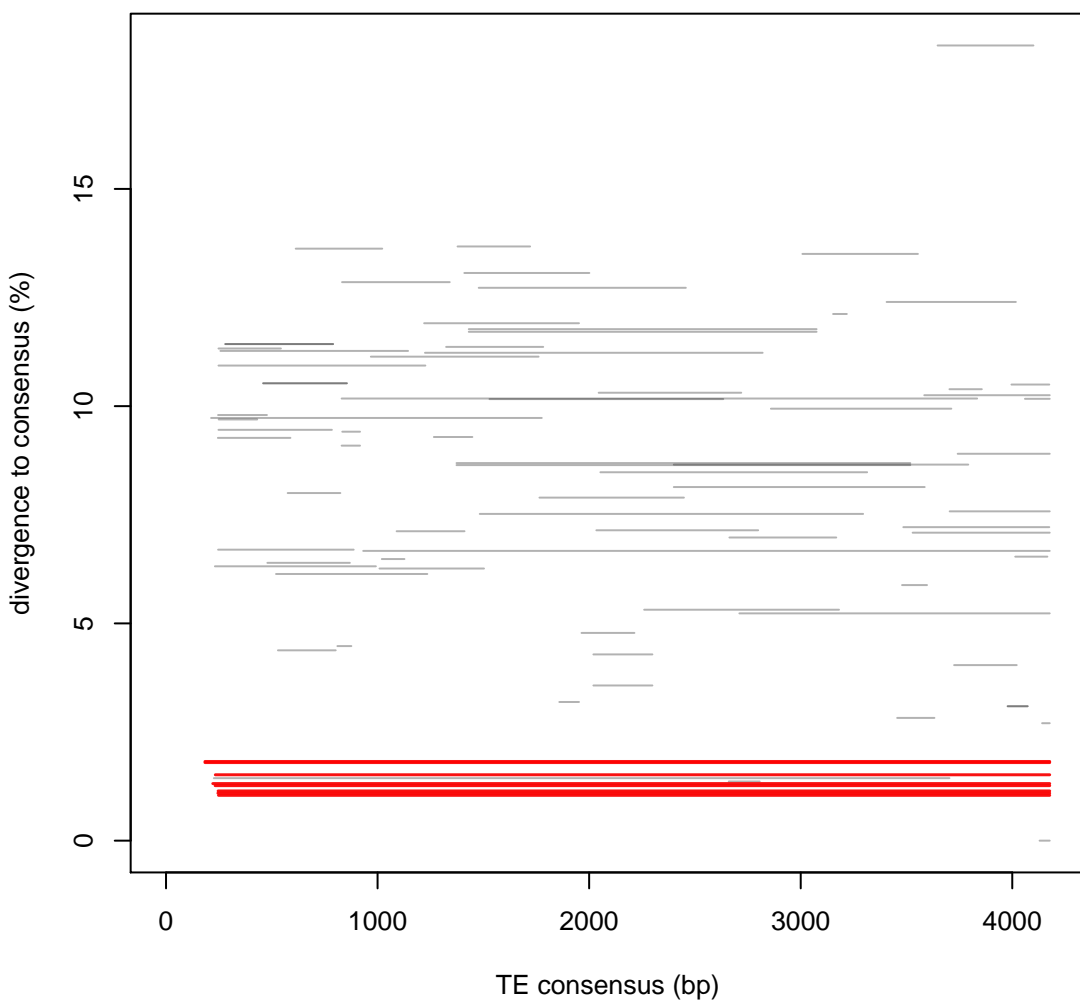
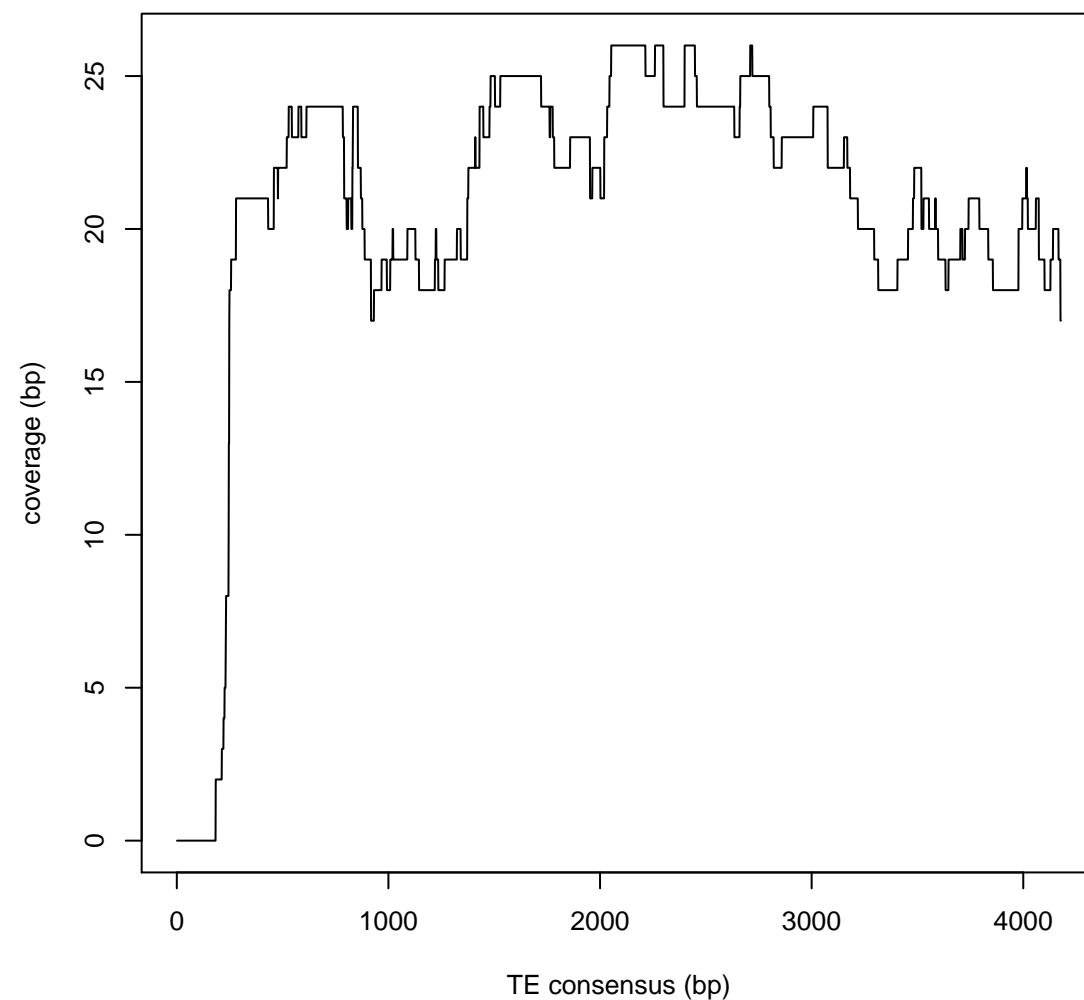


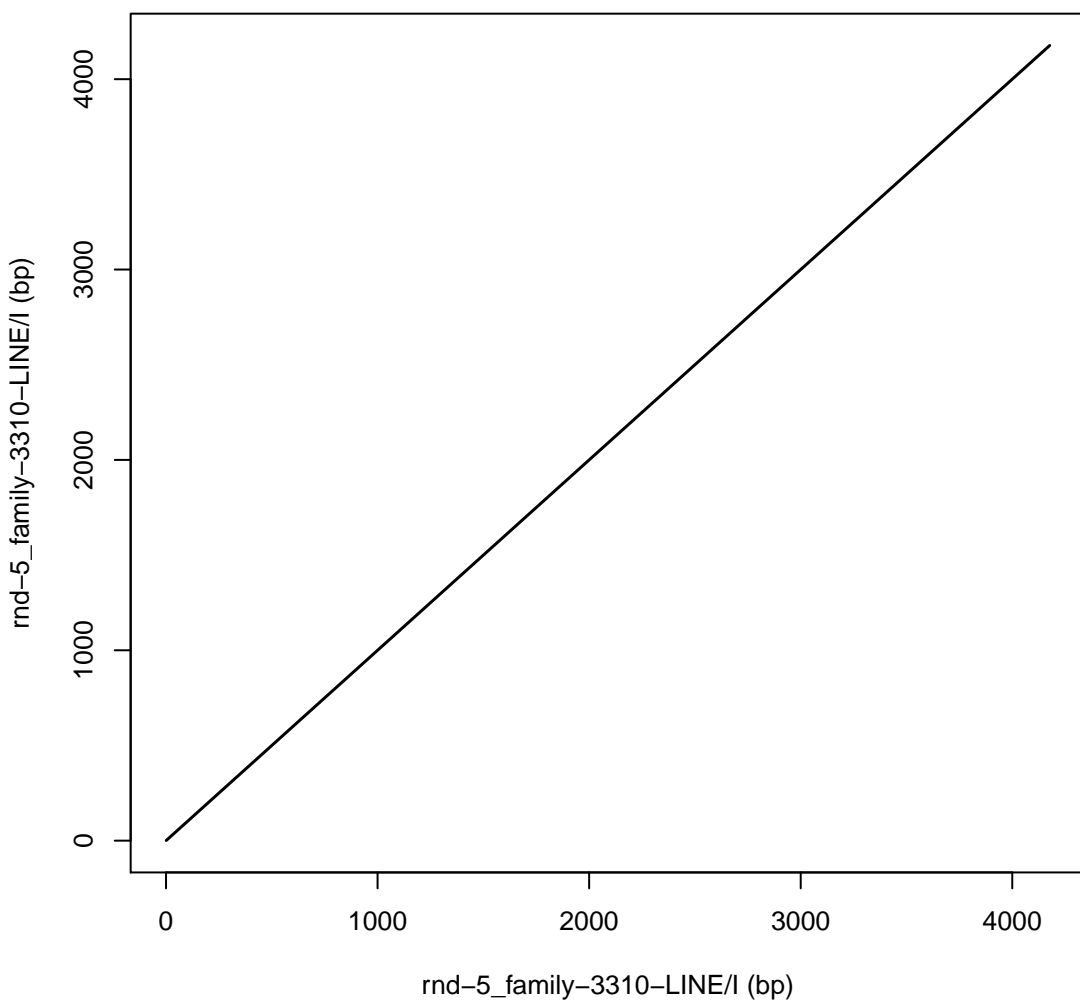
TE: rnd-5_family-3310-LINE/I
consensus size: 4177bp; fragments: 86; full length: 8 (>=3759.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

