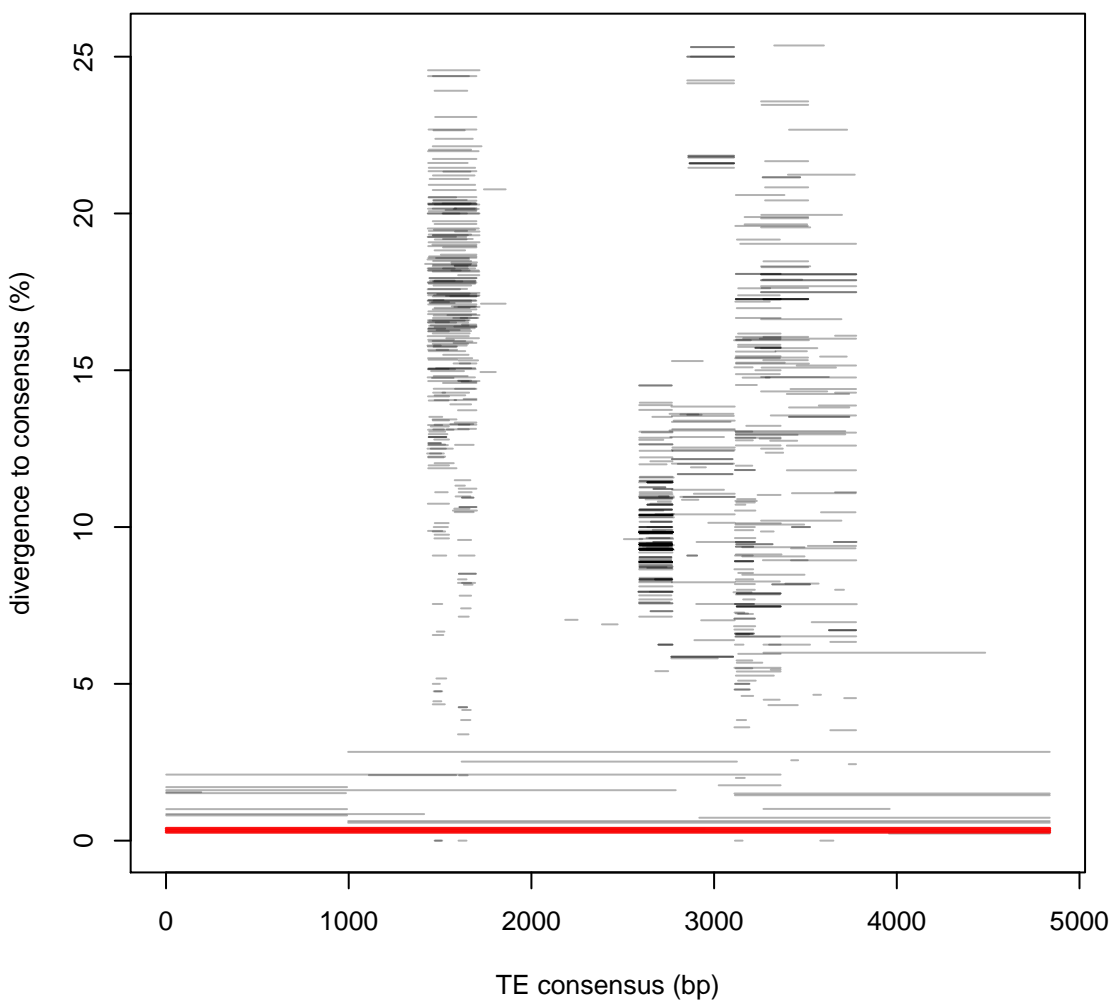
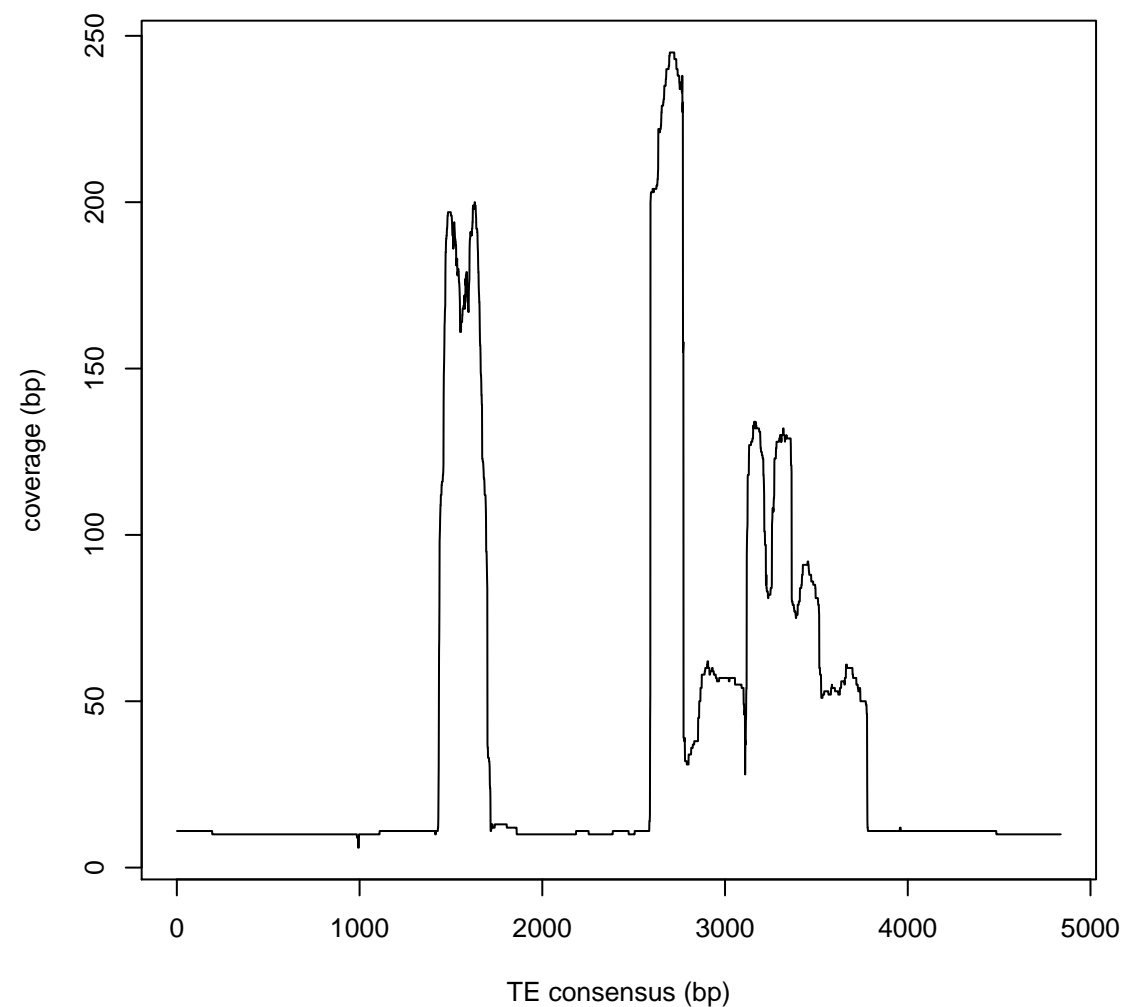


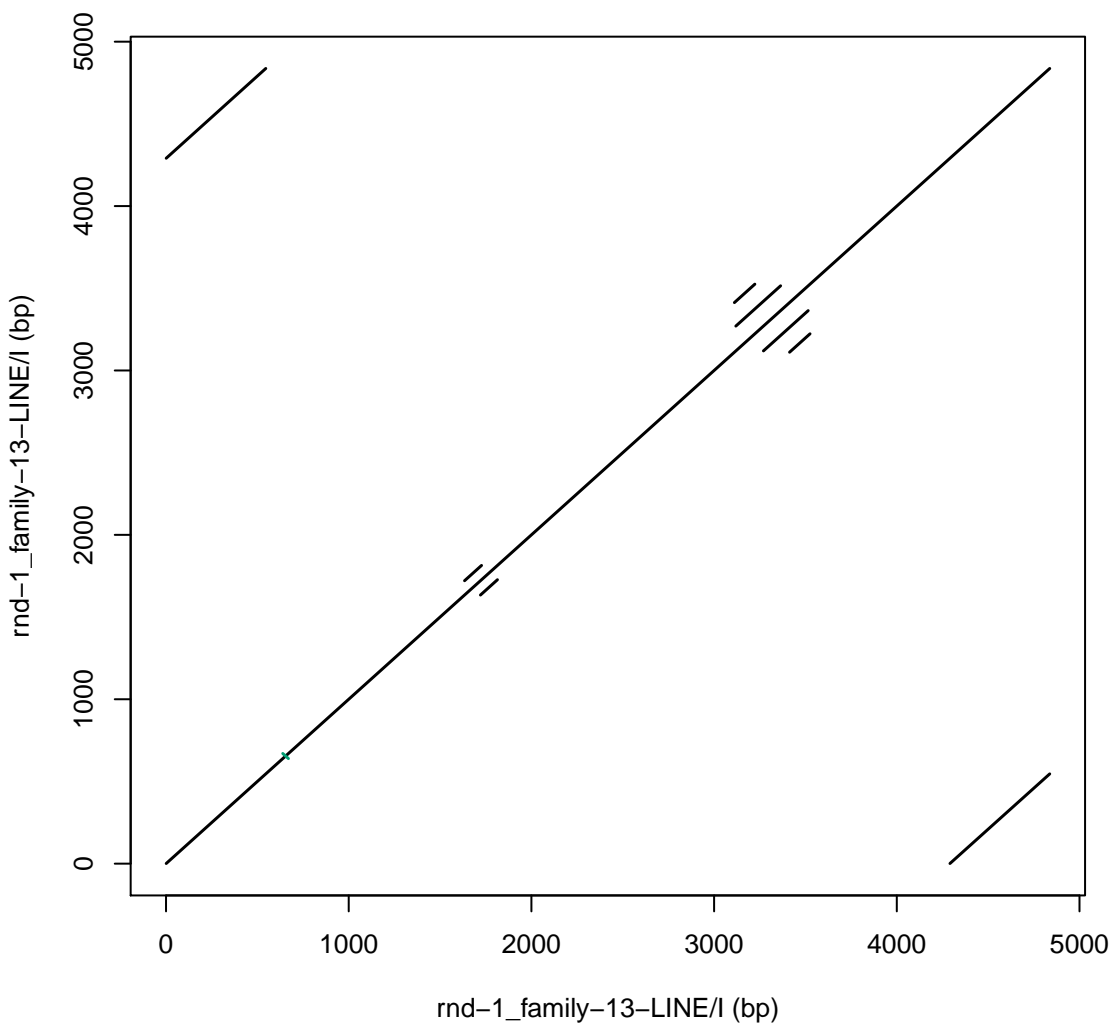
TE: rnd-1\_family-13-LINE/I  
consensus size: 4837bp; fragments: 811; full length: 3 (>=4353.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

