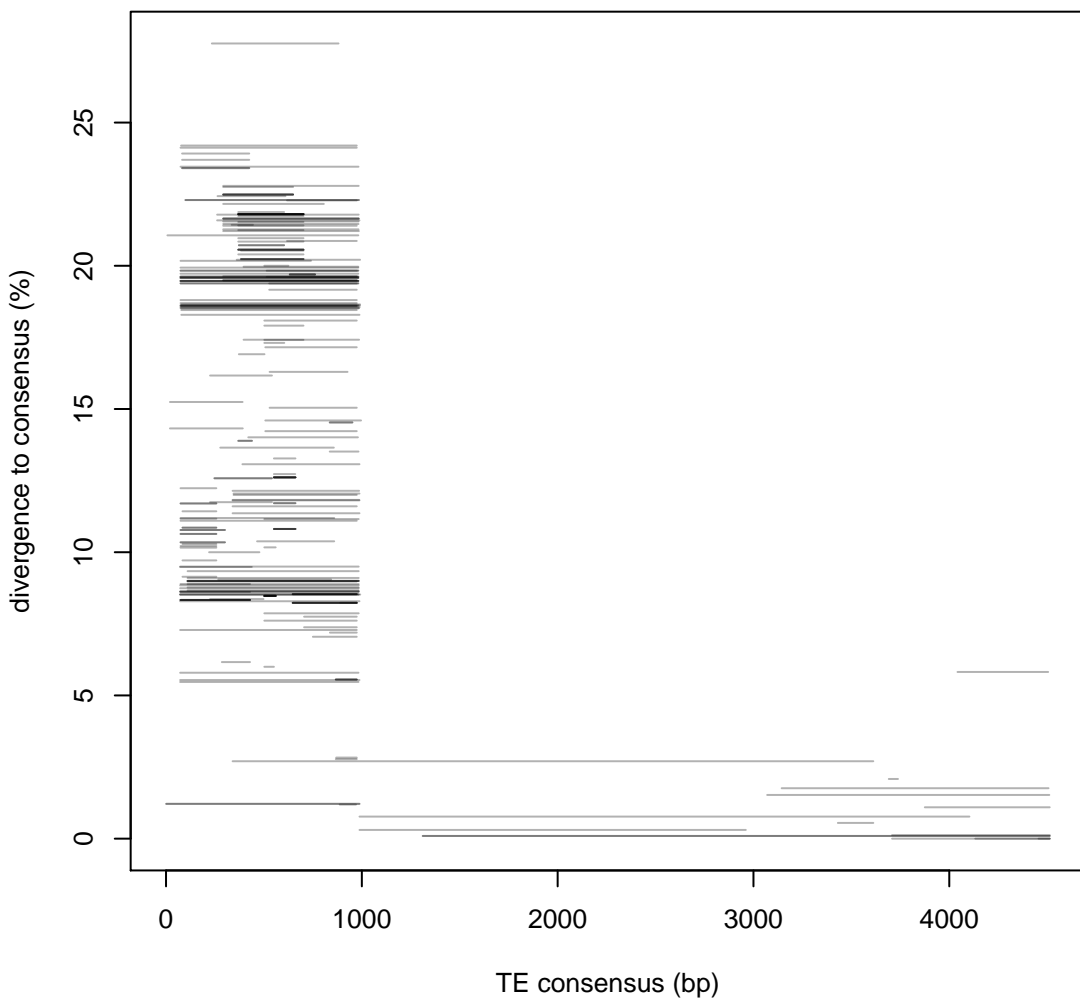
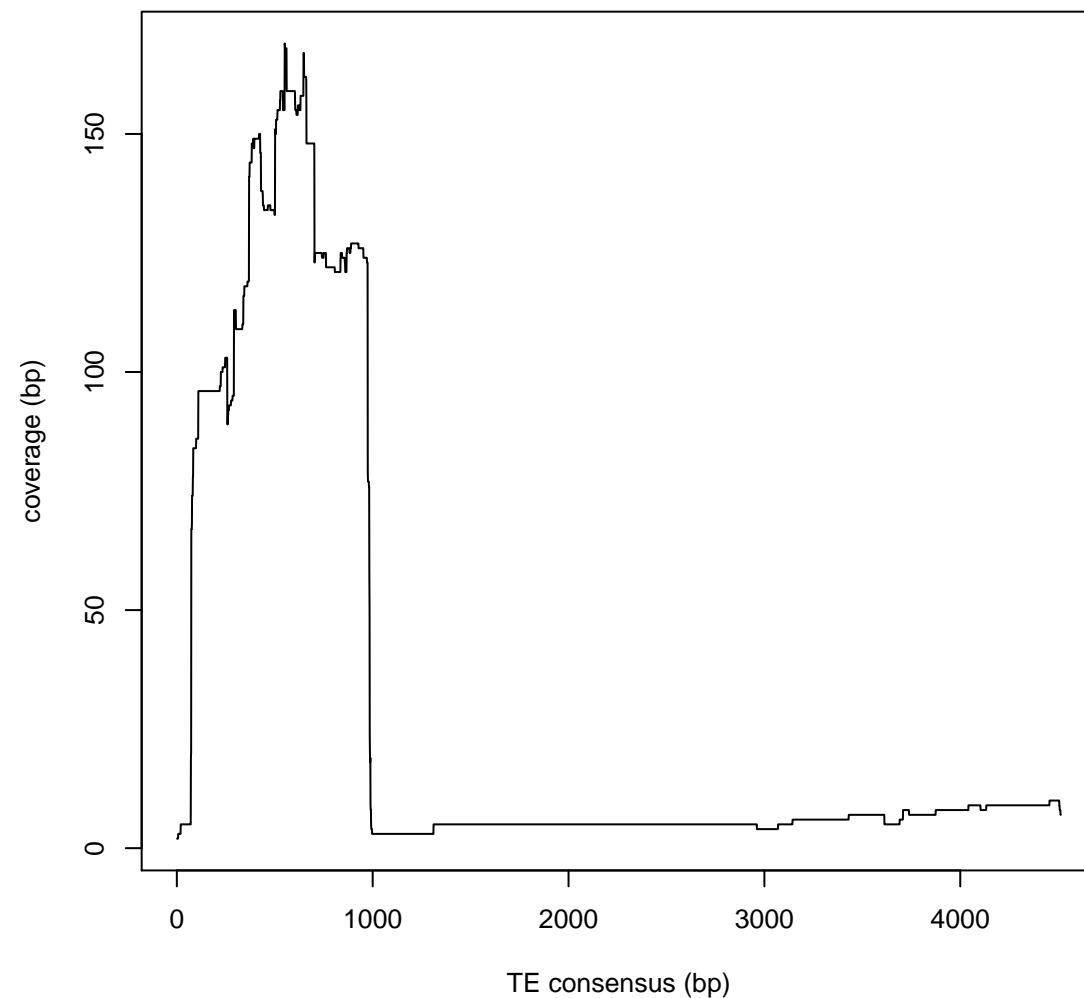


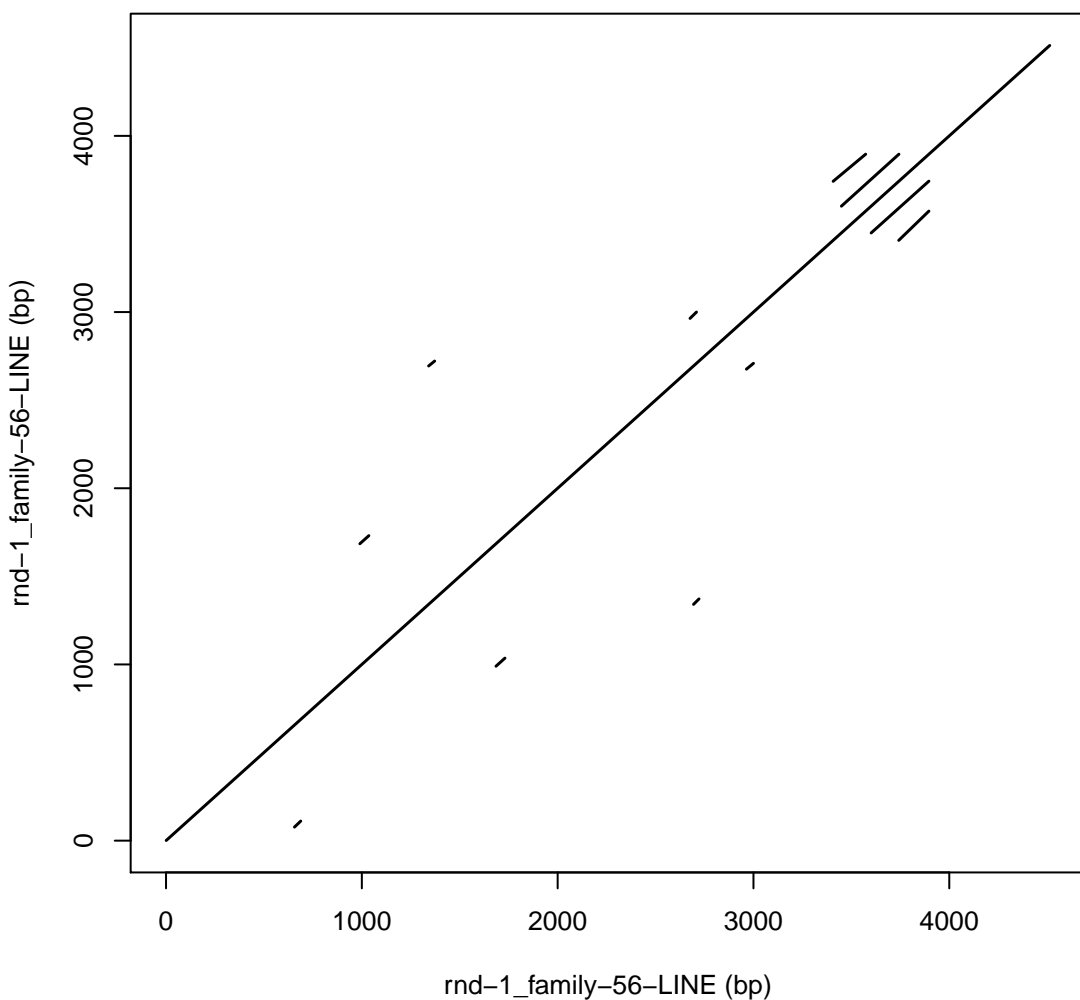
TE: rnd-1_family-56-LINE
consensus size: 4513bp; fragments: 254; full length: 0 (>=4061.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

