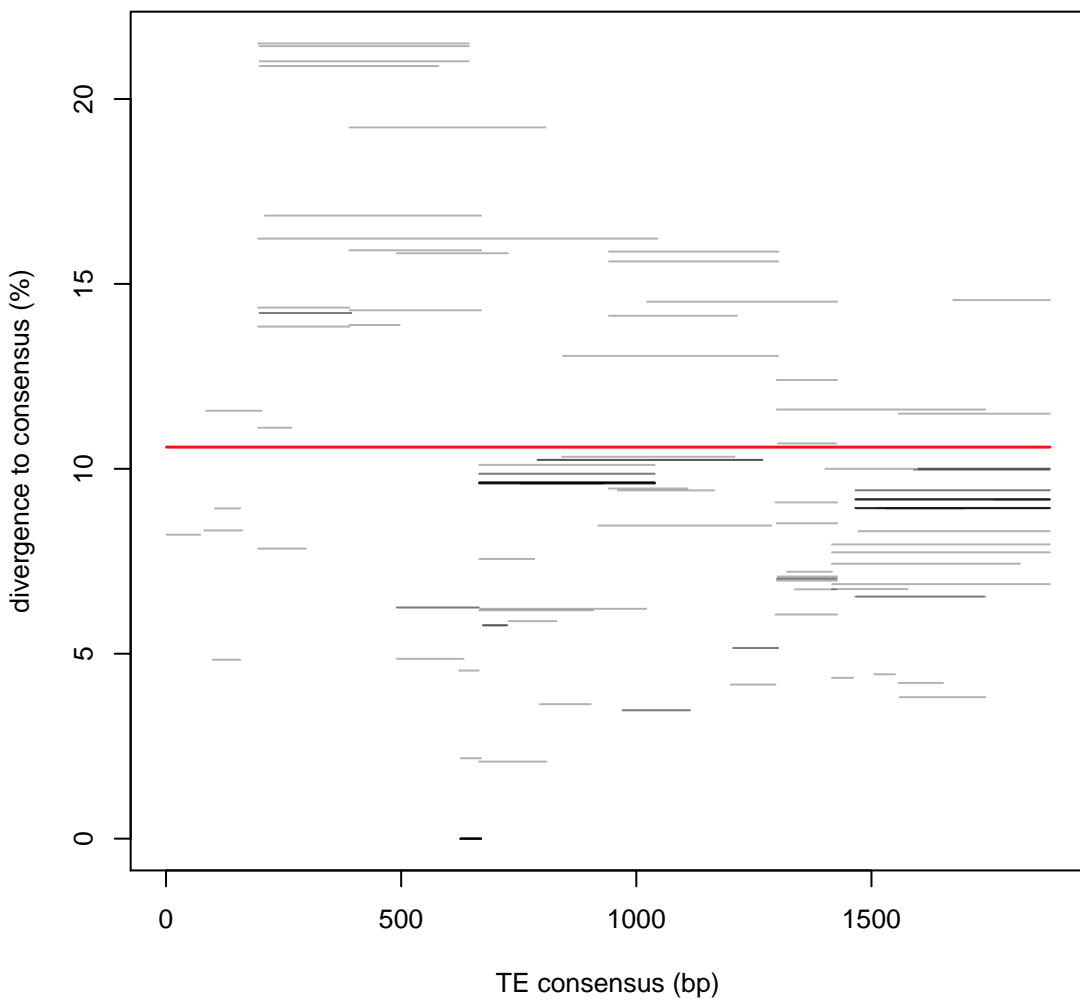
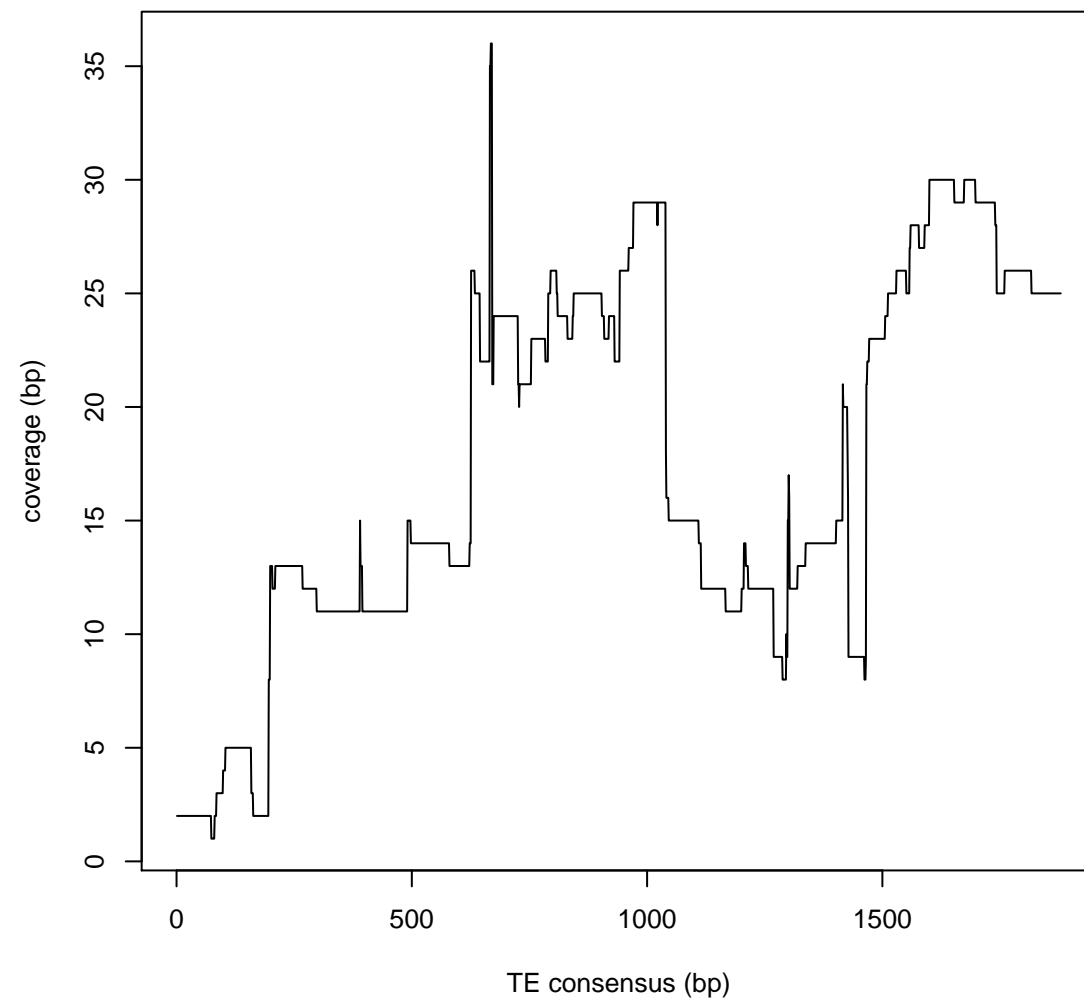


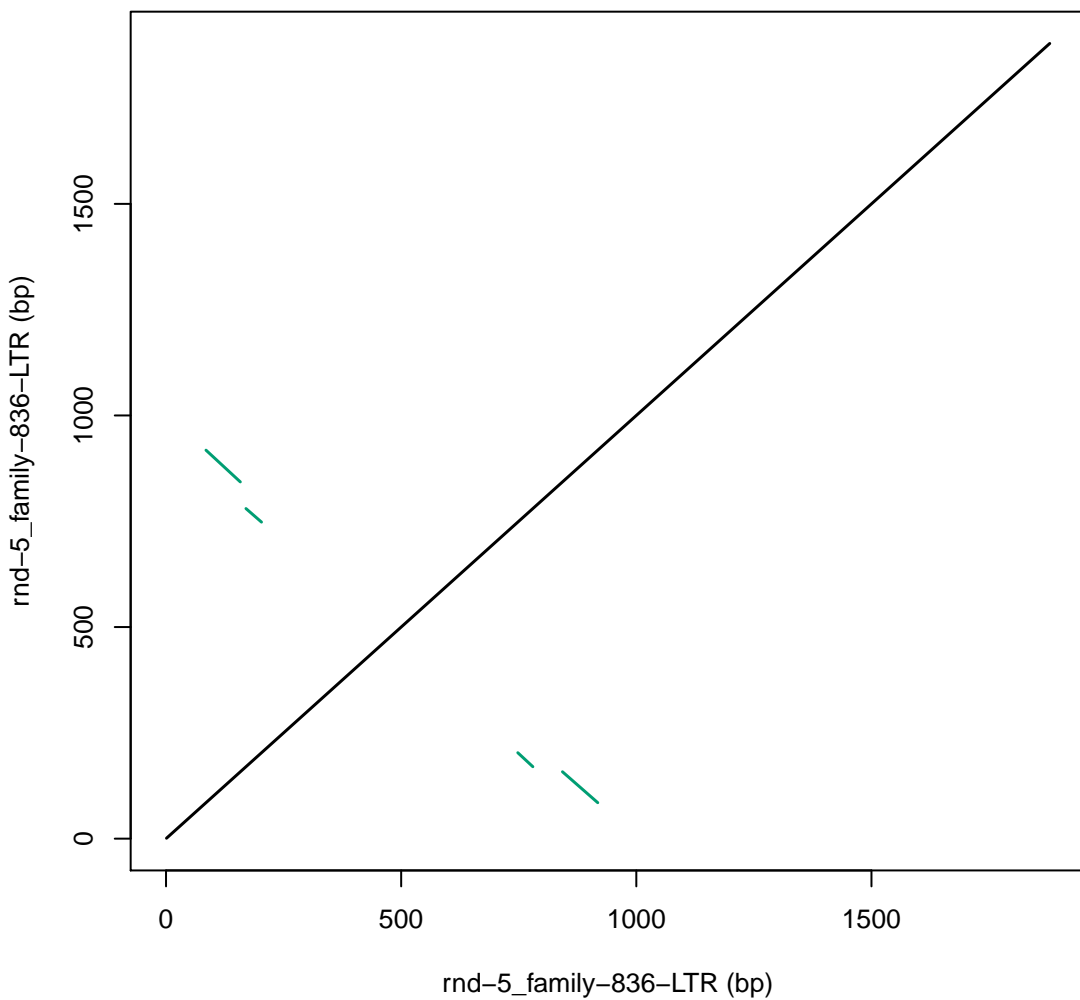
TE: rnd-5_family-836-LTR
consensus size: 1879bp; fragments: 125; full length: 1 (>=1691.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

