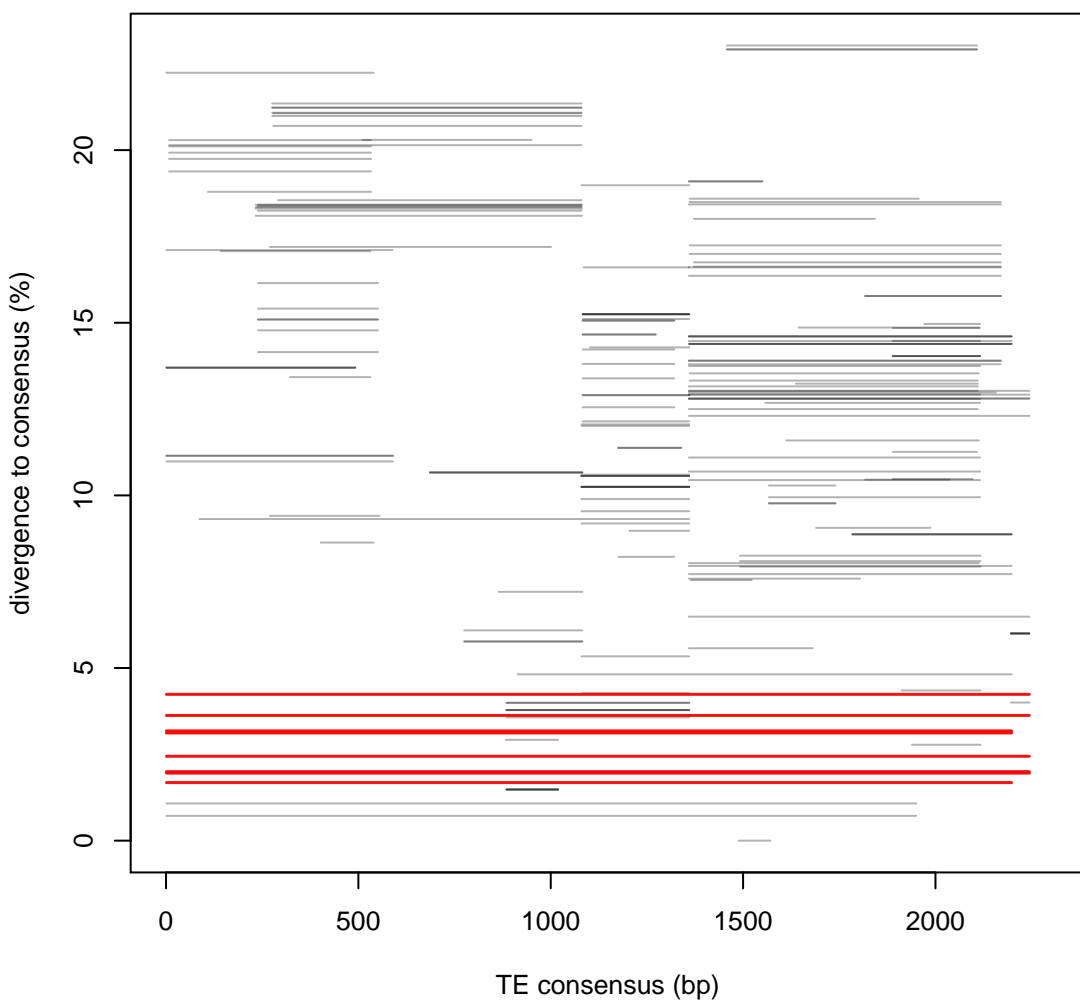
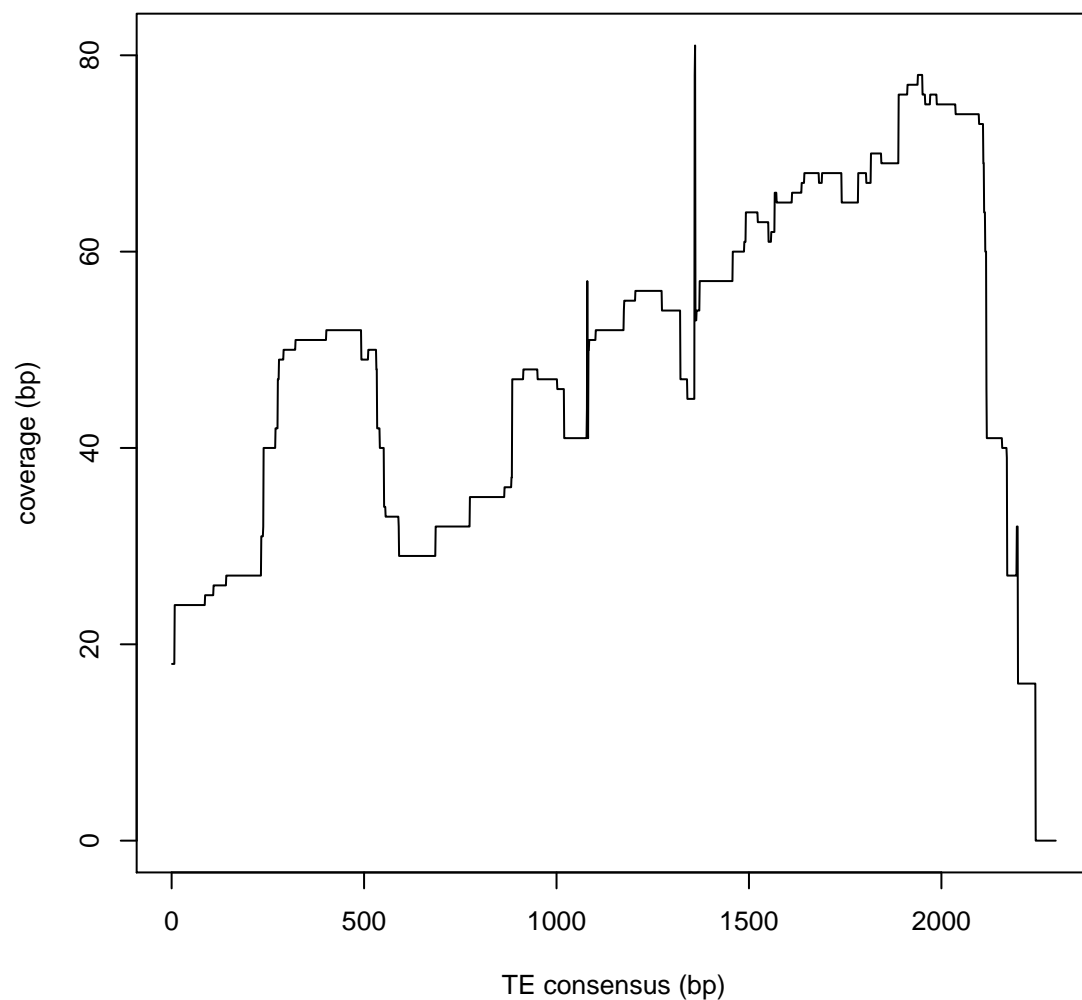


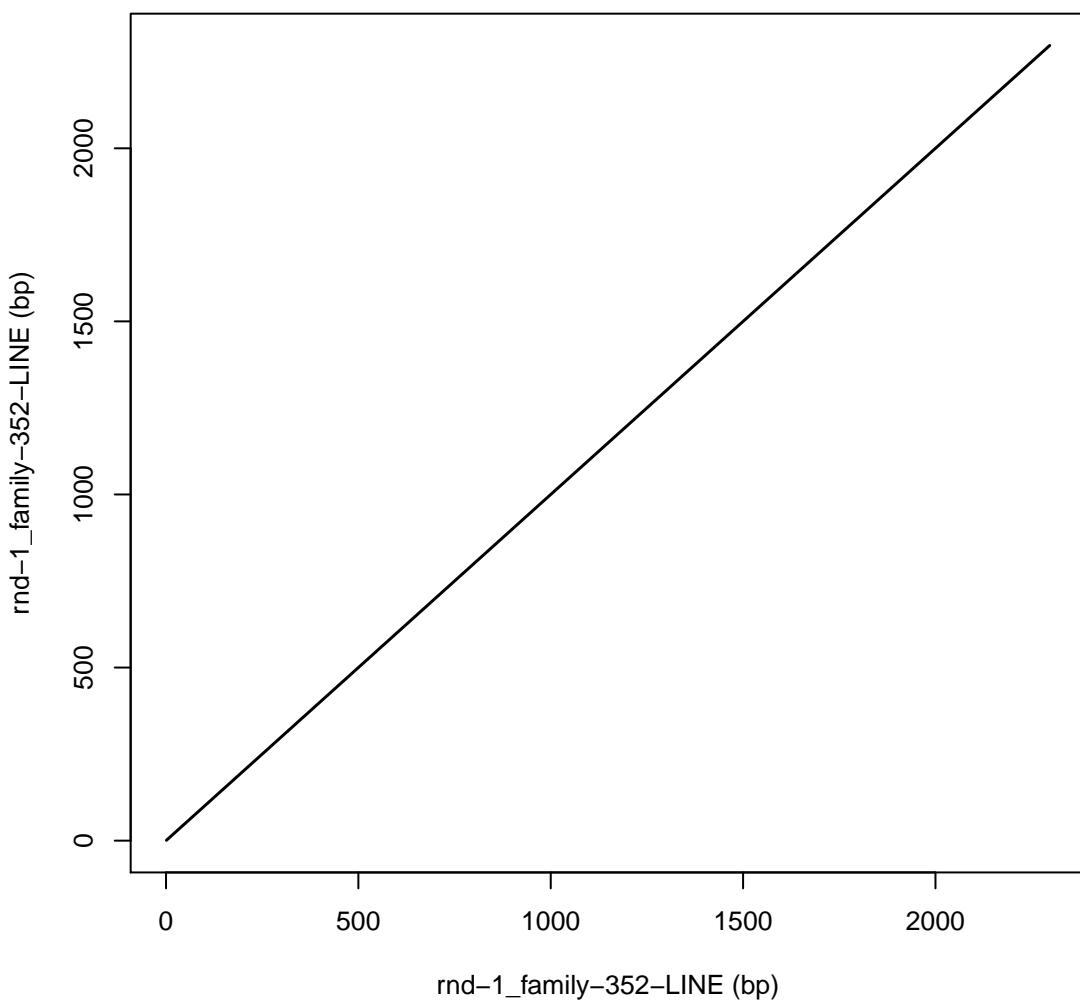
TE: rnd-1_family-352-LINE
consensus size: 2297bp; fragments: 193; full length: 8 (>=2067.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

