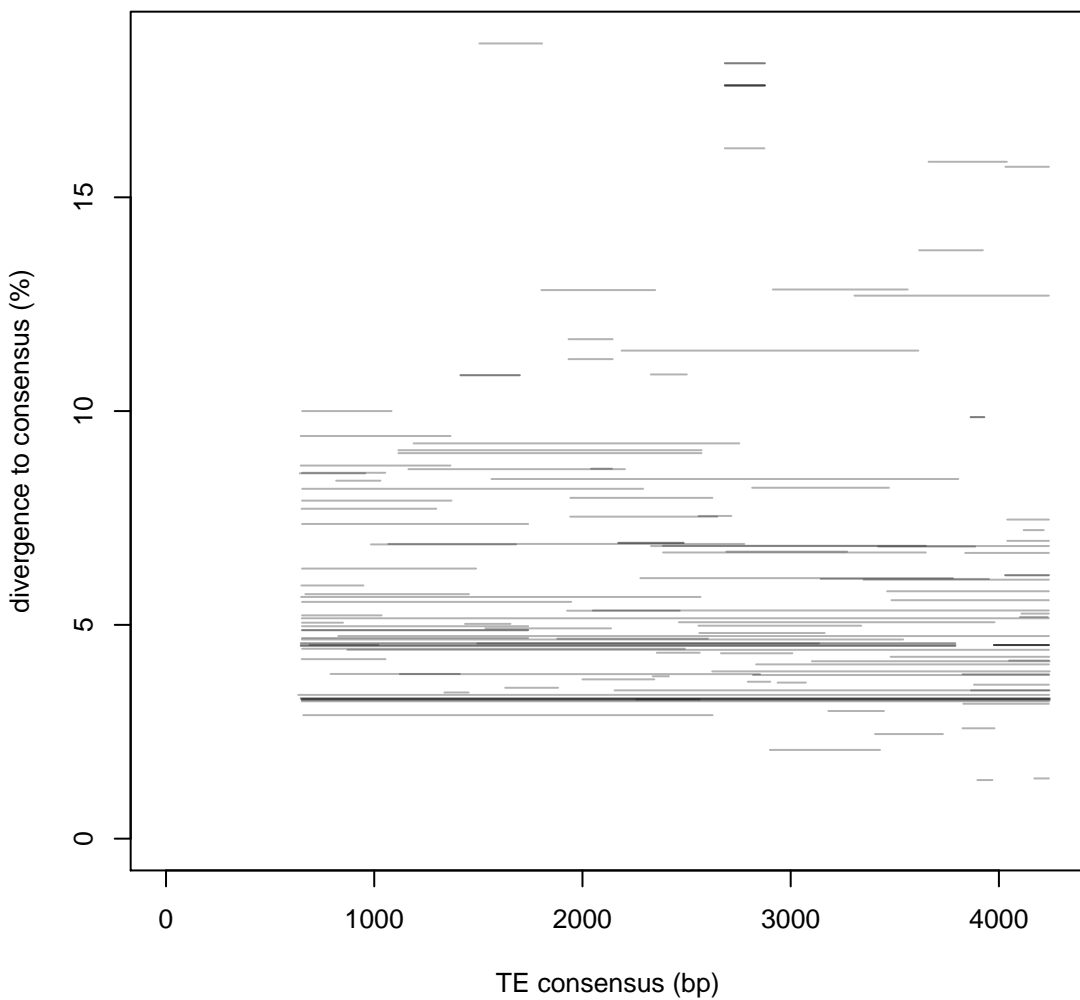
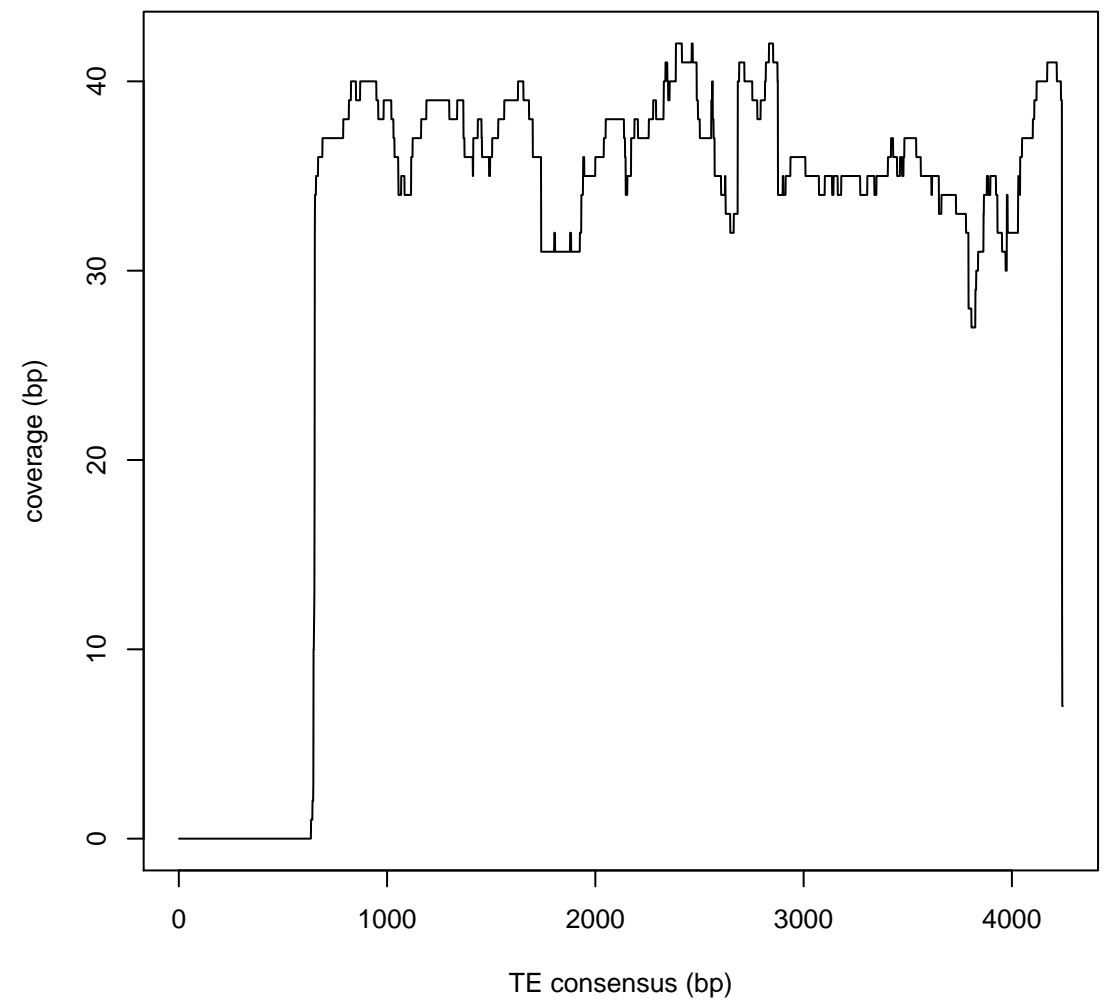


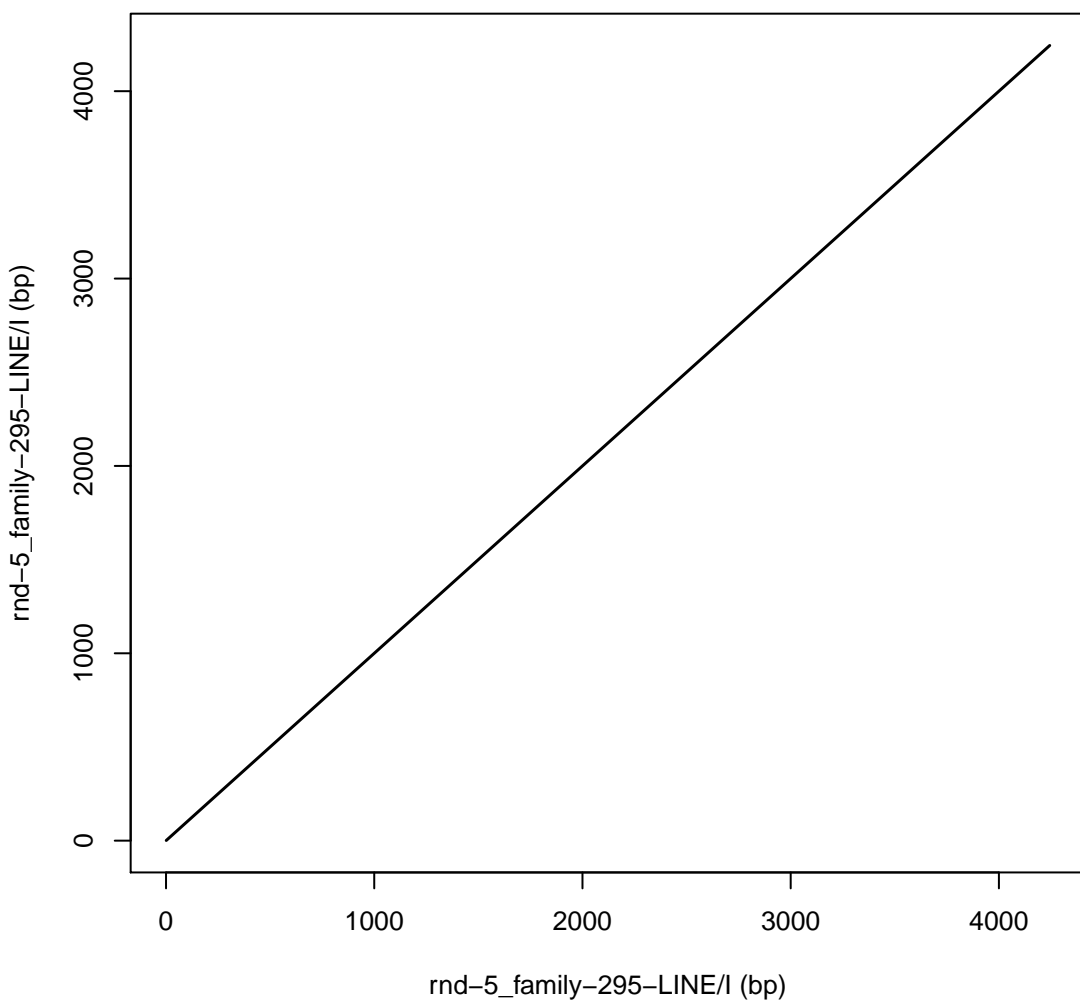
TE: rnd-5\_family-295-LINE/I  
consensus size: 4244bp; fragments: 135; full length: 0 (>=3819.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

