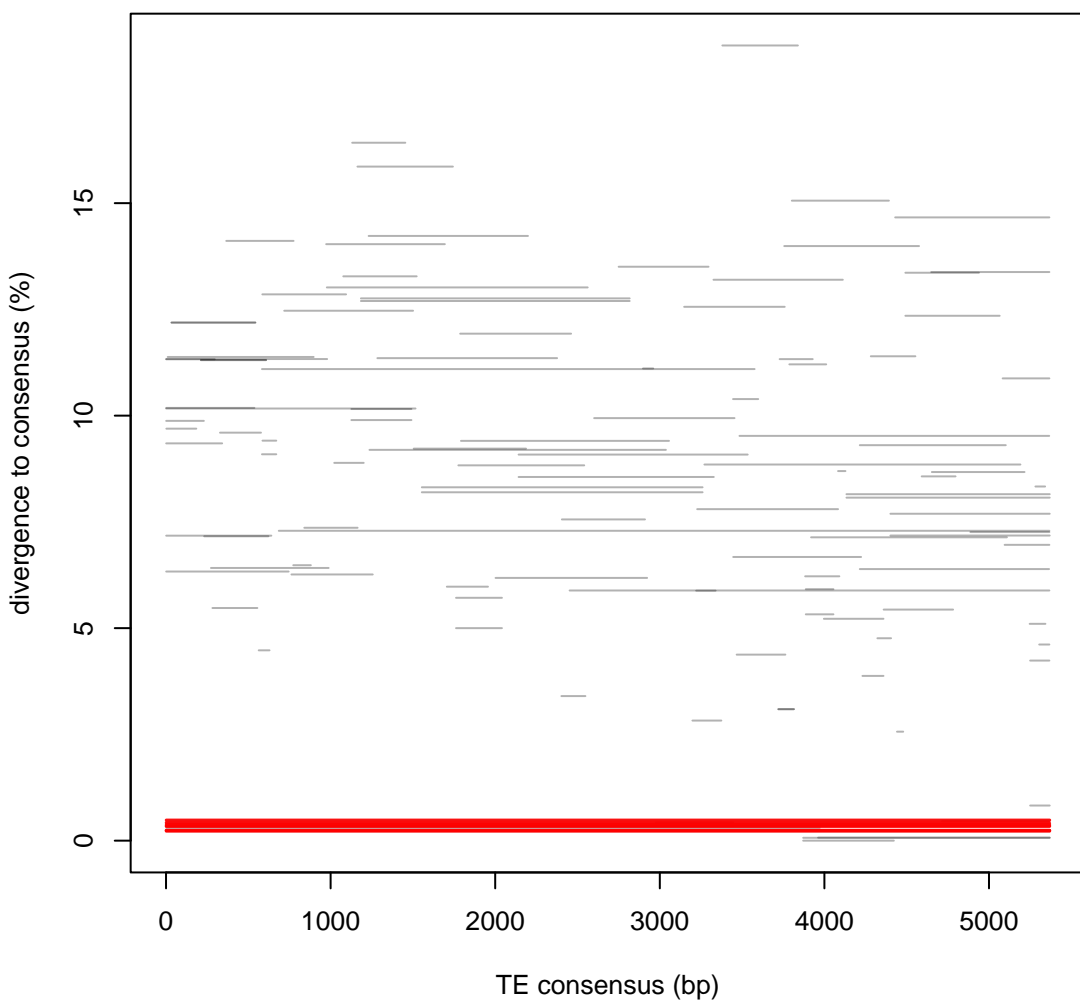
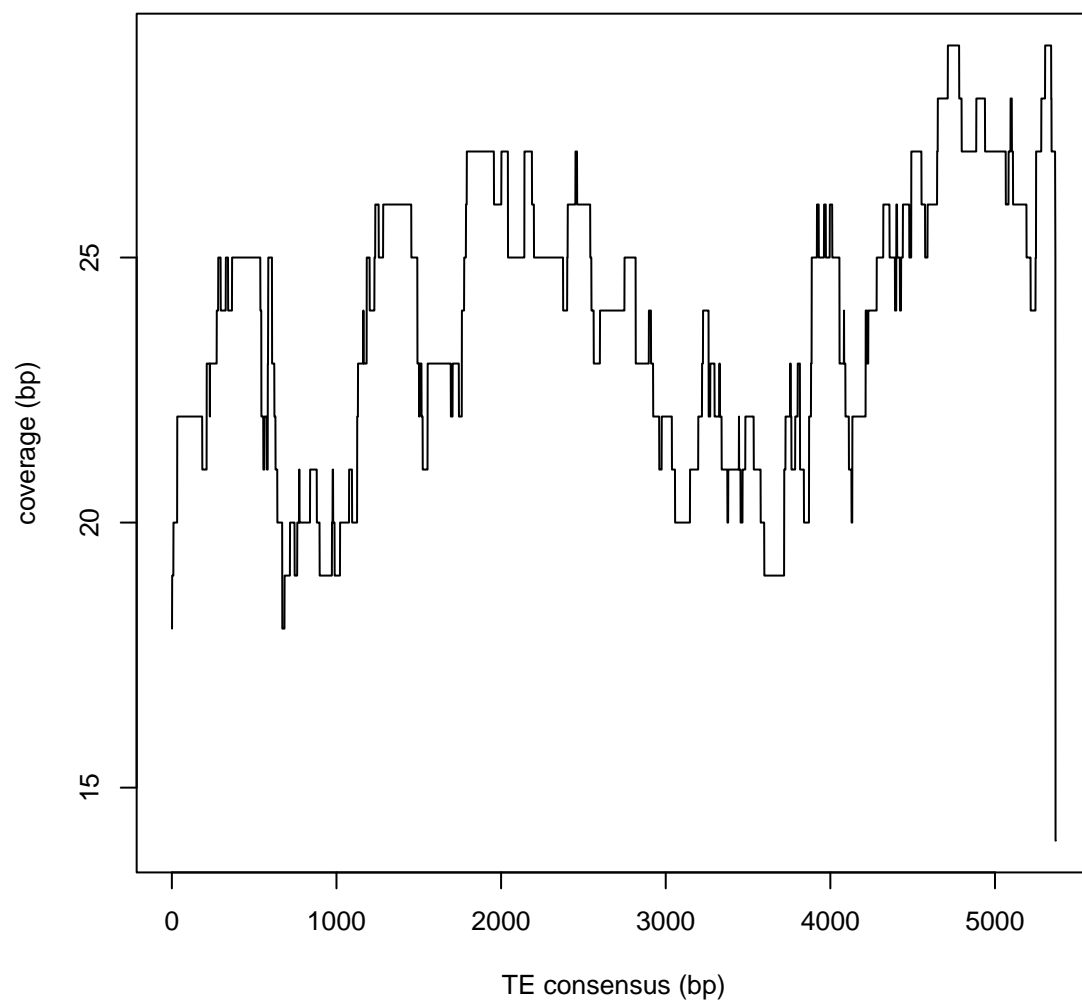


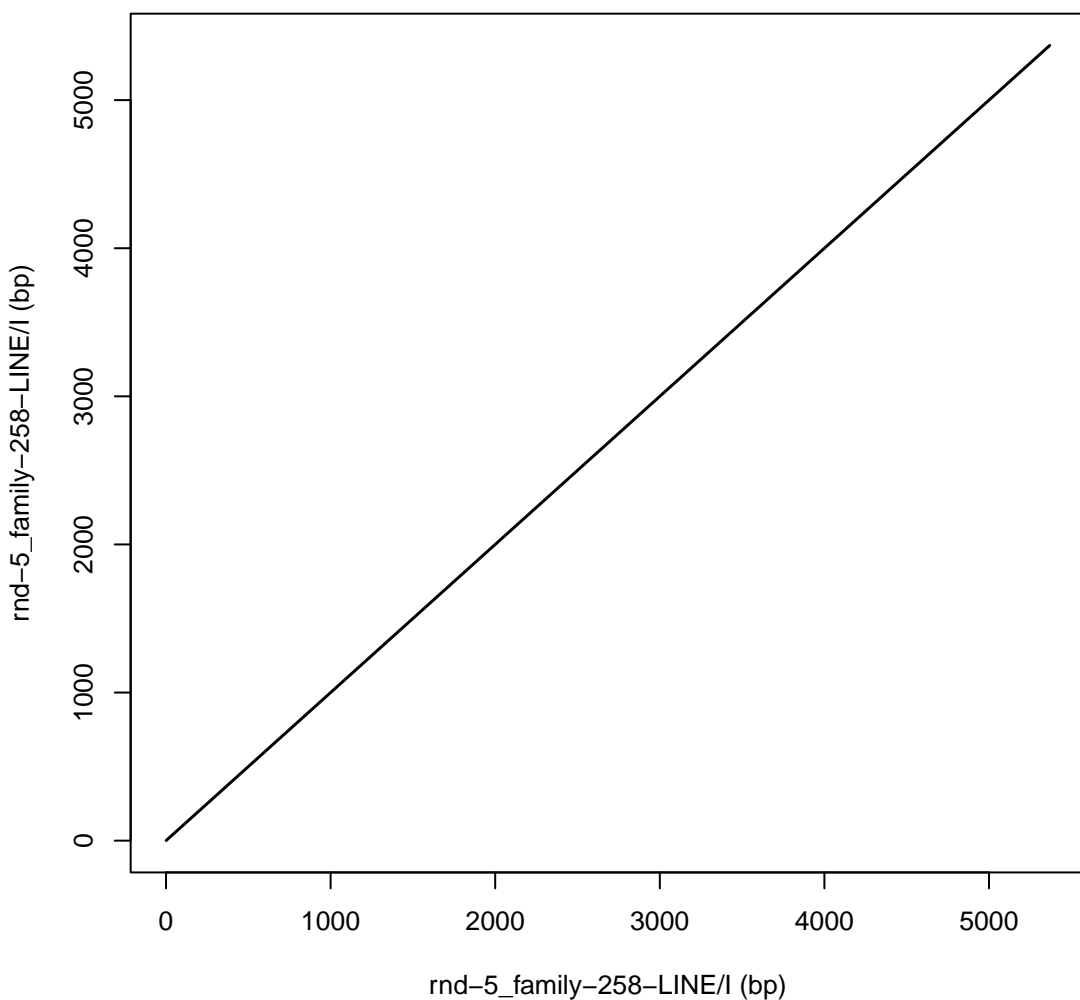
TE: rnd-5_family-258-LINE/I
consensus size: 5369bp; fragments: 123; full length: 8 (>=4832.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

