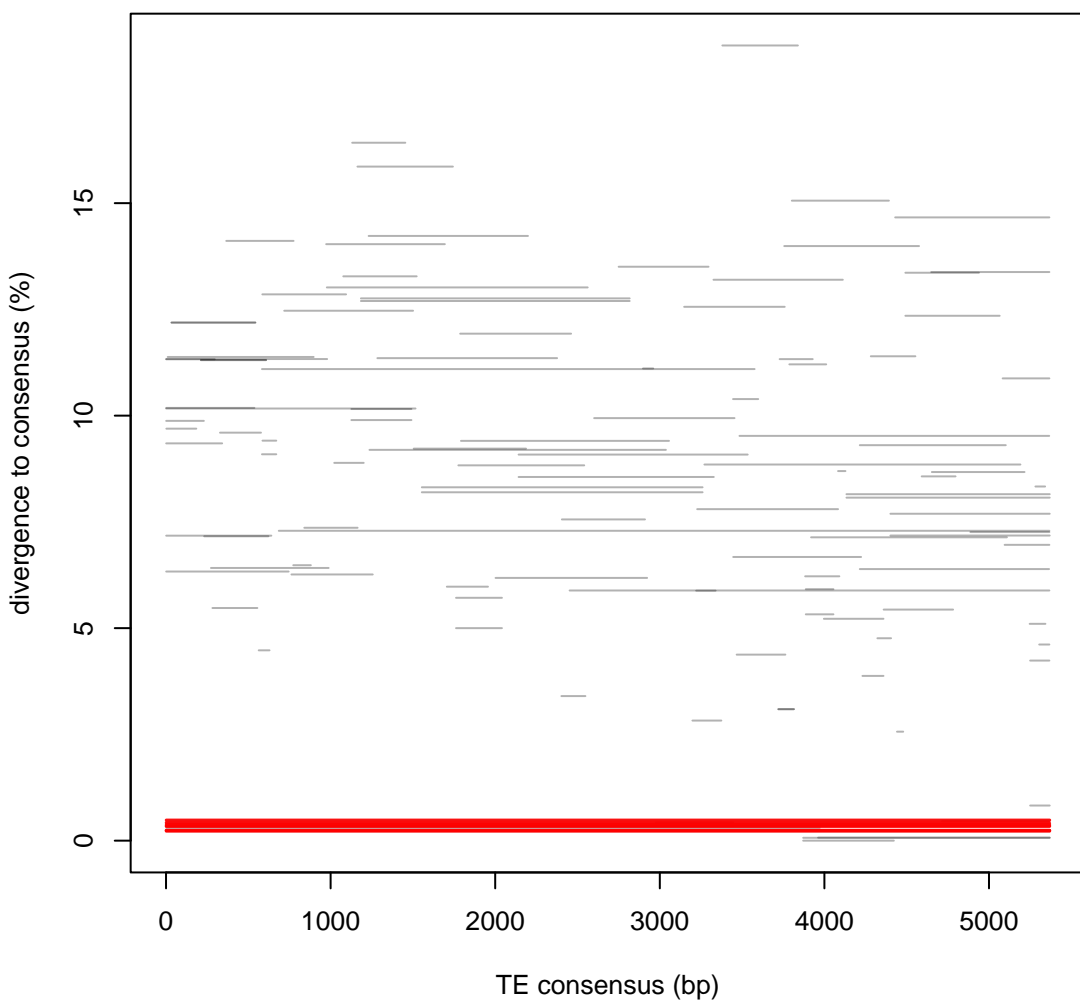
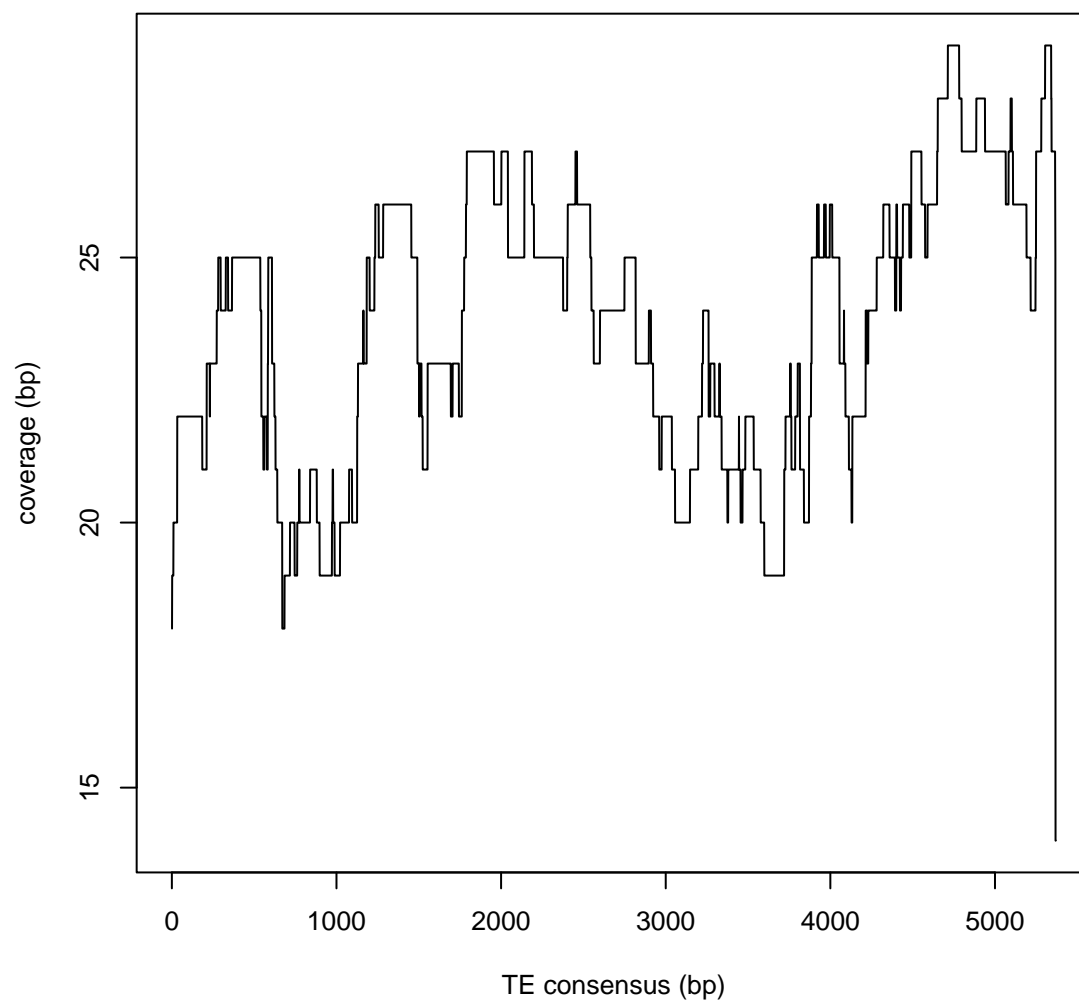


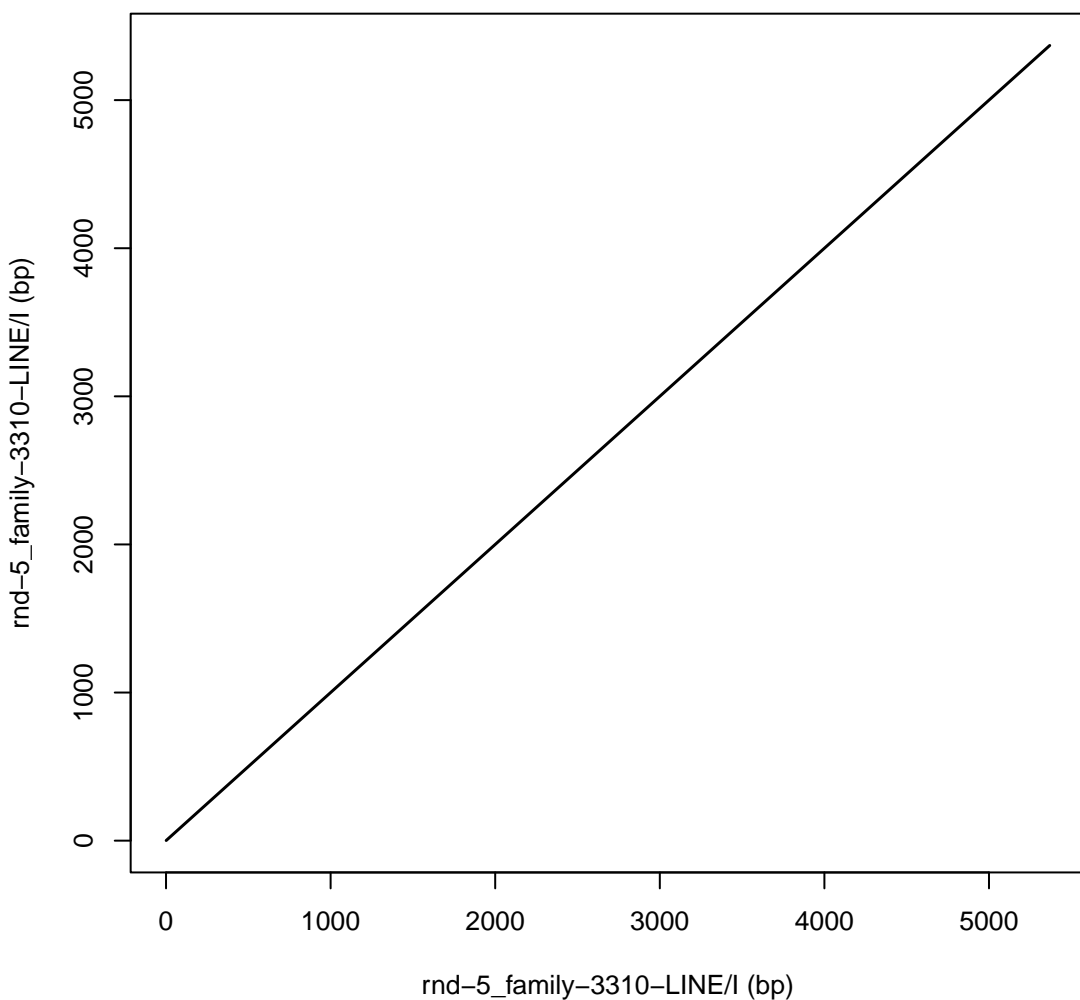
TE: rnd-5\_family-3310-LINE/I  
consensus size: 5369bp; fragments: 123; full length: 8 (>=4832.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

