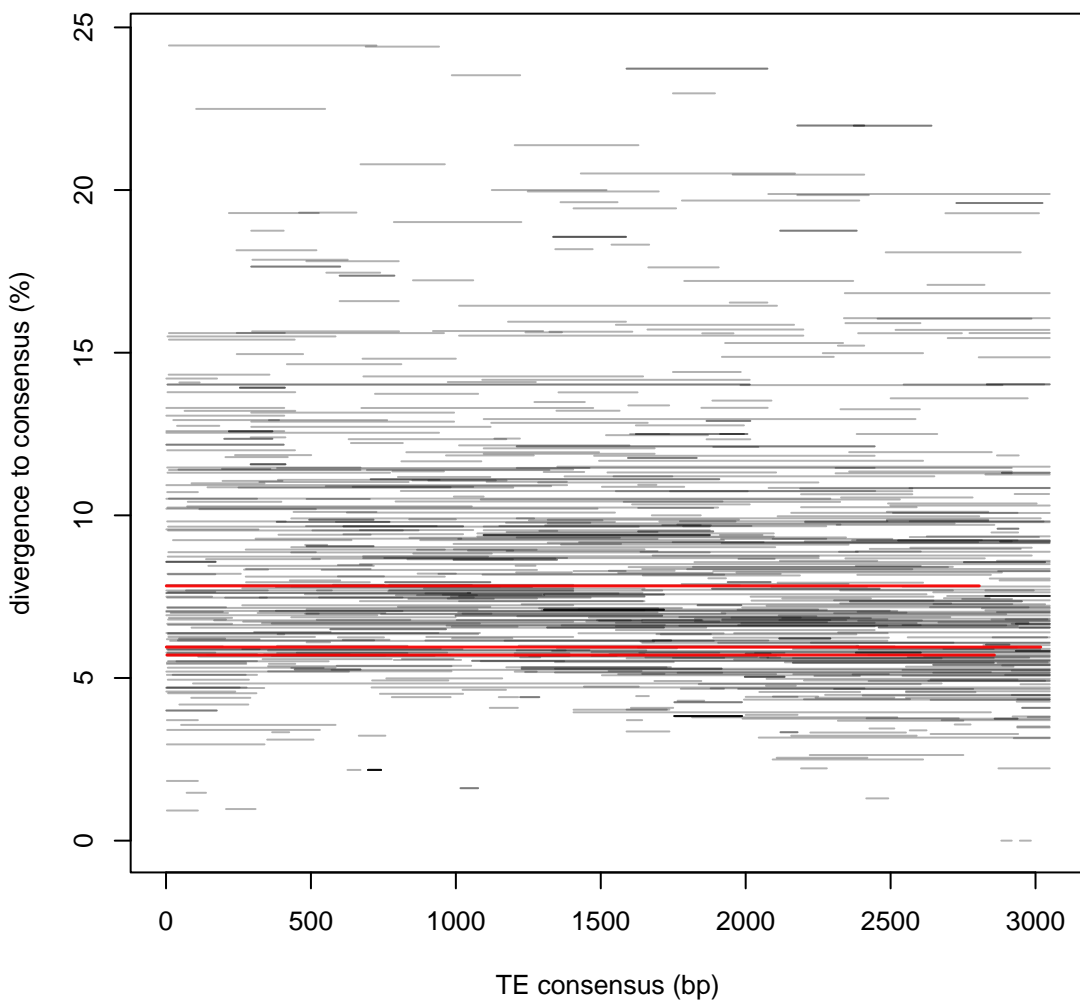
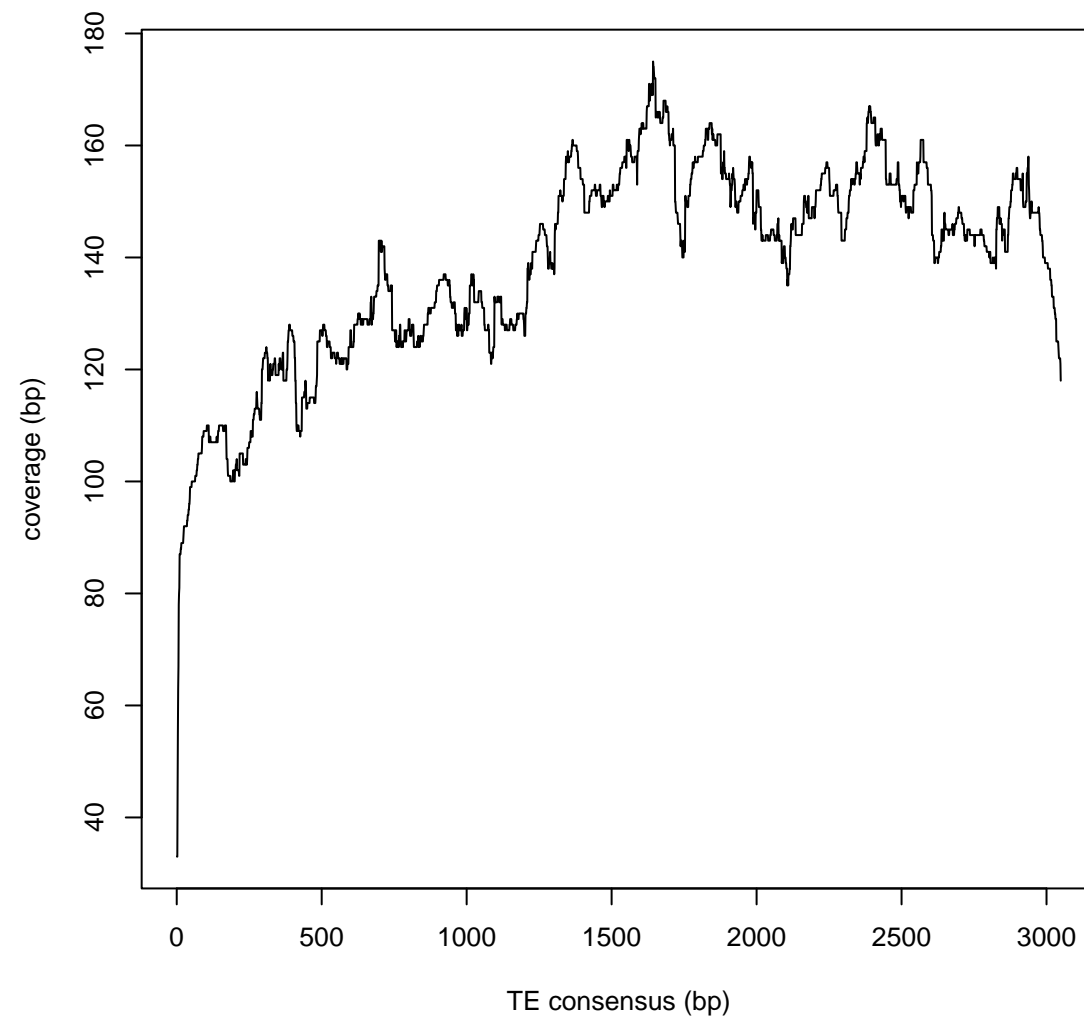


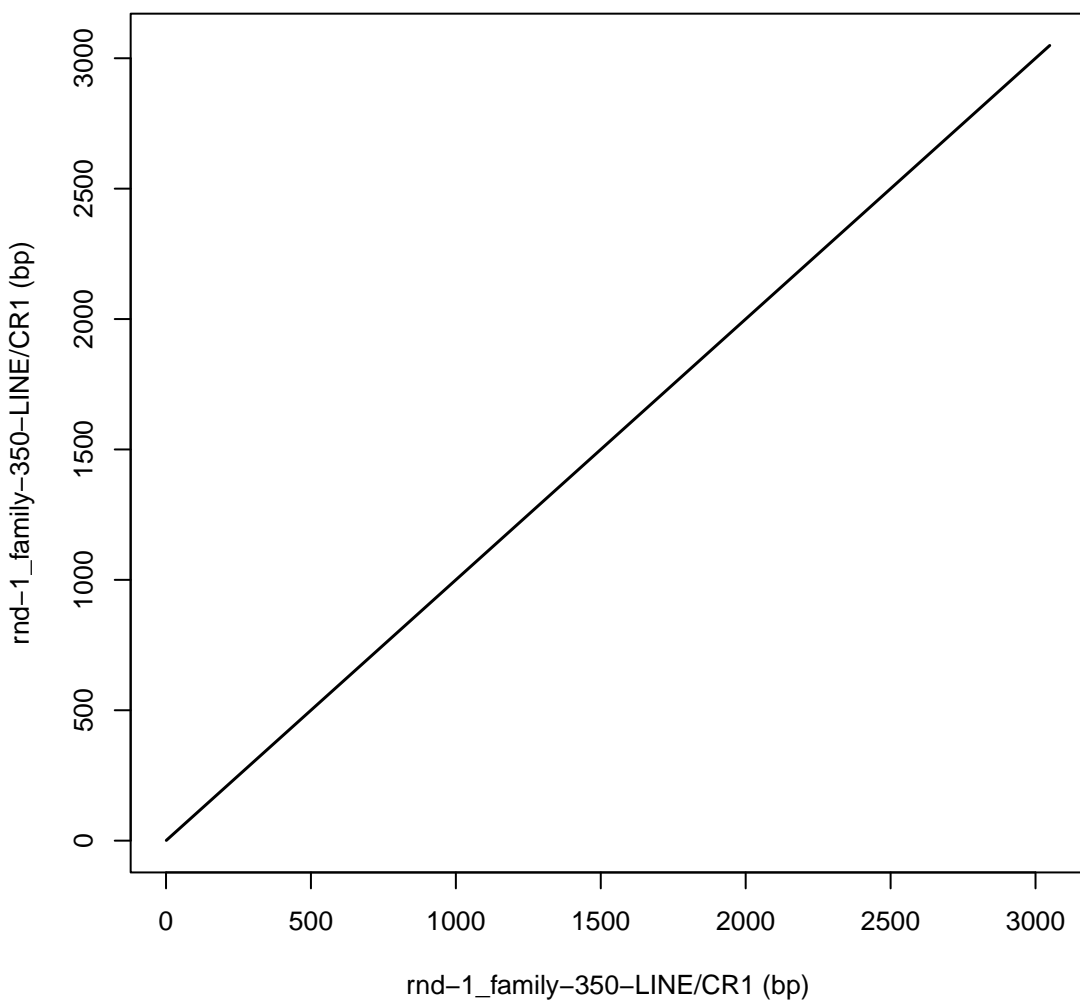
TE: rnd-1_family-350-LINE/CR1
consensus size: 3049bp; fragments: 809; full length: 3 (>=2744.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

