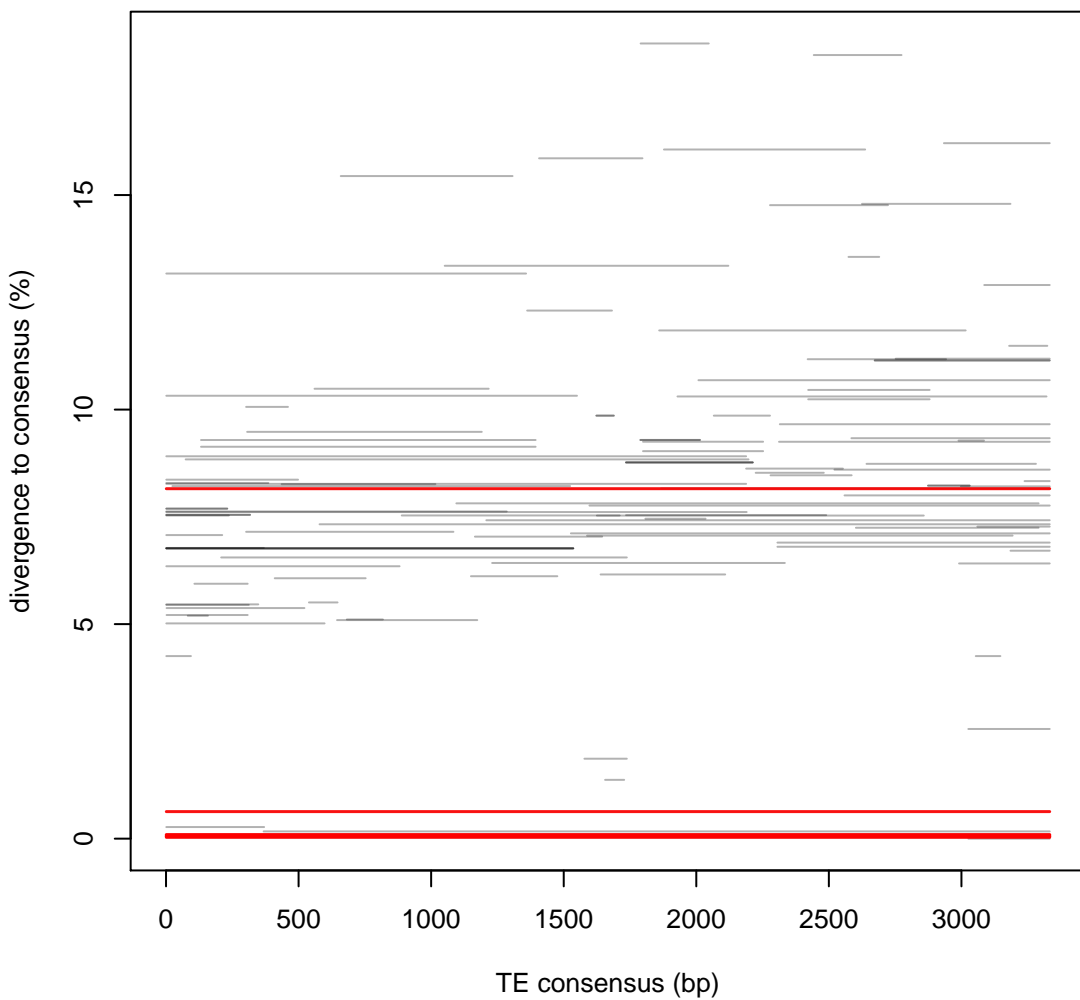
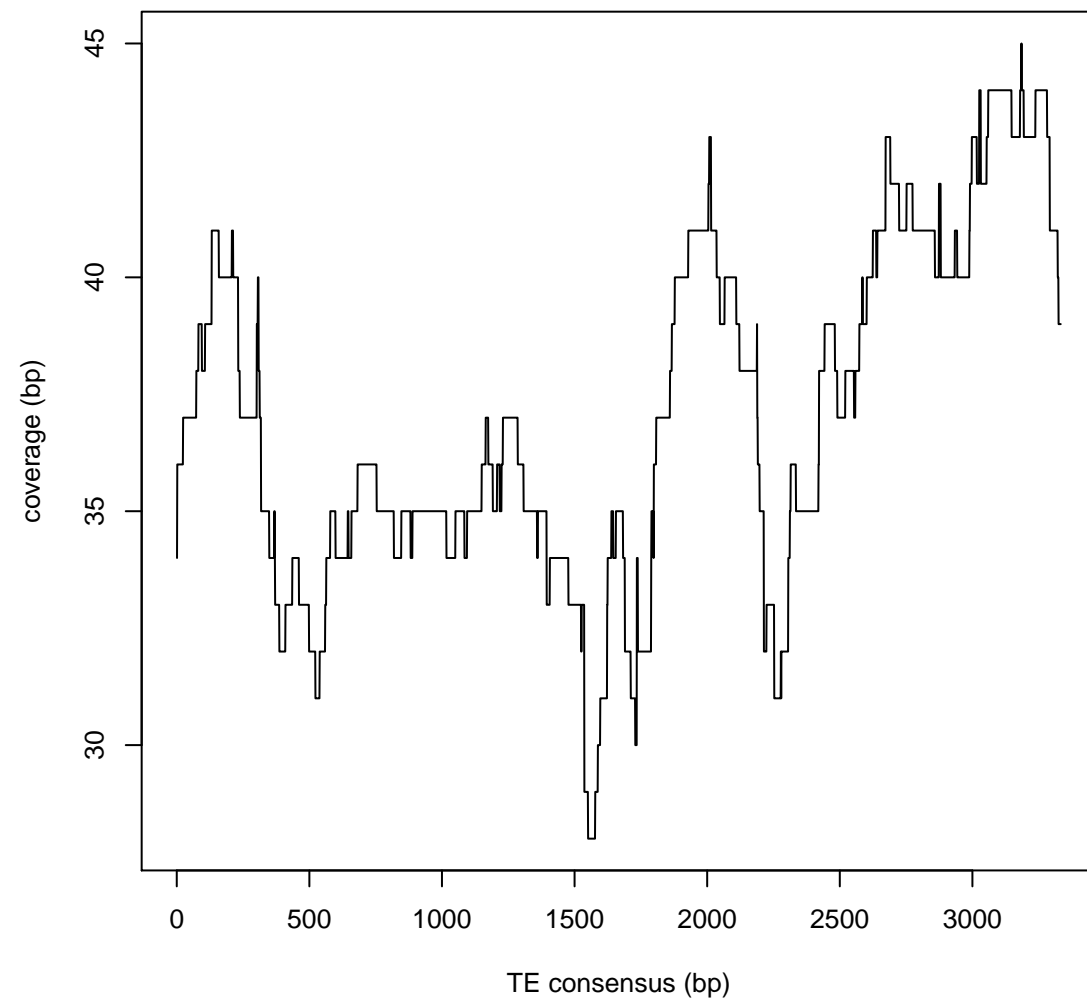


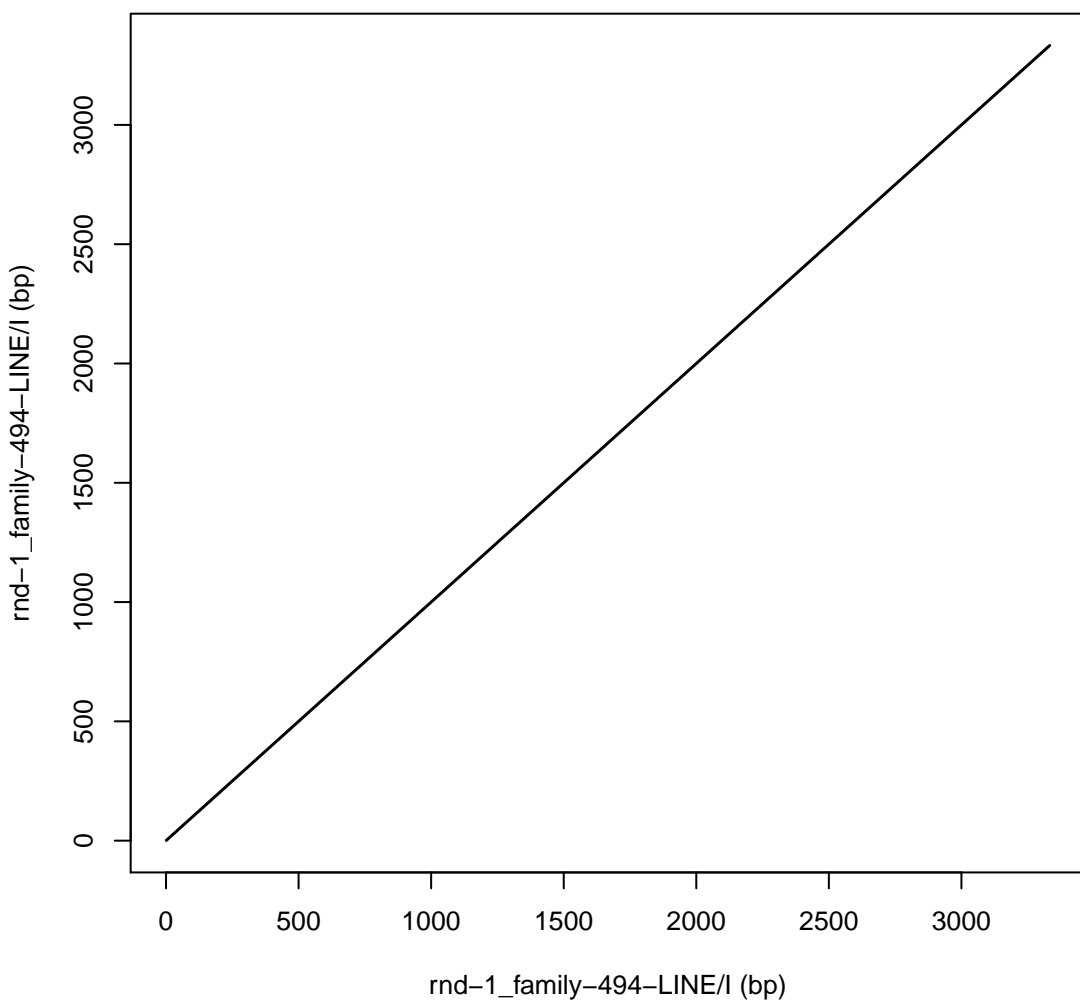
TE: rnd-1\_family-494-LINE/I  
consensus size: 3333bp; fragments: 129; full length: 9 (>=2999.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

