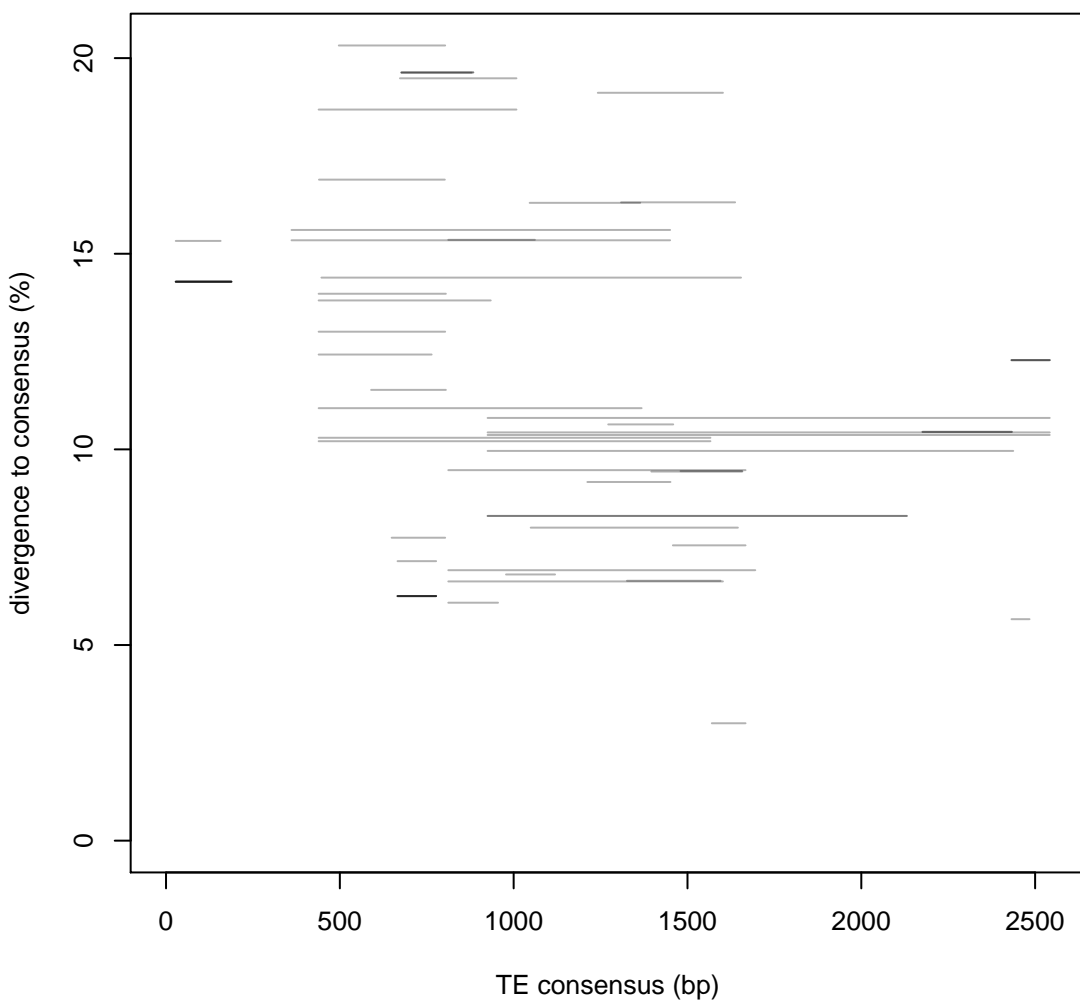
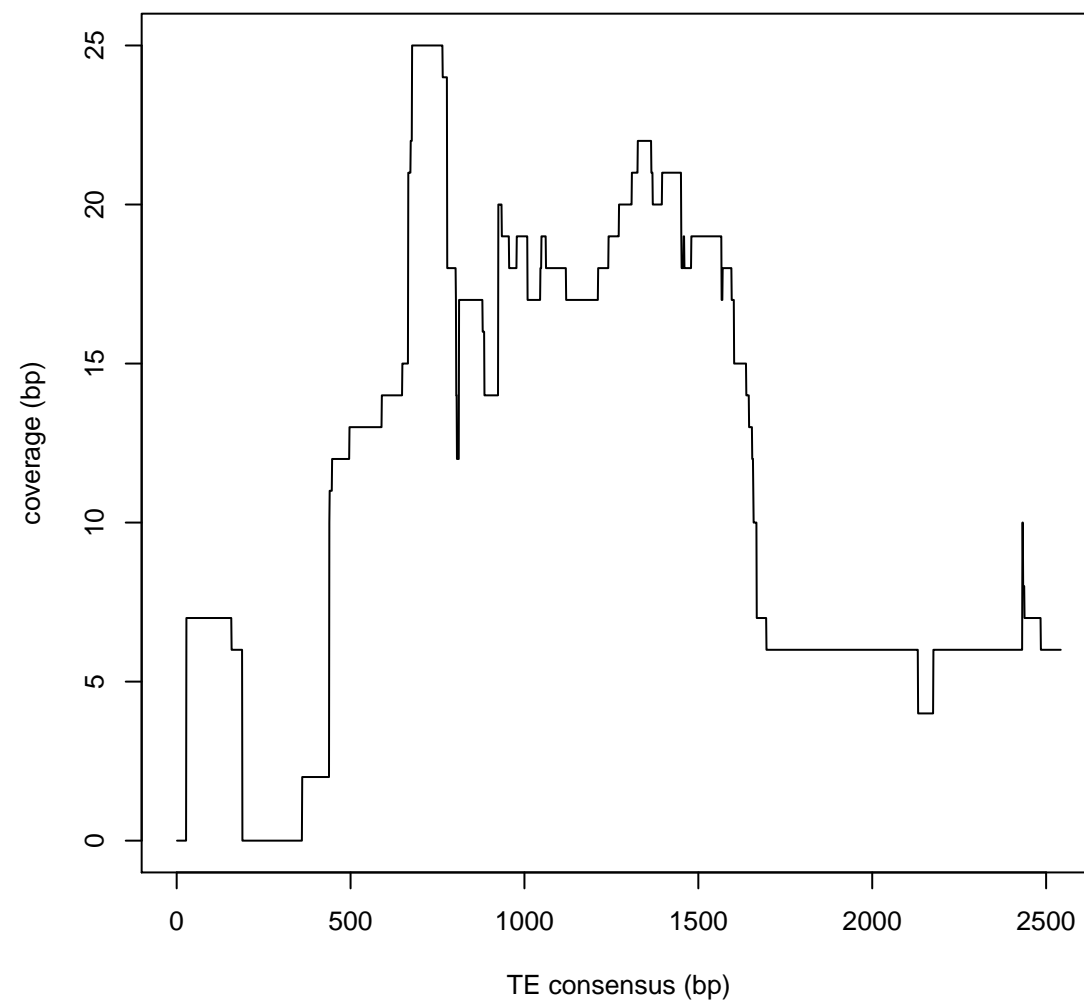


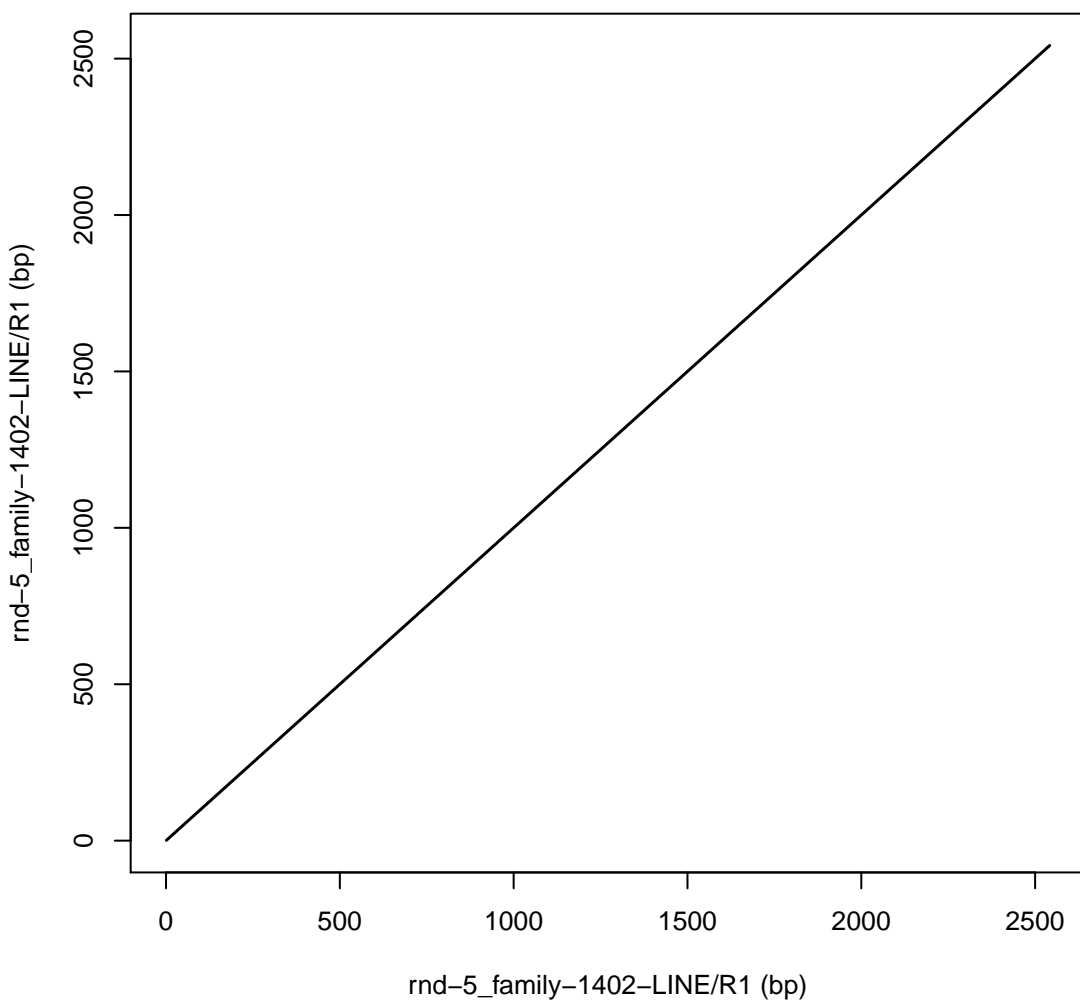
TE: rnd-5_family-1402-LINE/R1
consensus size: 2542bp; fragments: 61; full length: 0 (≥ 2287.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

