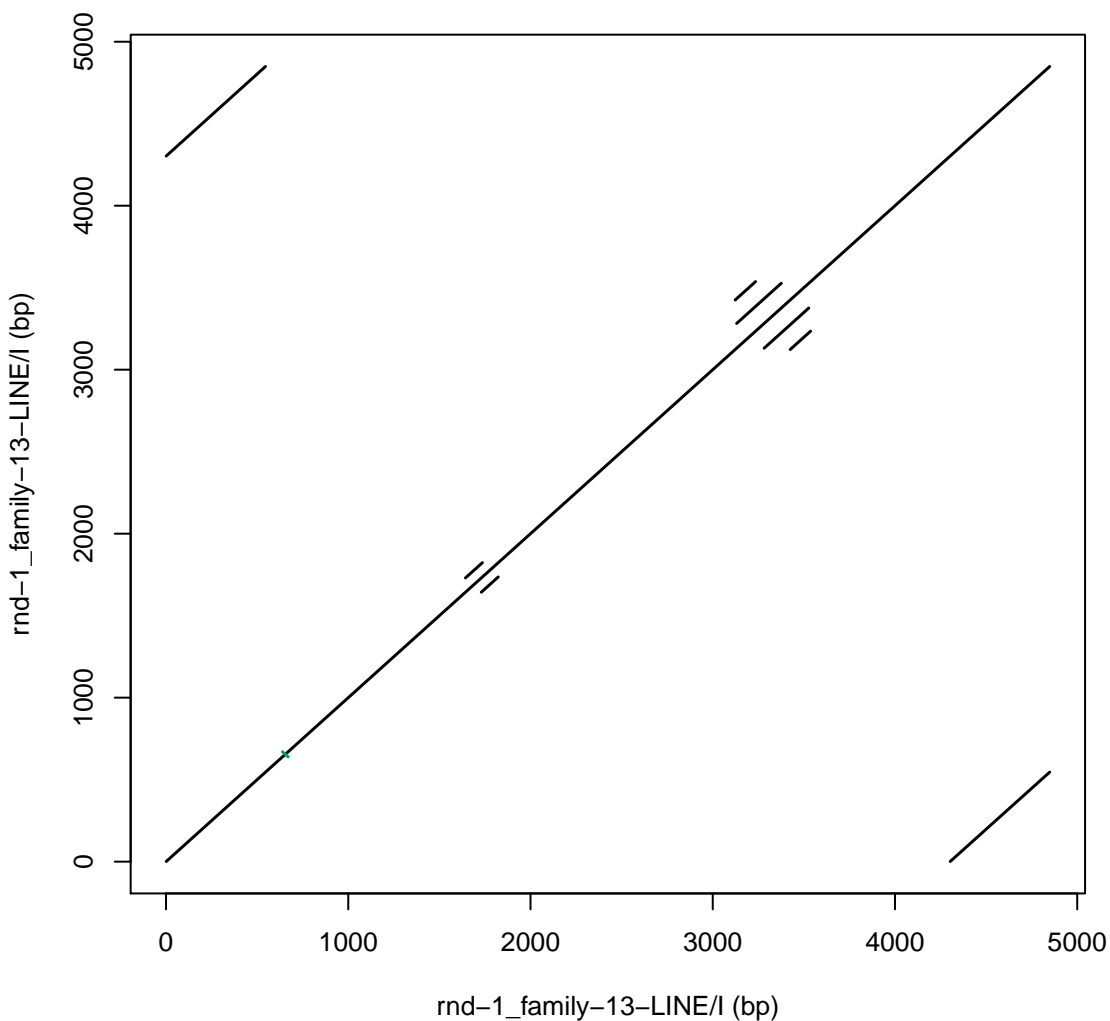
TE: rnd-1_family-13-LINE/I
consensus size: 4849bp; fragments: 811; full length: 3 (>=4364.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

