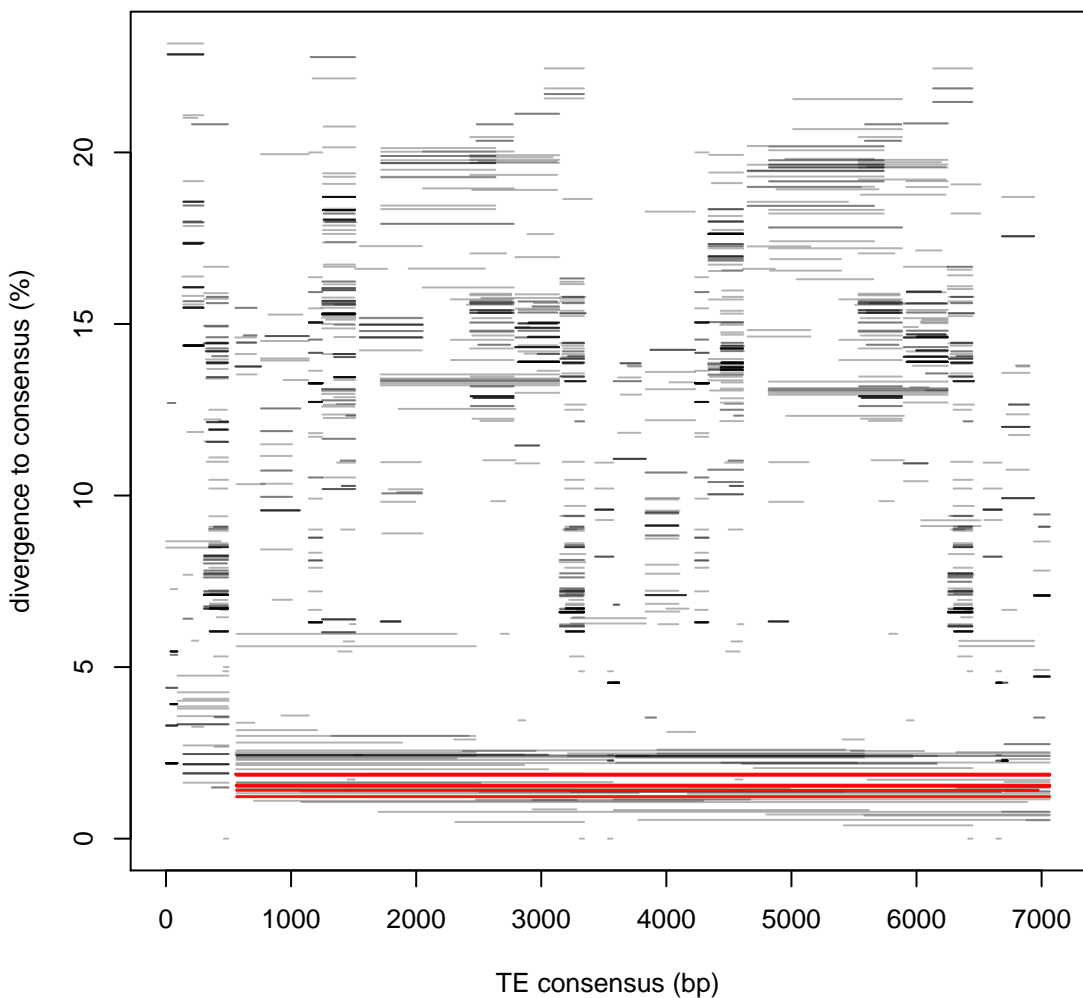
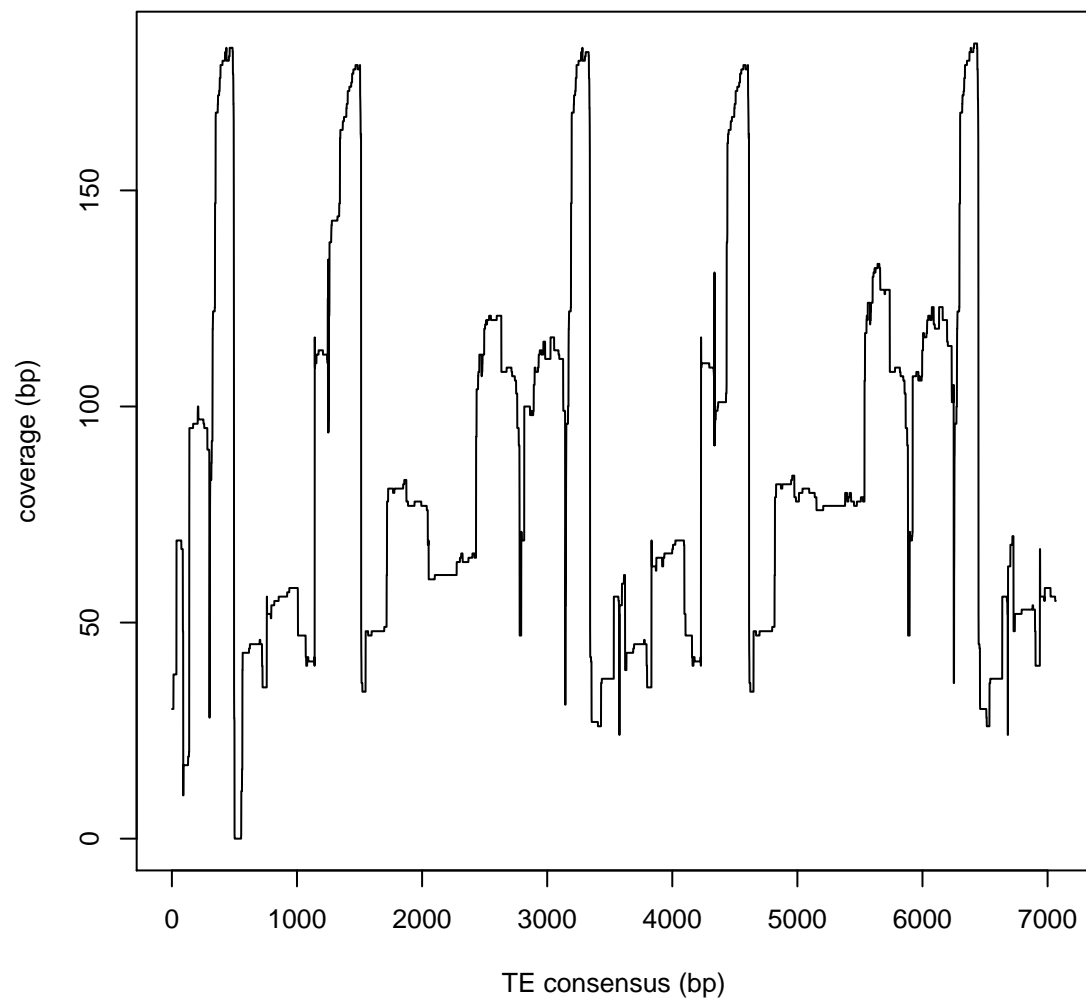


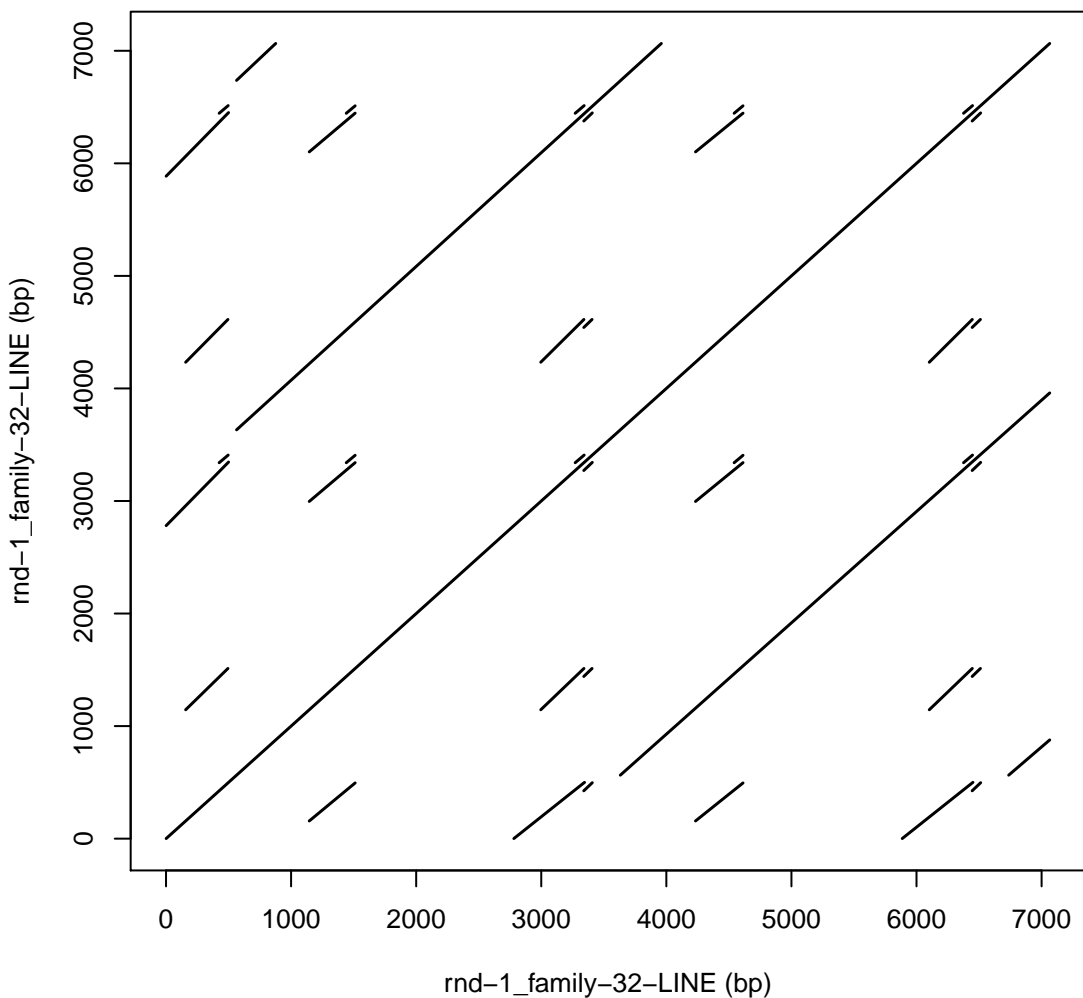
TE: rnd-1_family-32-LINE
consensus size: 7065bp; fragments: 1813; full length: 6 (>=6358.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

