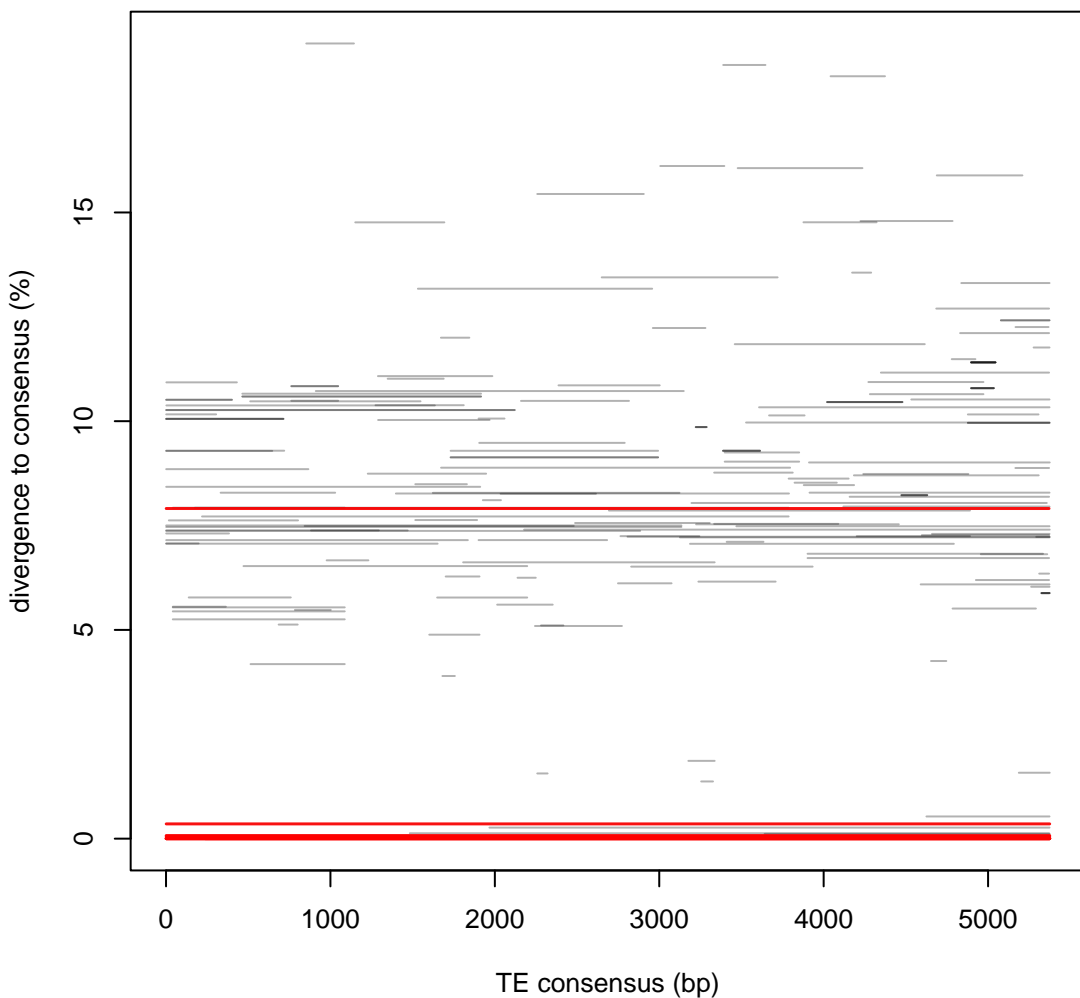
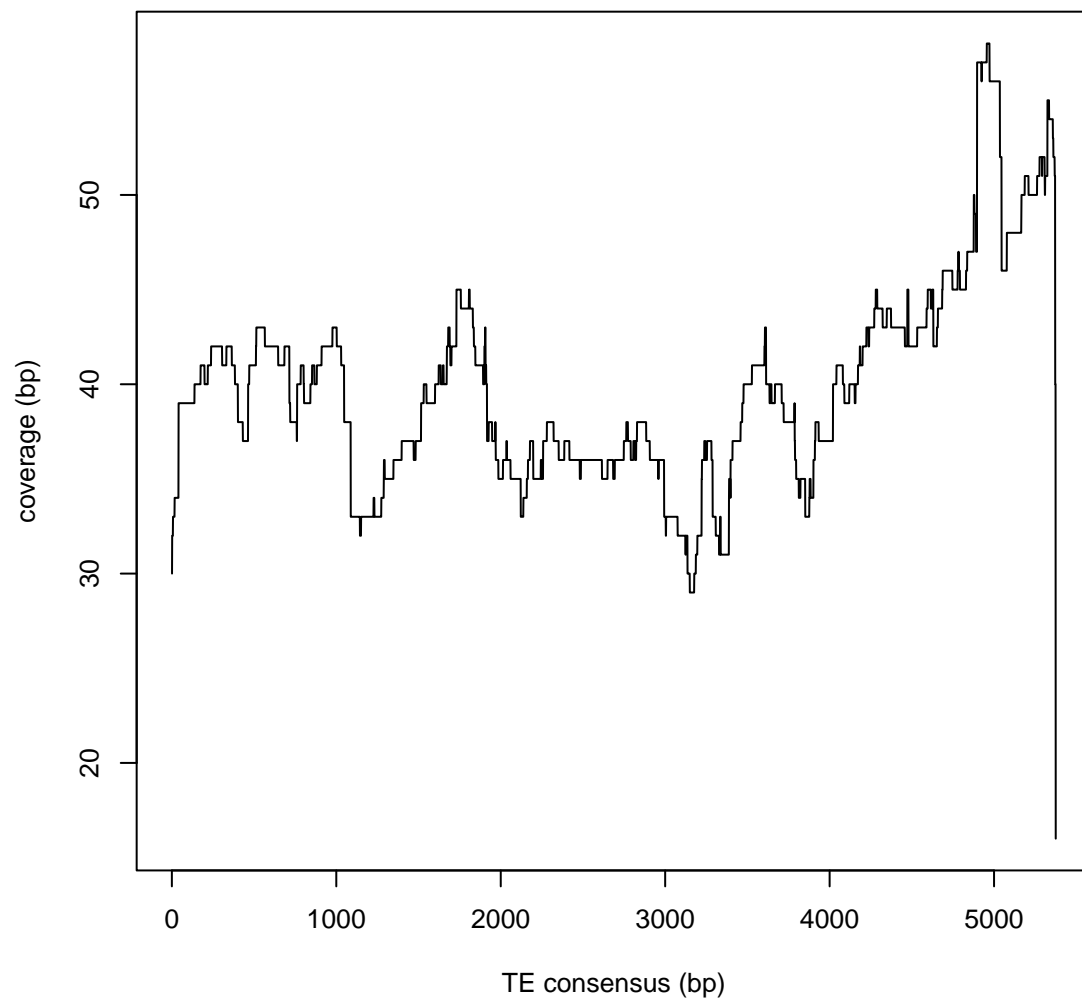


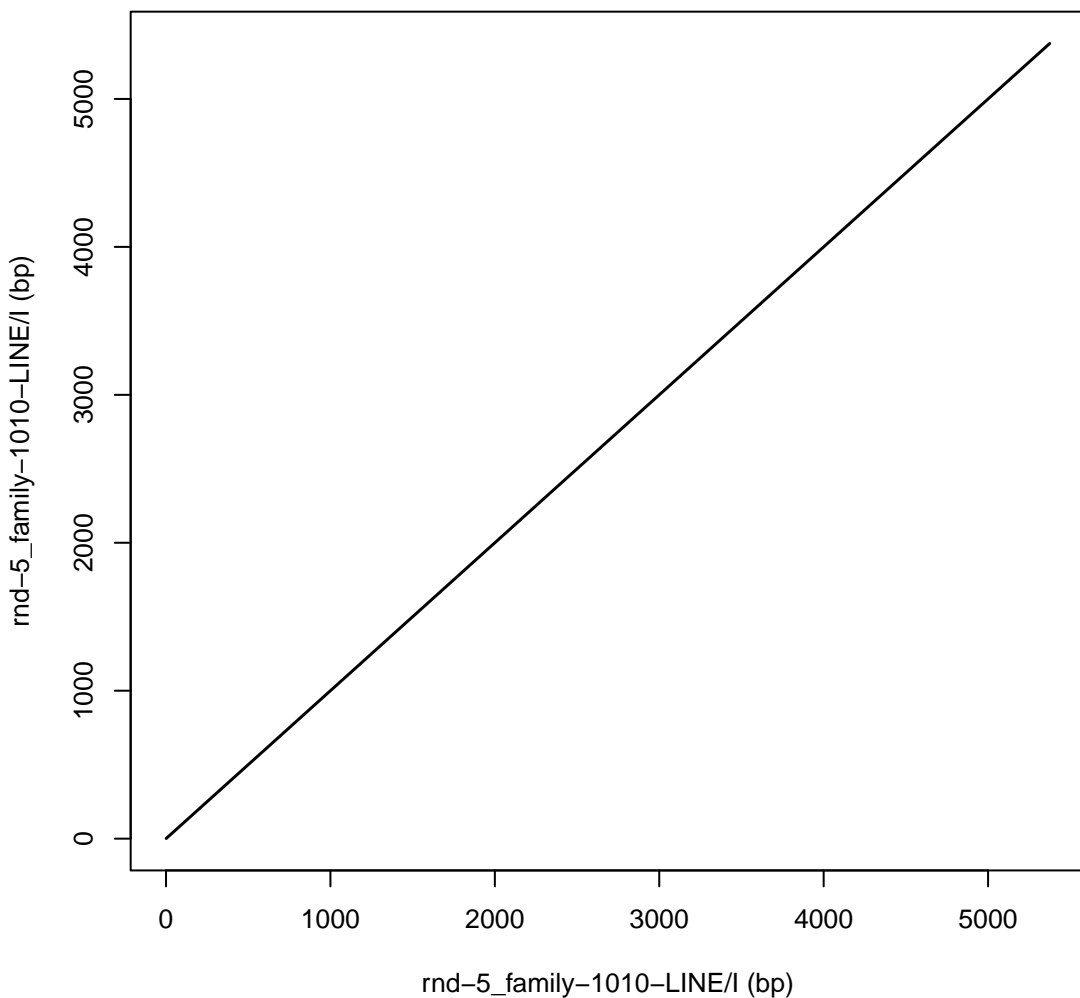
TE: rnd-5_family-1010-LINE/I
consensus size: 5375bp; fragments: 200; full length: 10 (>=4837.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

