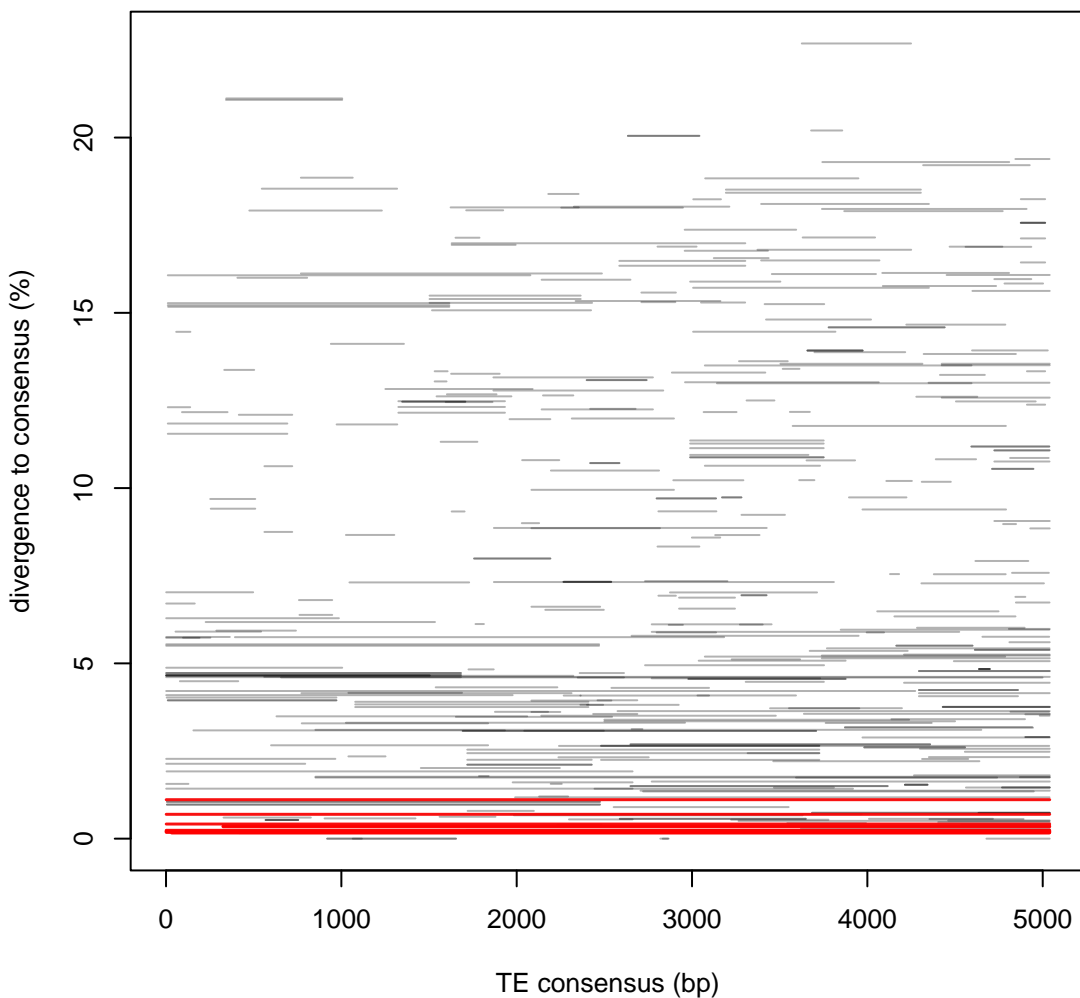
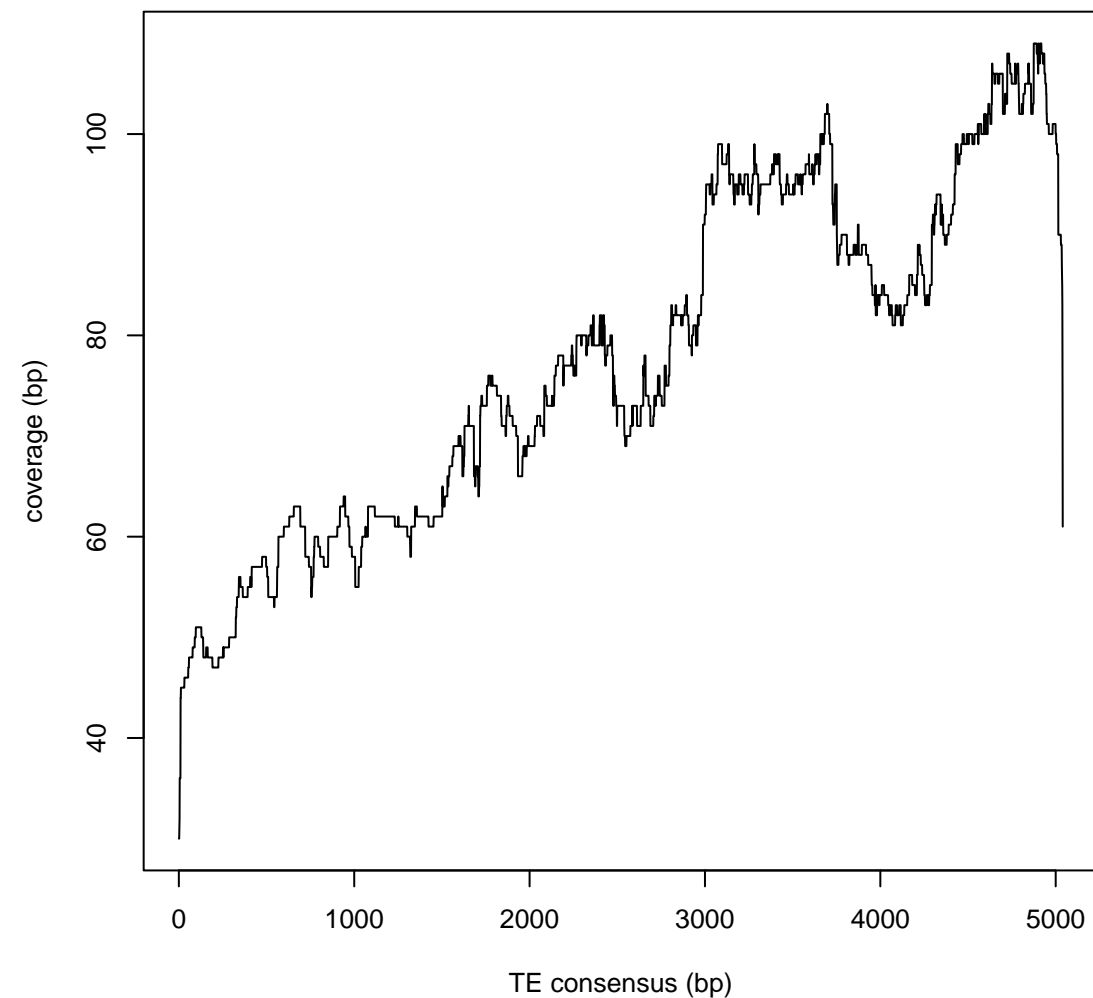


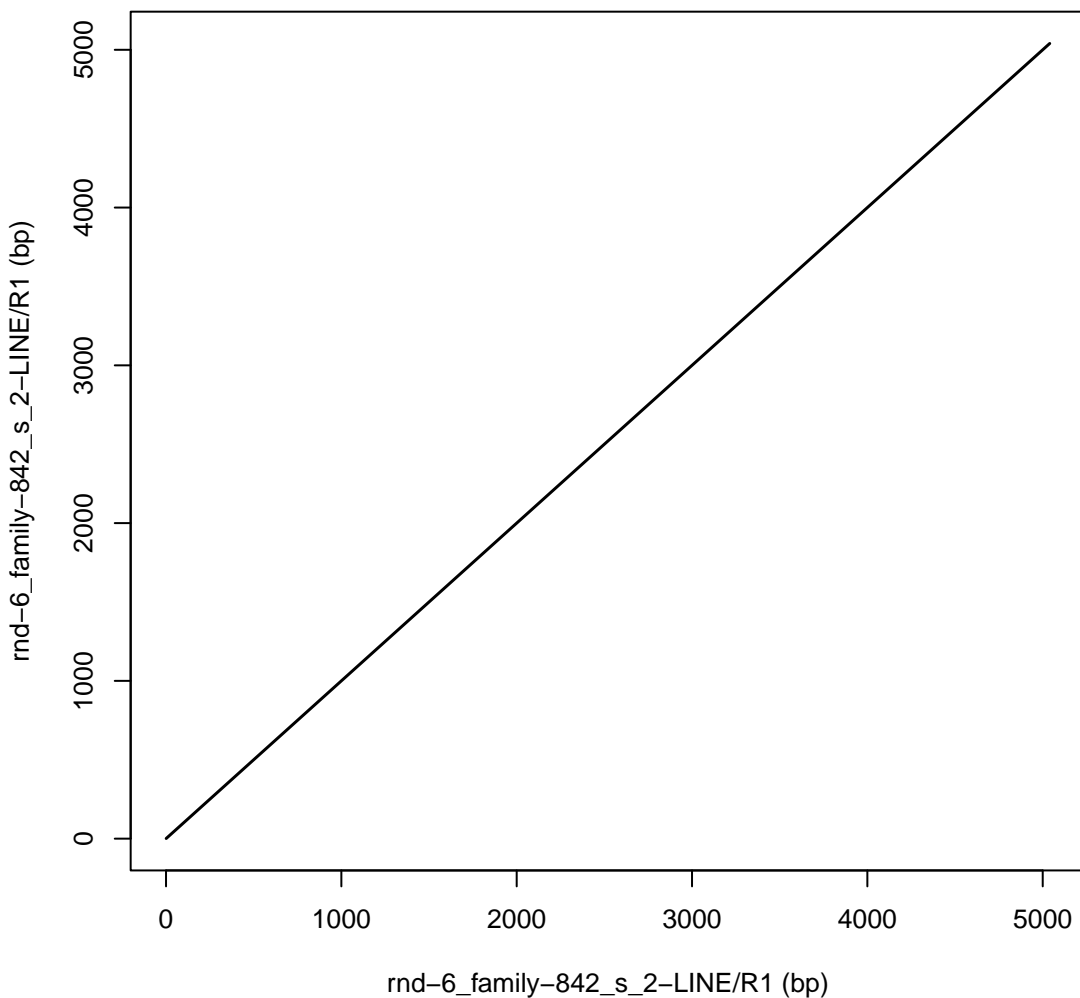
TE: rnd-6_family-842_s_2-LINE/R1
consensus size: 5040bp; fragments: 437; full length: 12 (>=4536bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

