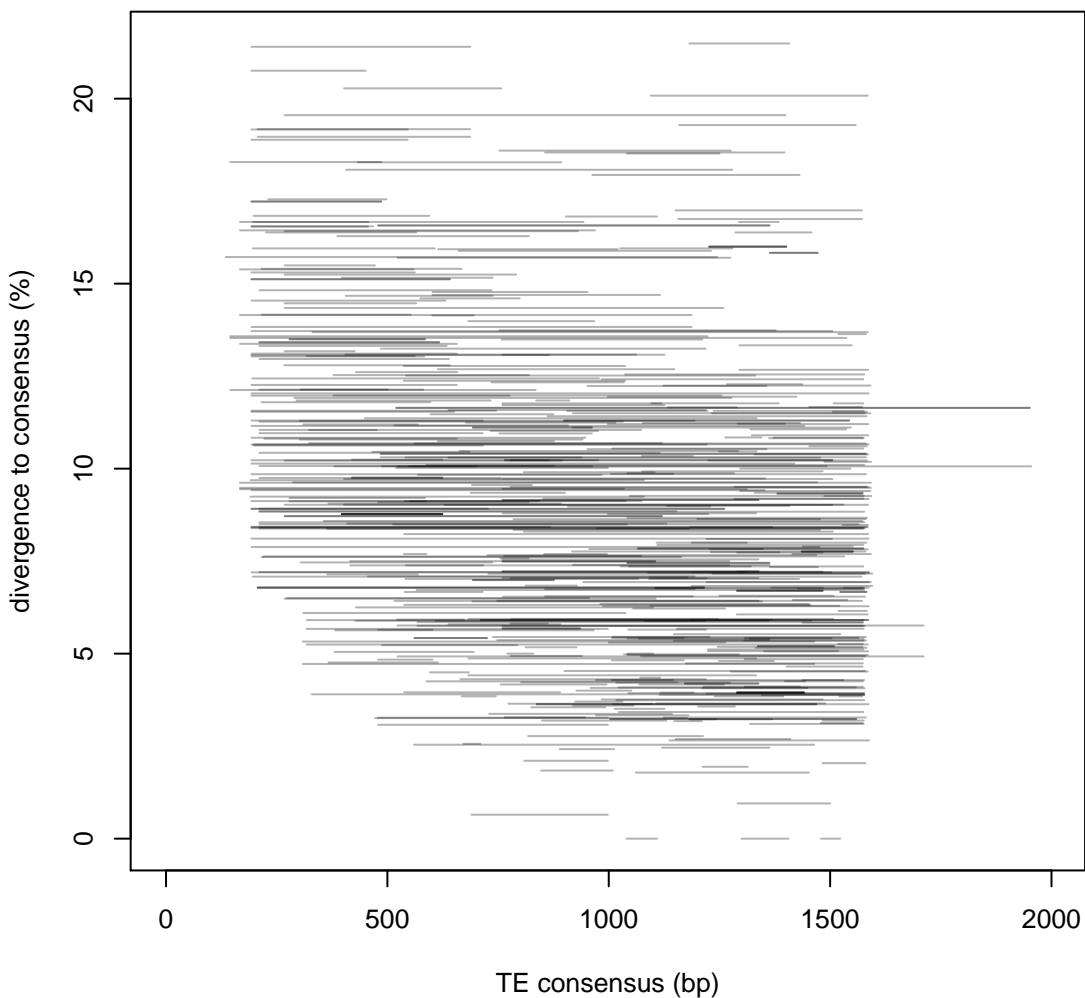
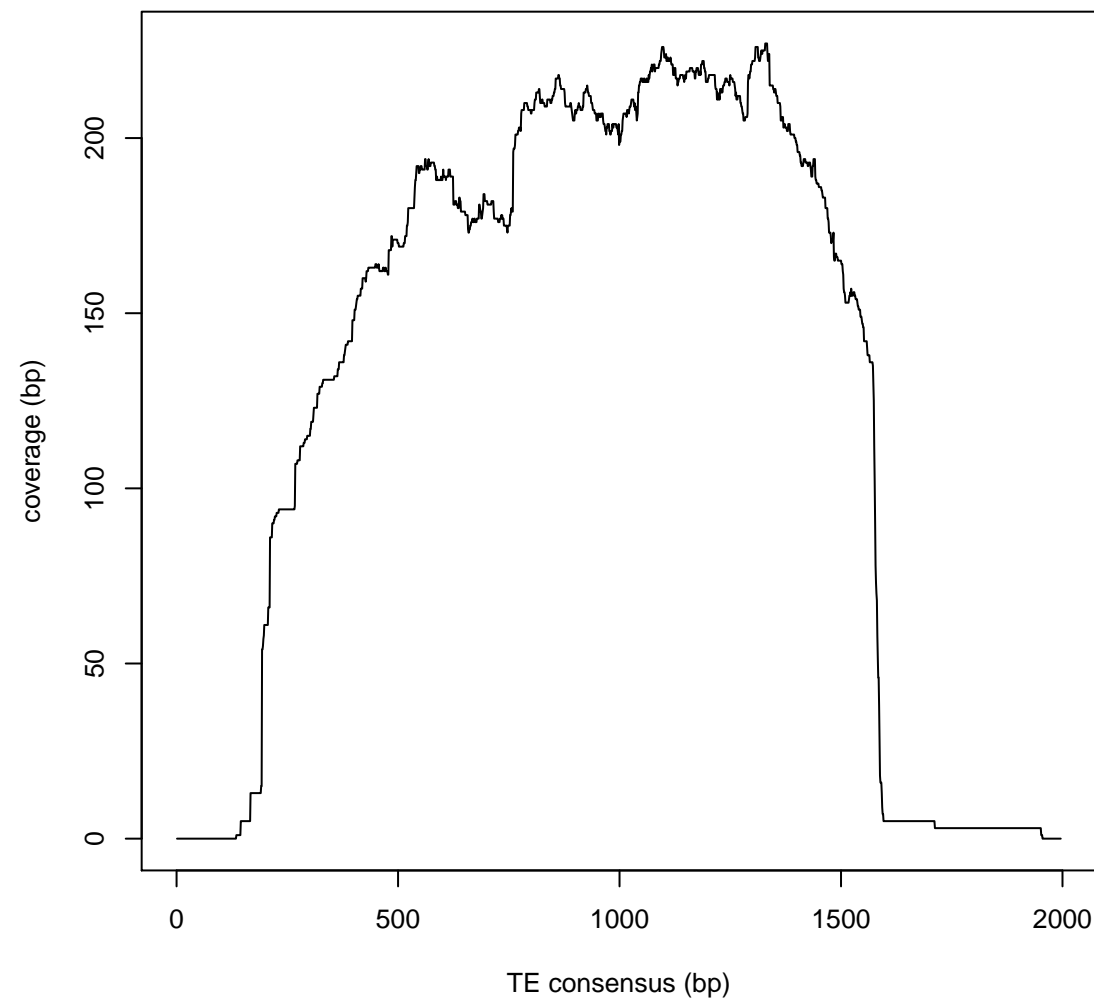


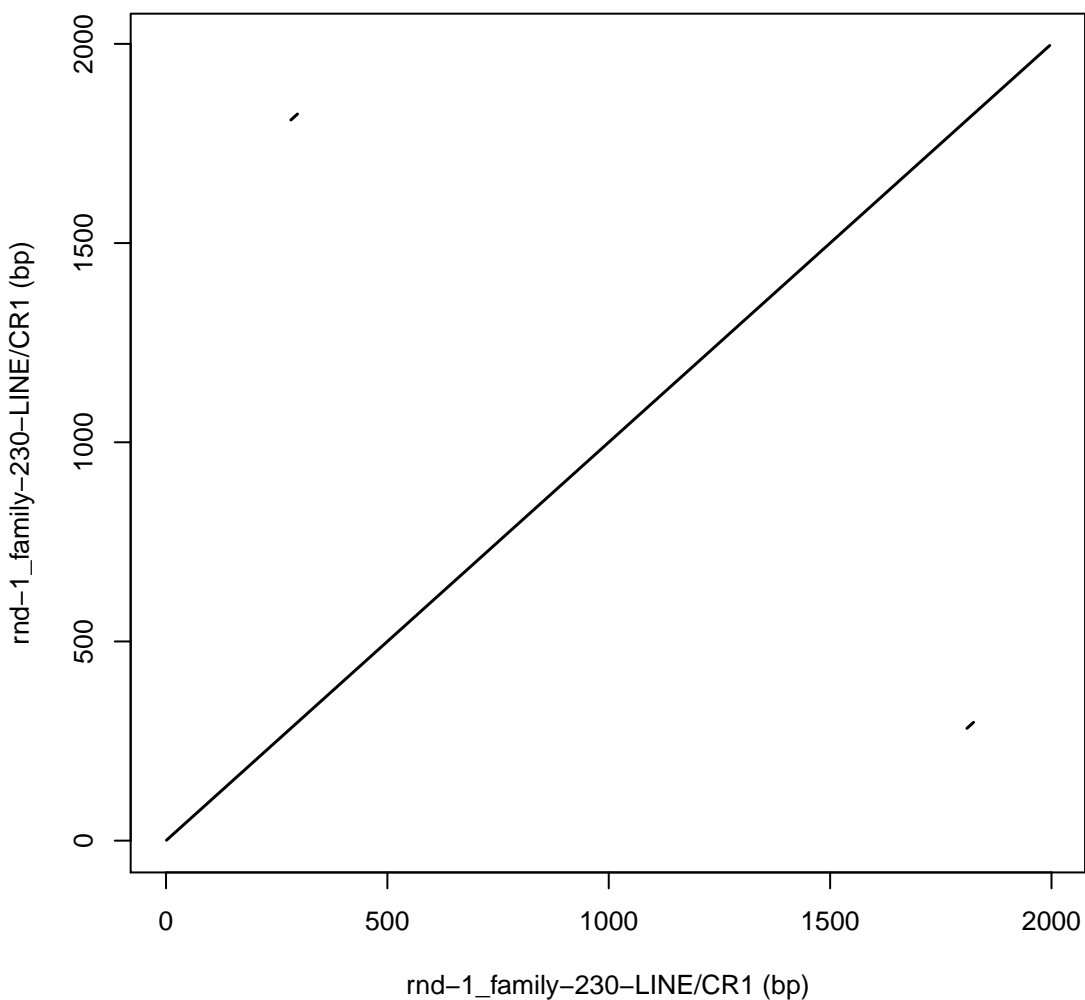
TE: rnd-1_family-230-LINE/CR1
consensus size: 1996bp; fragments: 528; full length: 0 (>=1796.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

