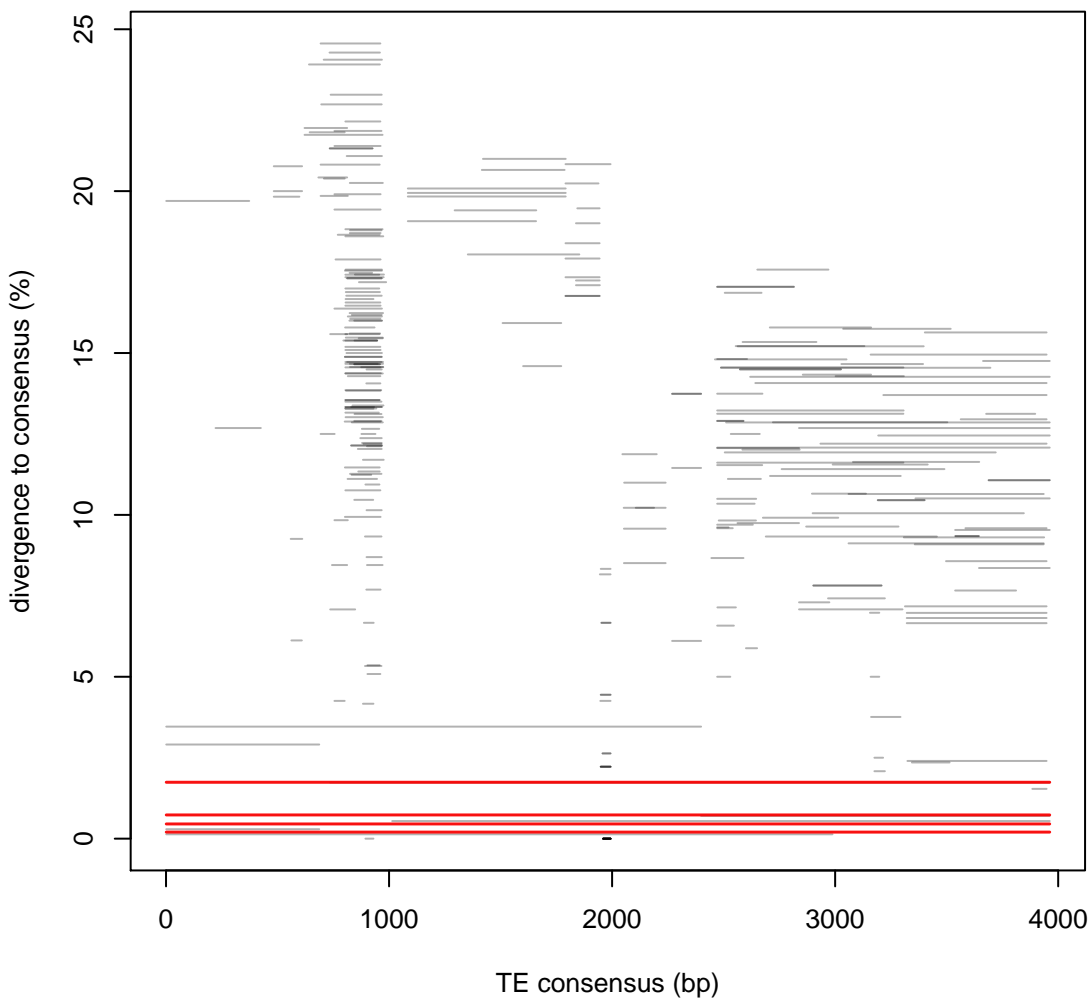
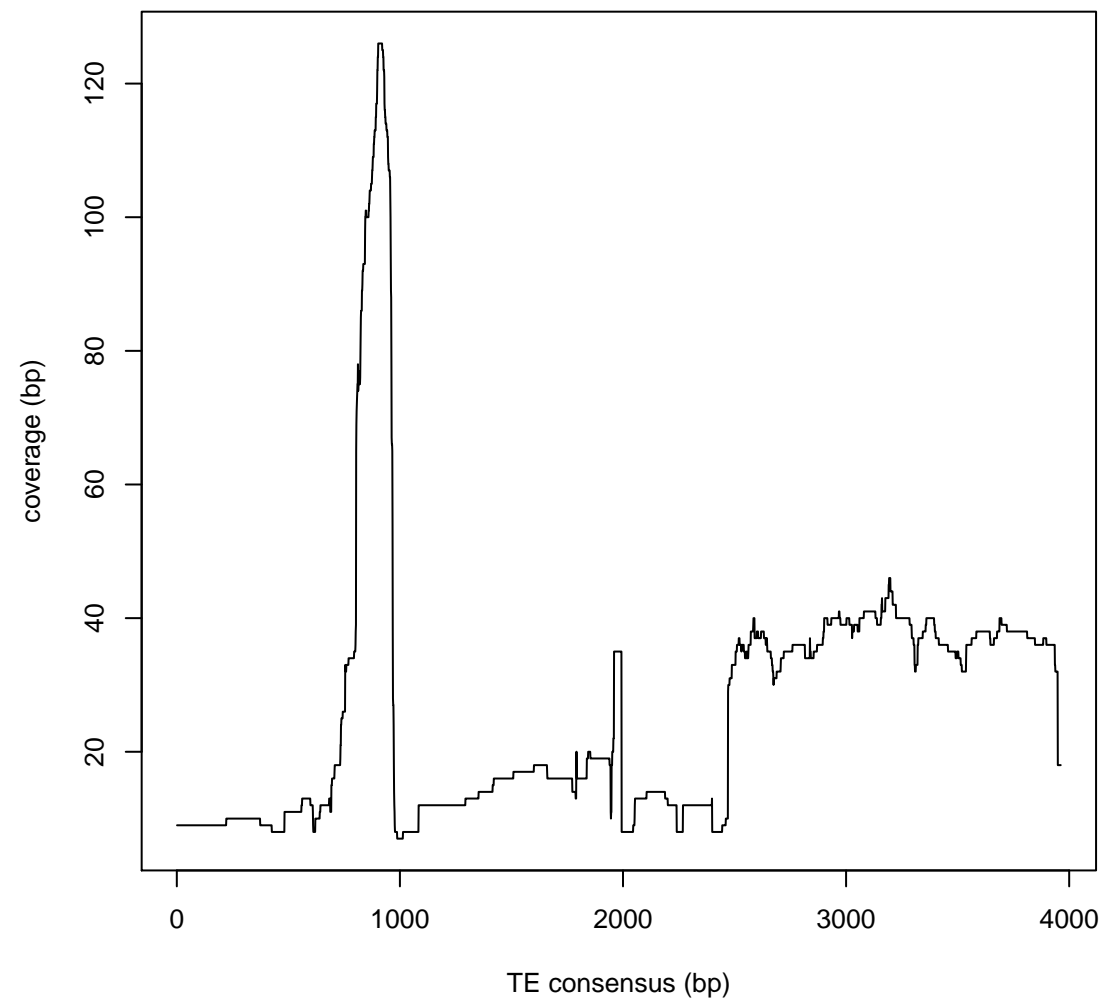


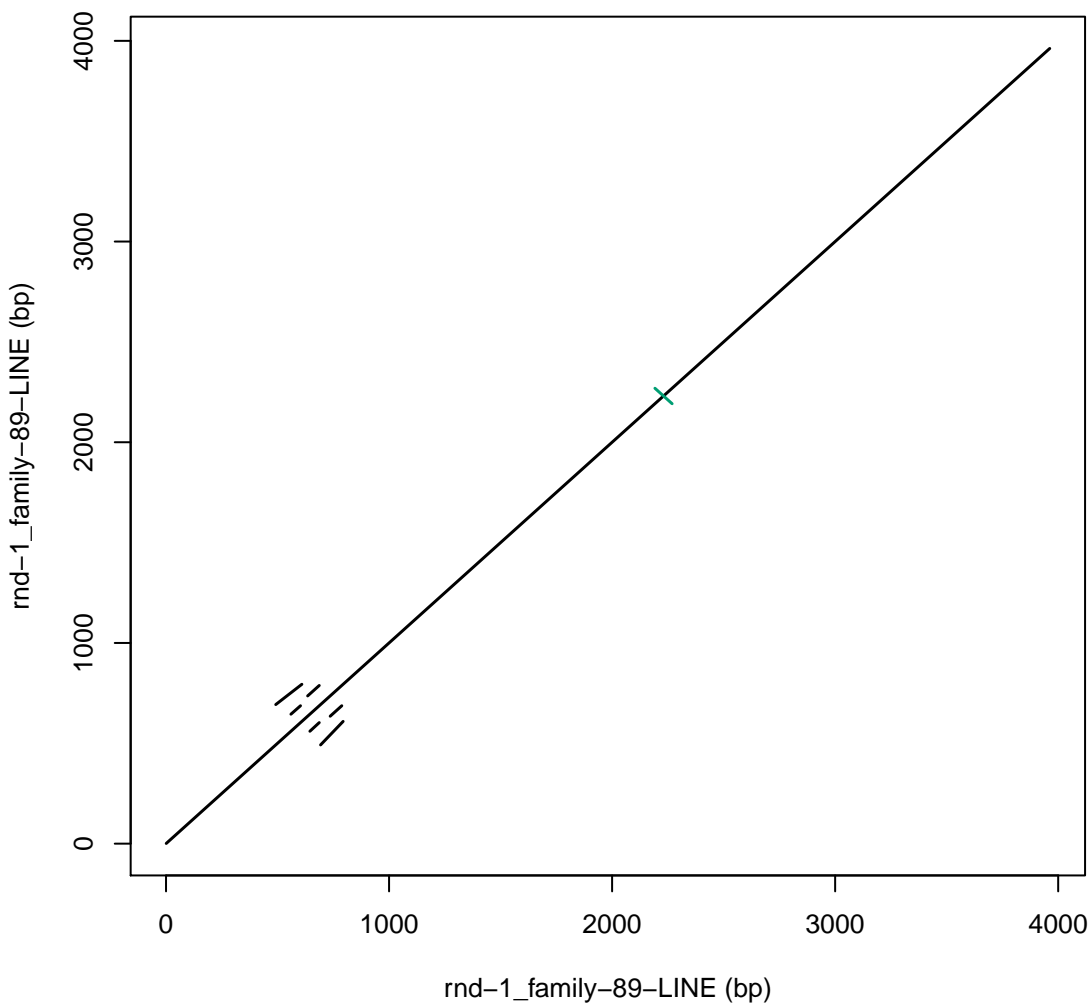
TE: rnd-1\_family-89-LINE  
consensus size: 3962bp; fragments: 303; full length: 4 (>=3565.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

