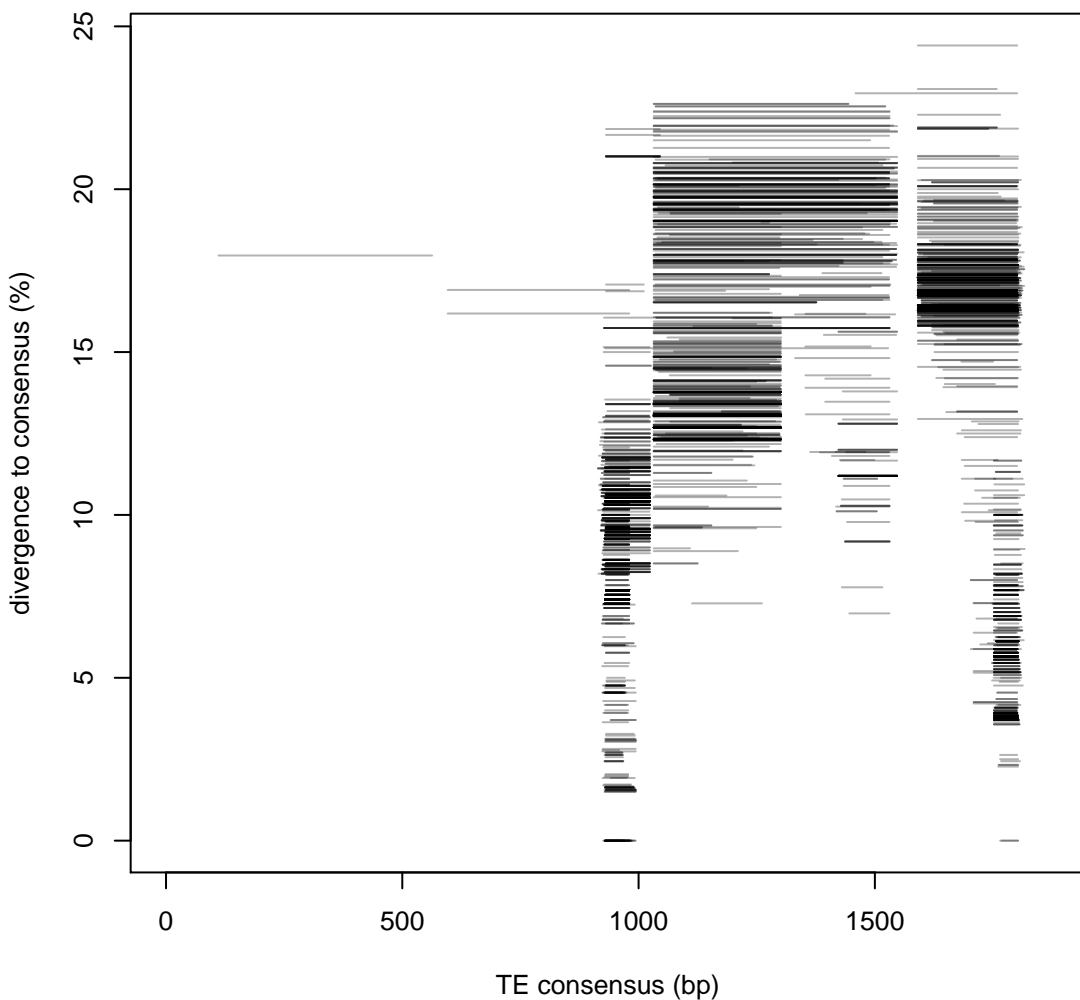
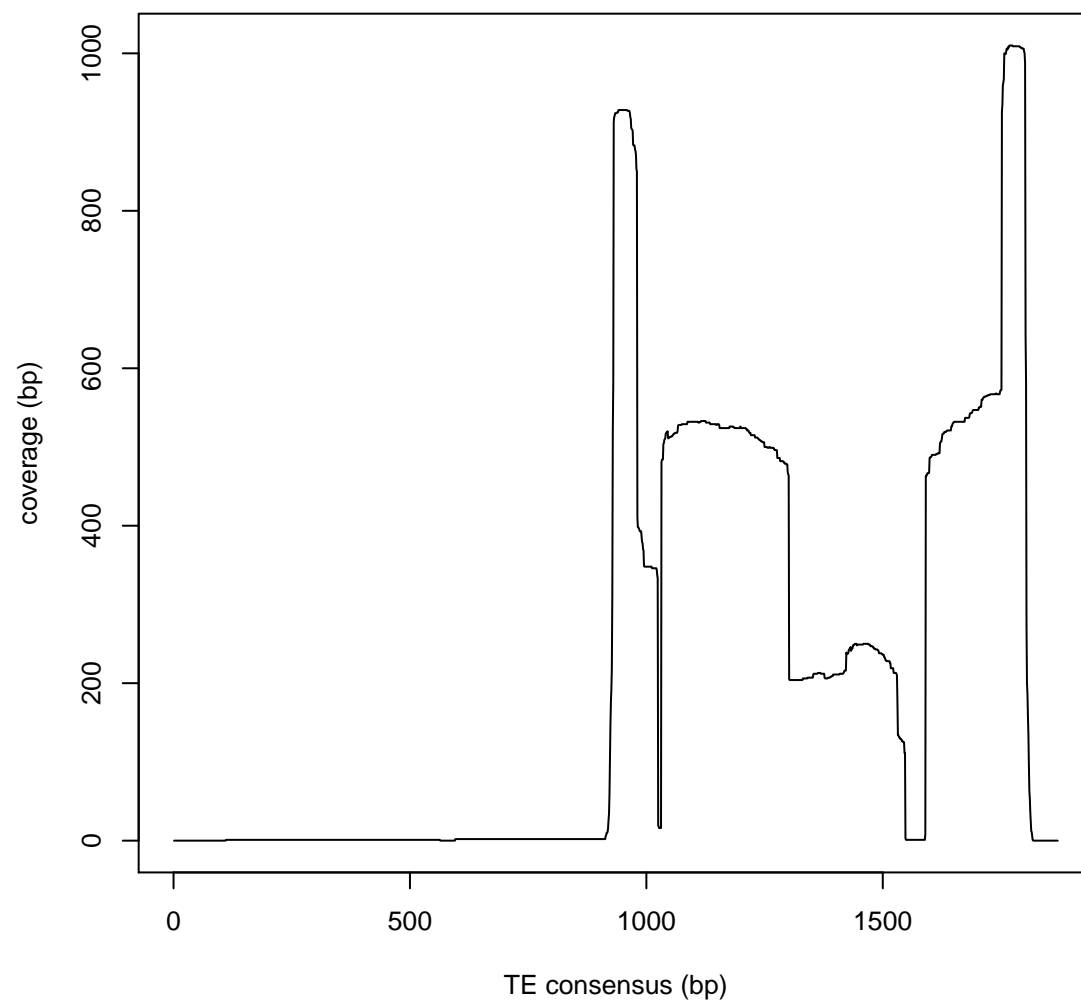


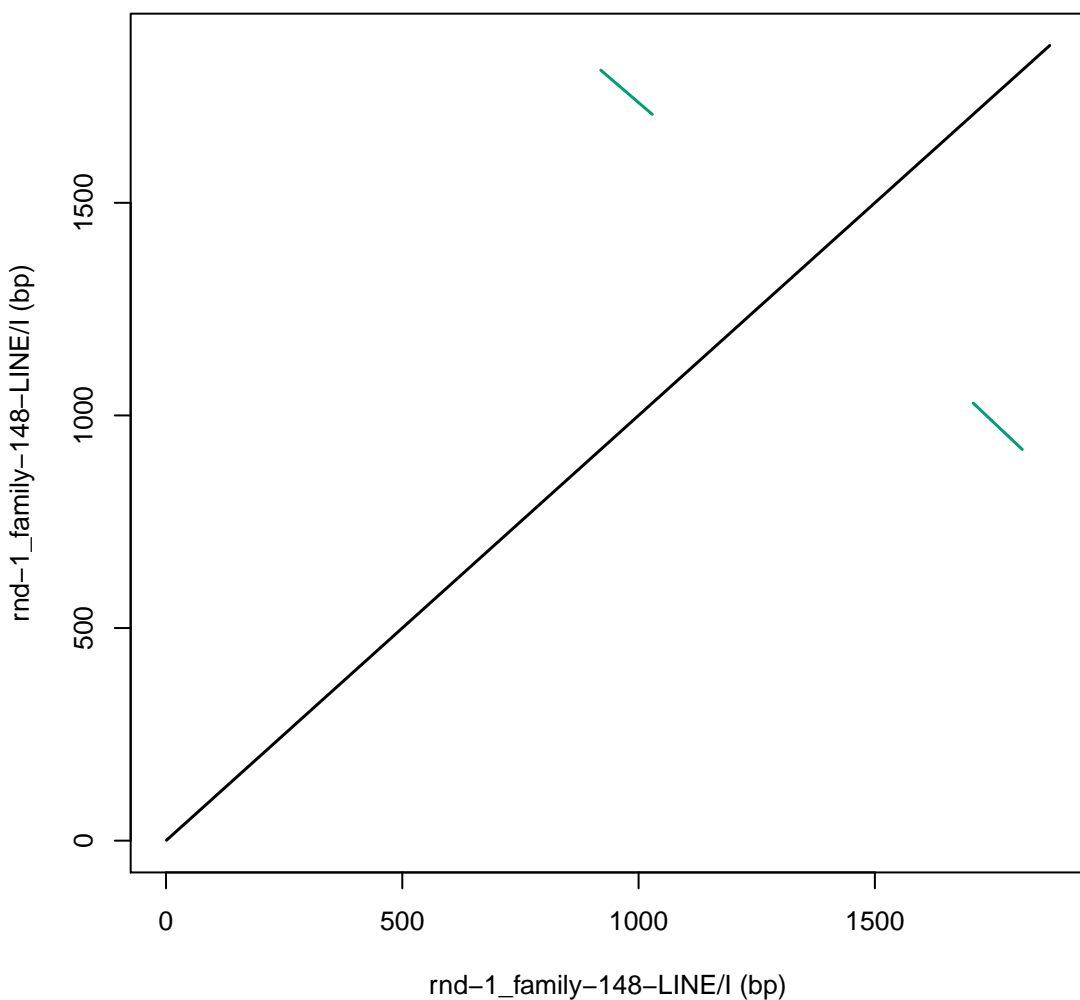
TE: rnd-1\_family-148-LINE/I  
consensus size: 1870bp; fragments: 2548; full length: 0 (>=1683bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

