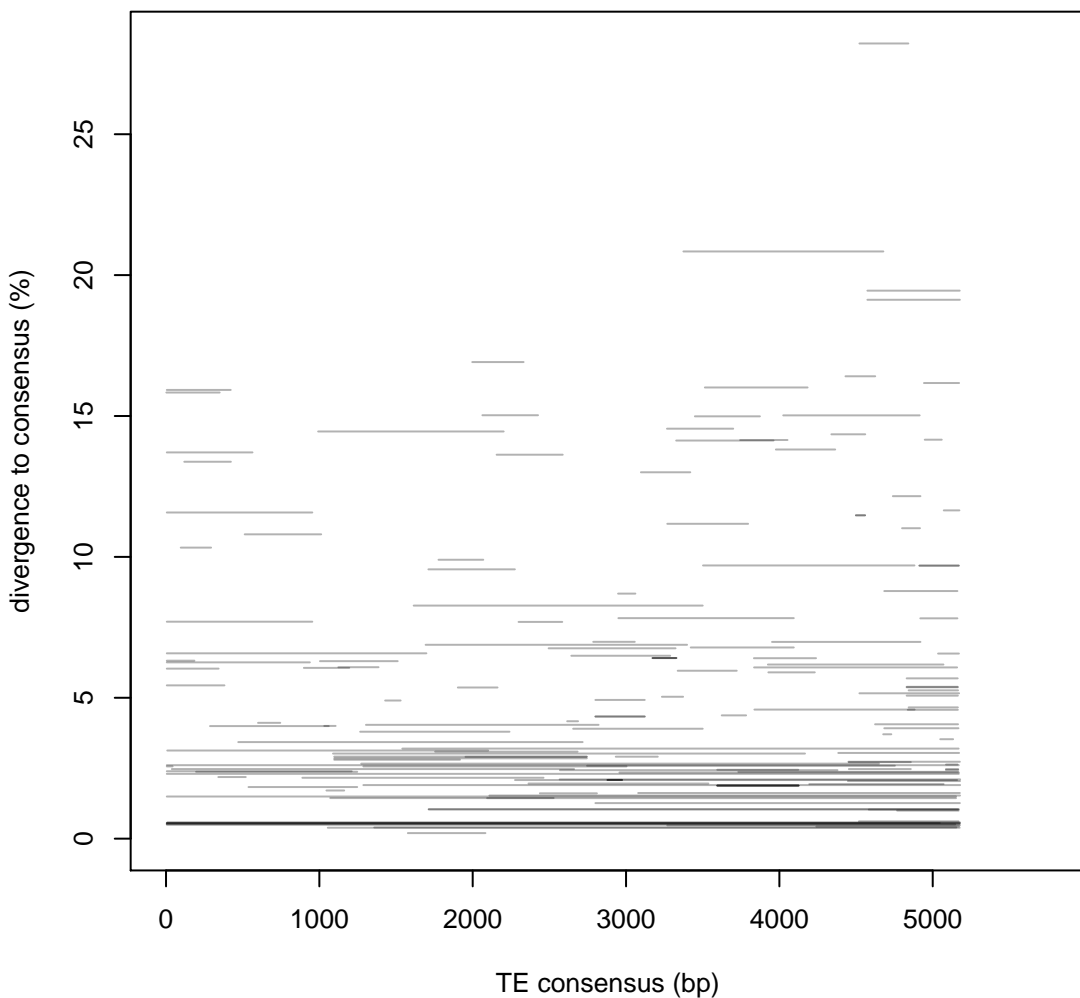
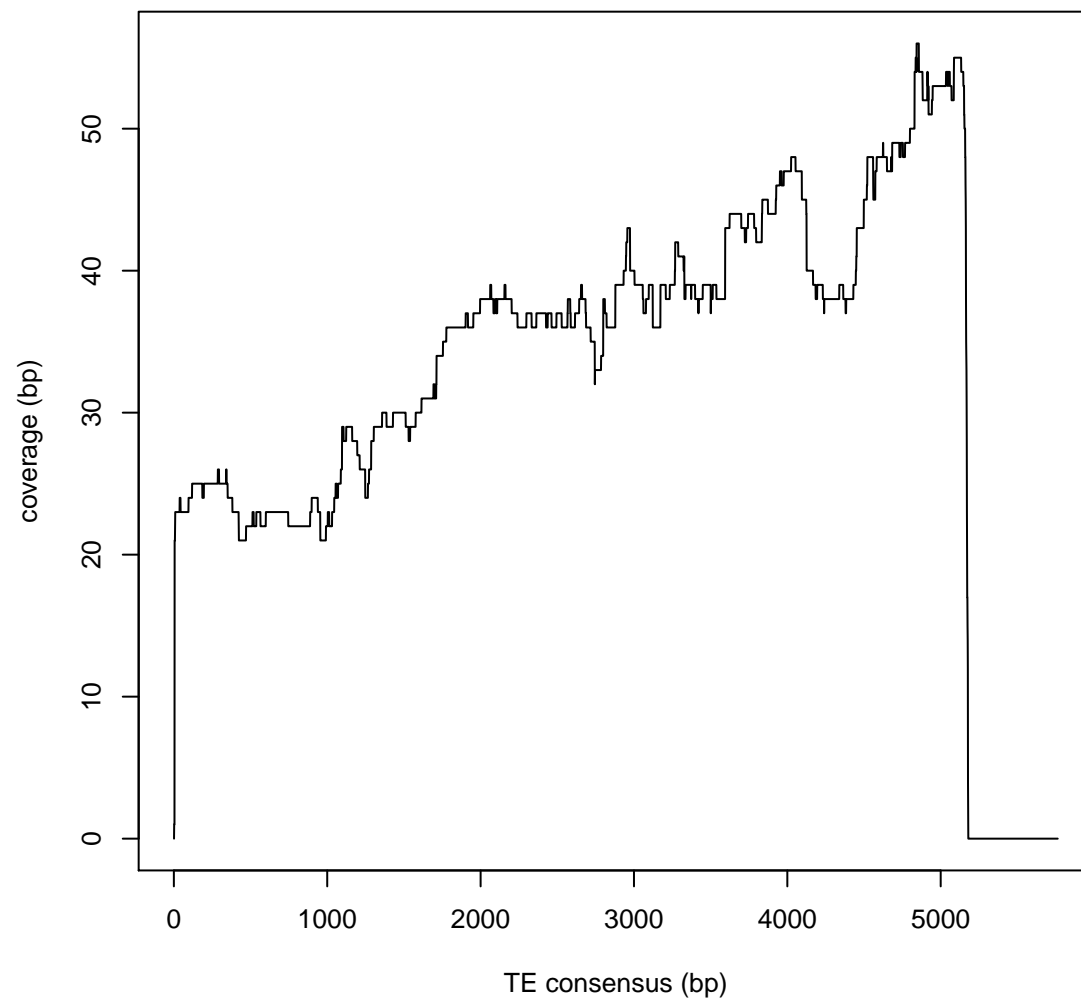


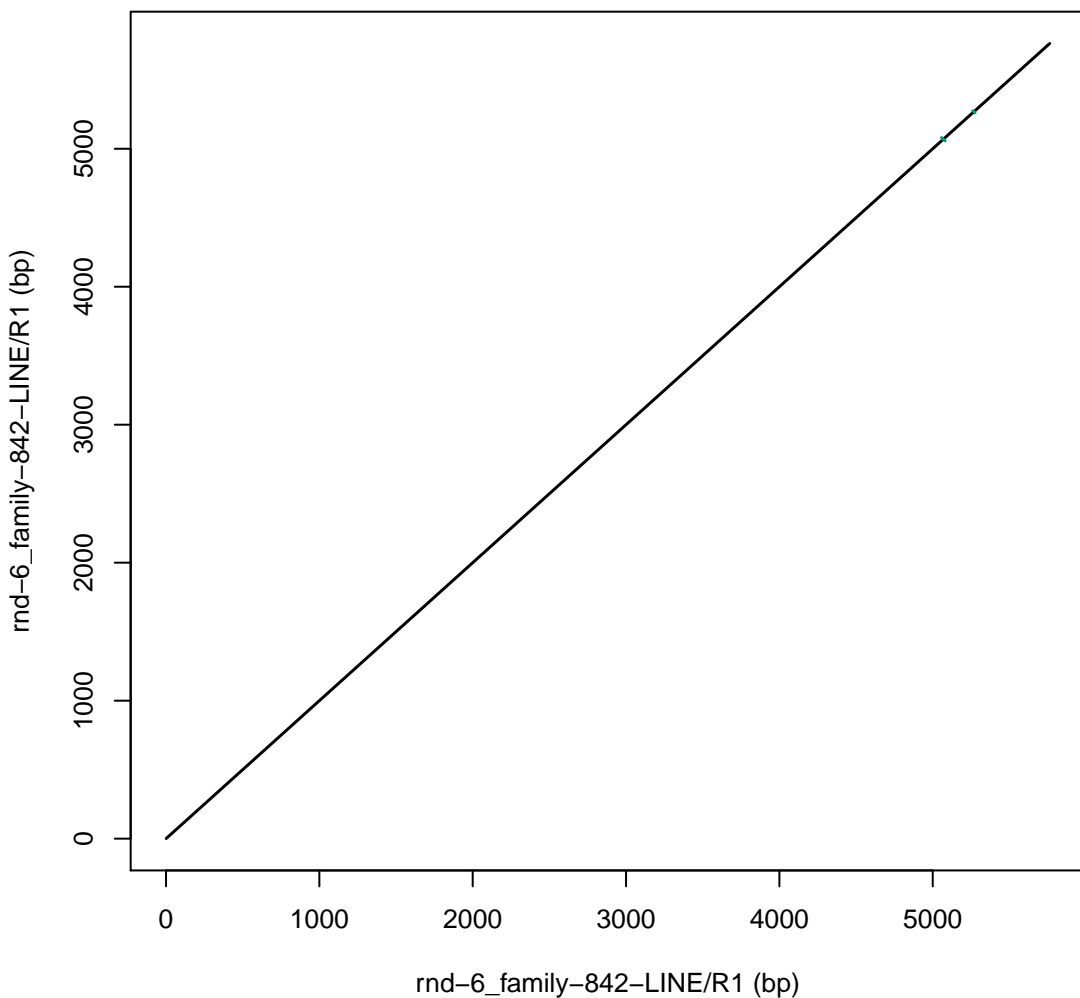
TE: rnd-6_family-842-LINE/R1
consensus size: 5763bp; fragments: 166; full length: 0 (>=5186.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

