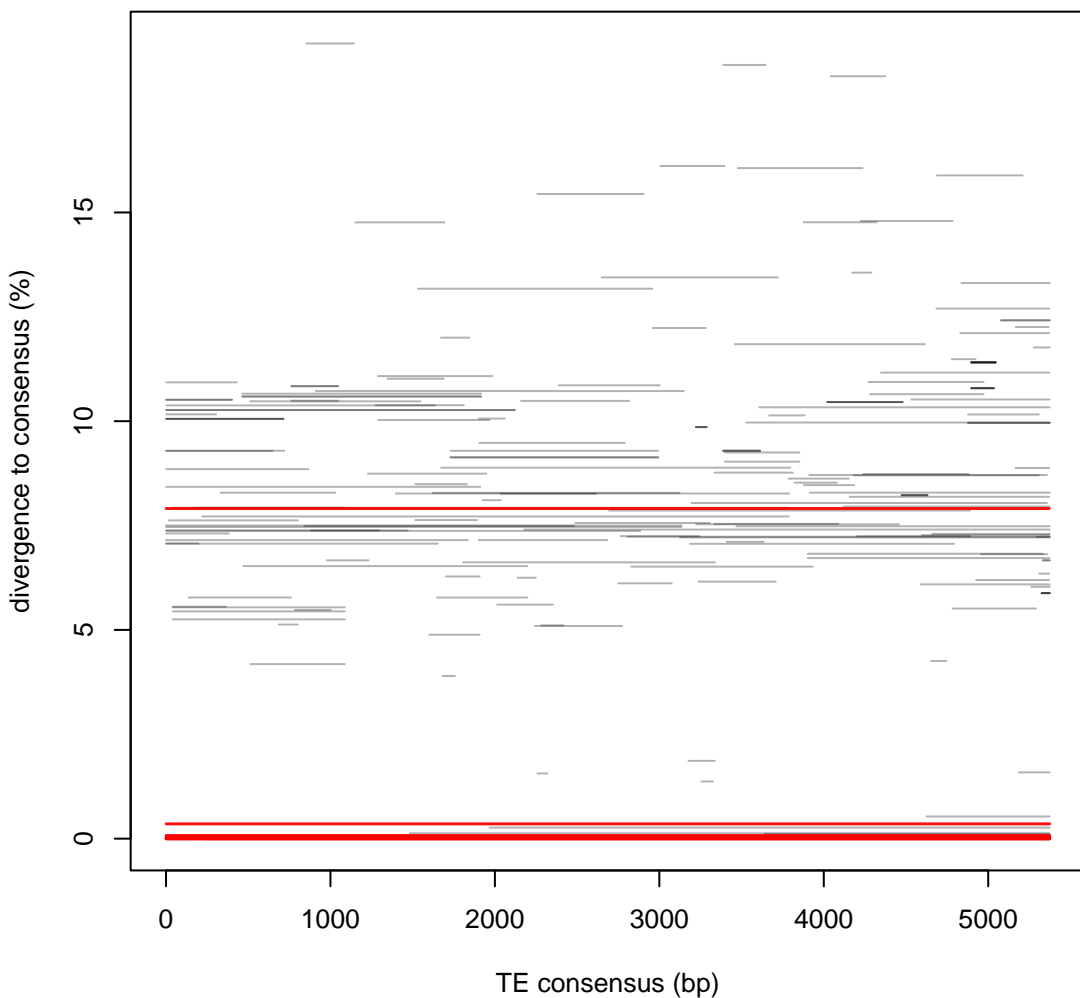
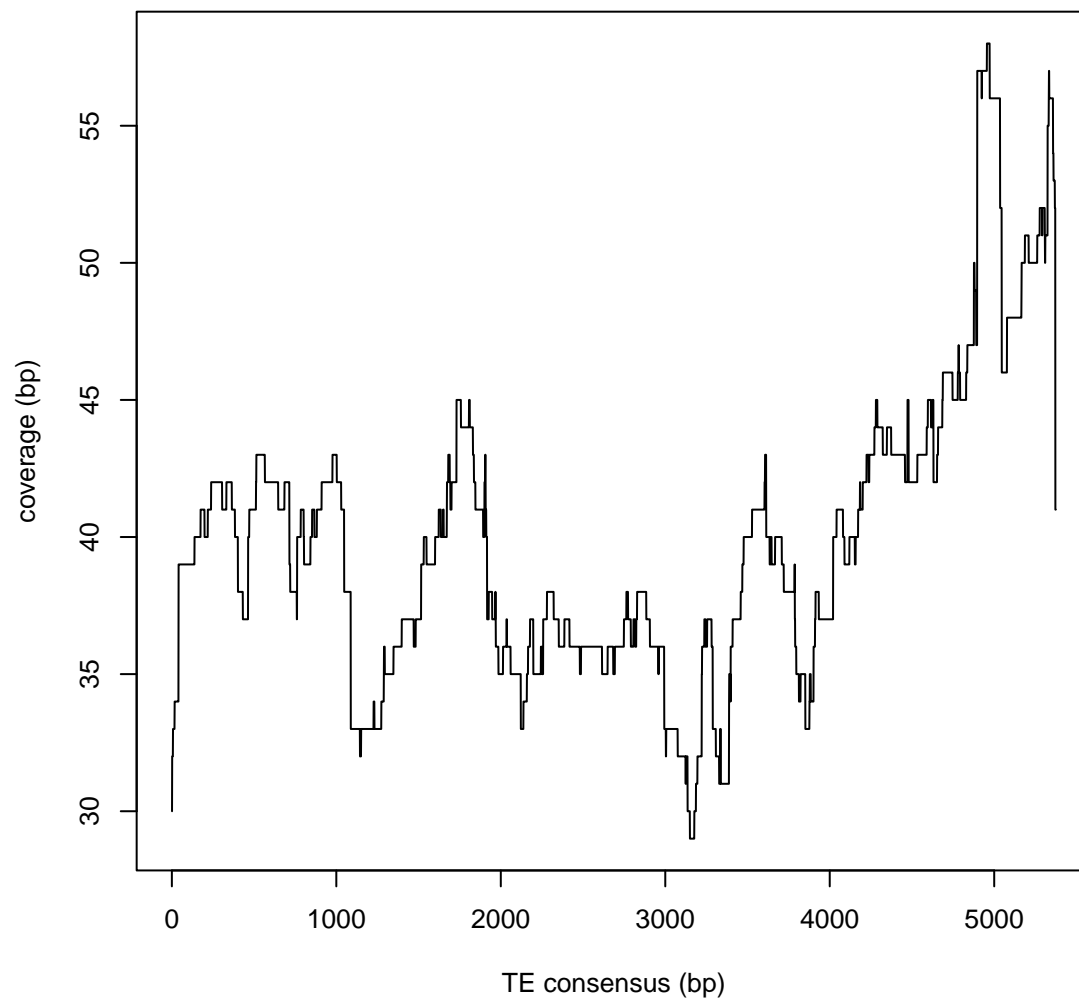


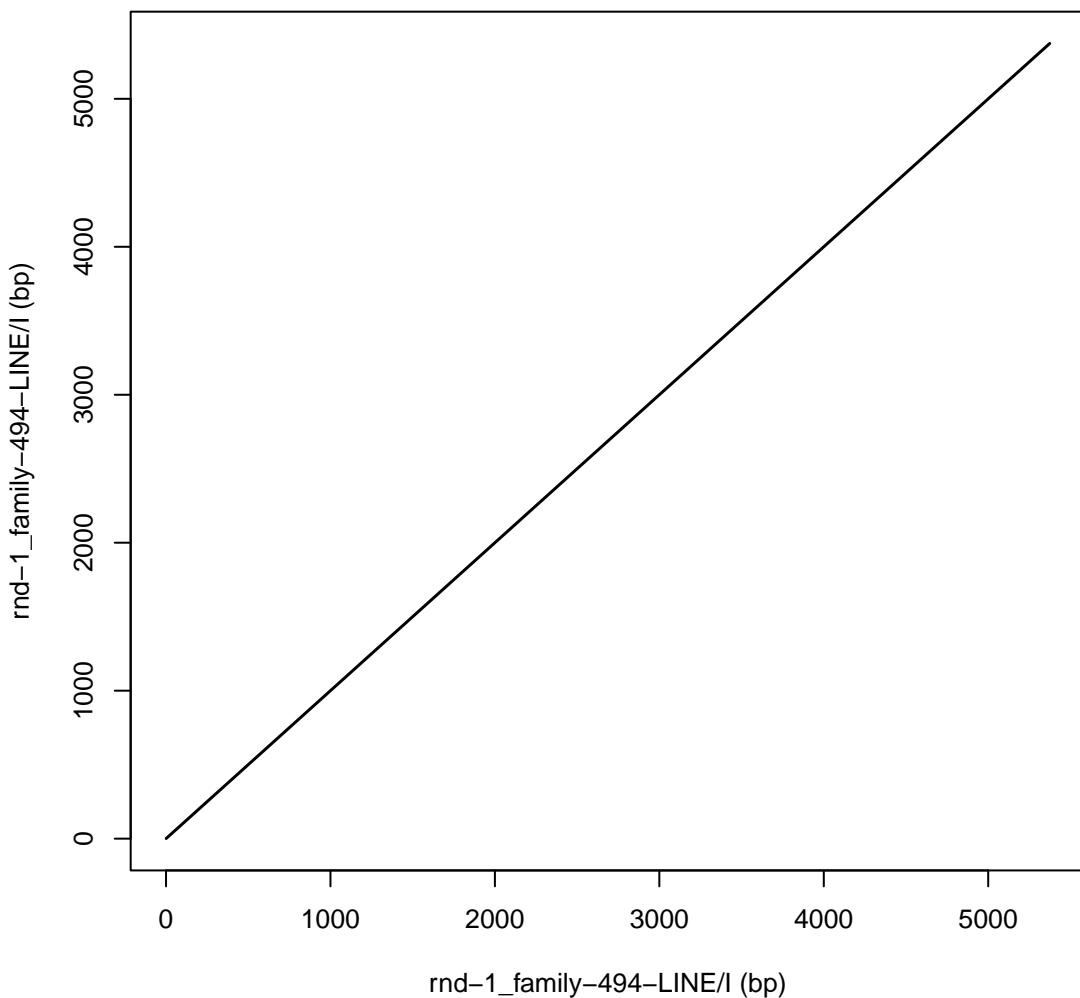
TE: rnd-1_family-494-LINE/I
consensus size: 5374bp; fragments: 202; full length: 10 (>=4836.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

