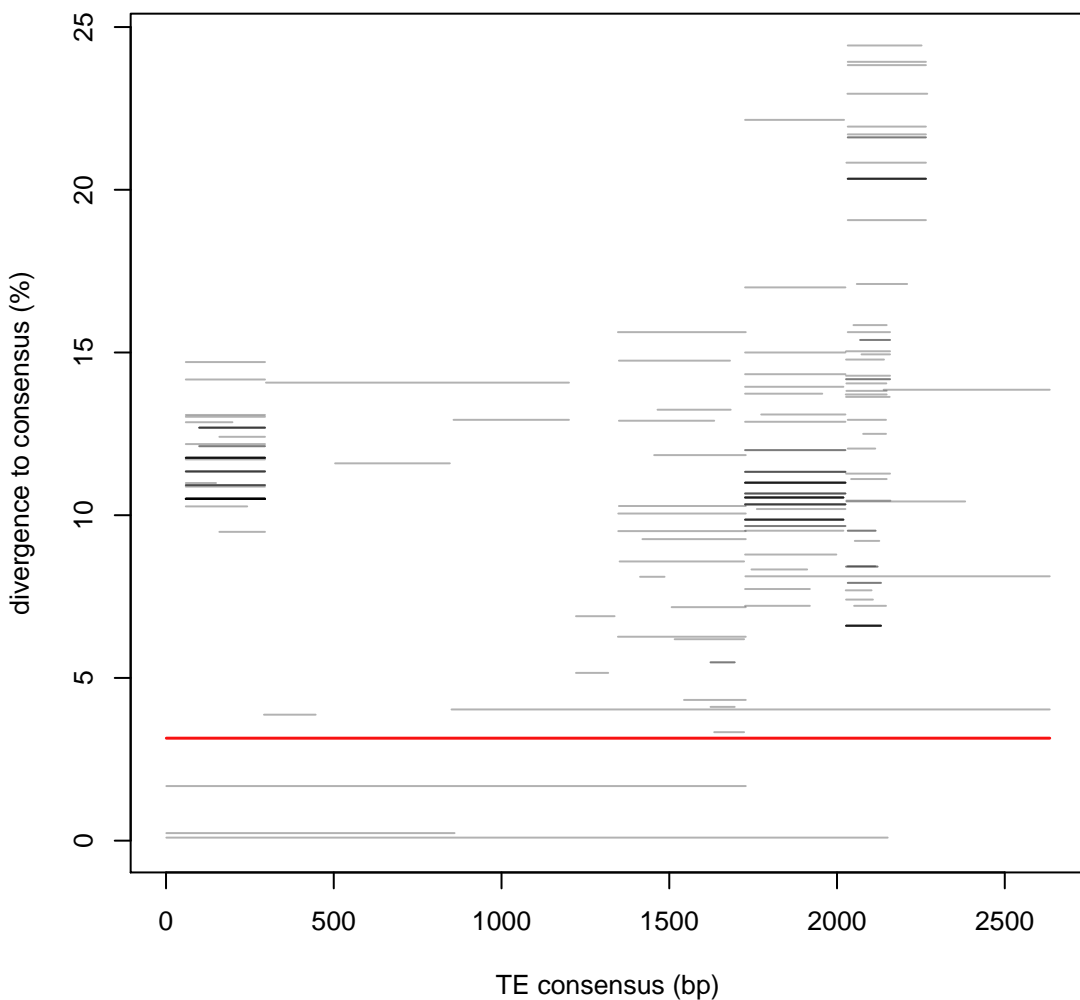
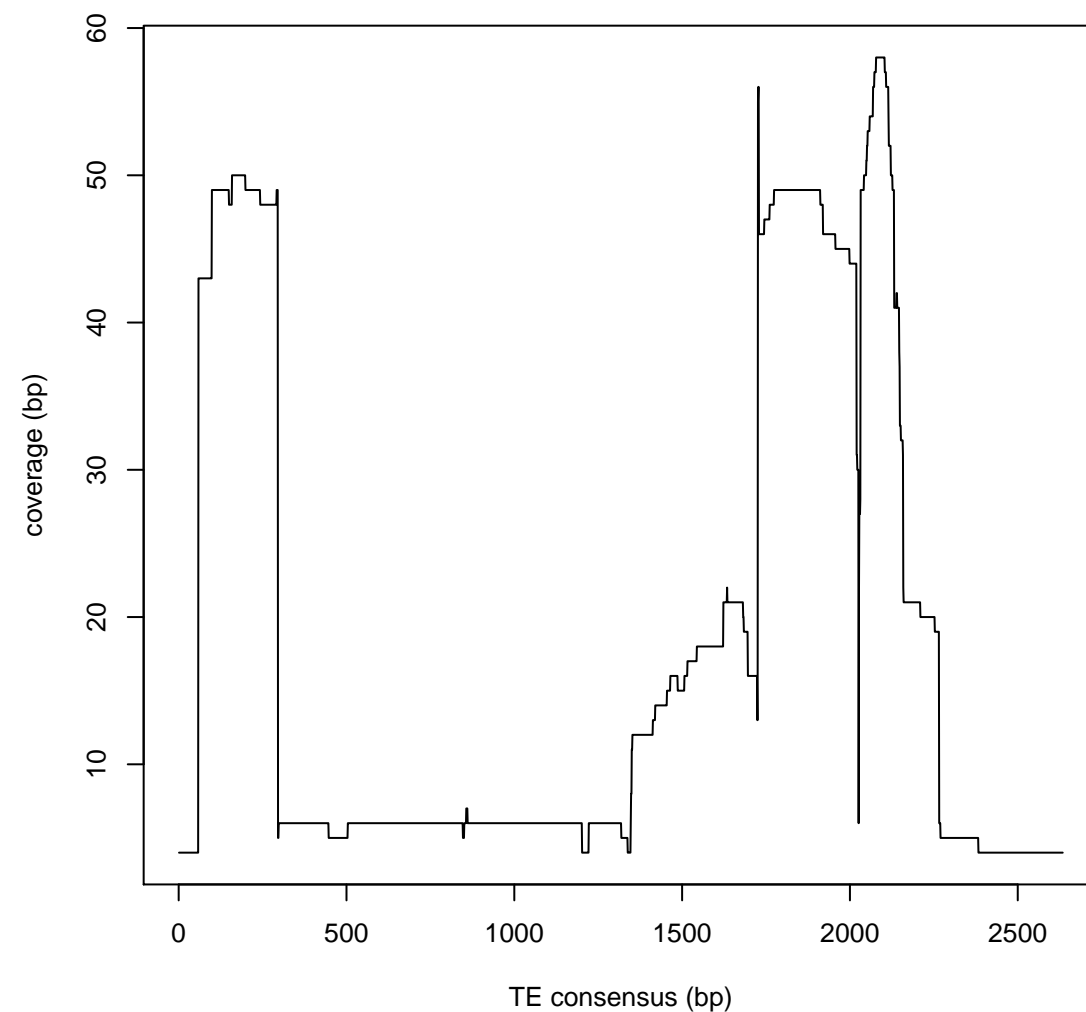


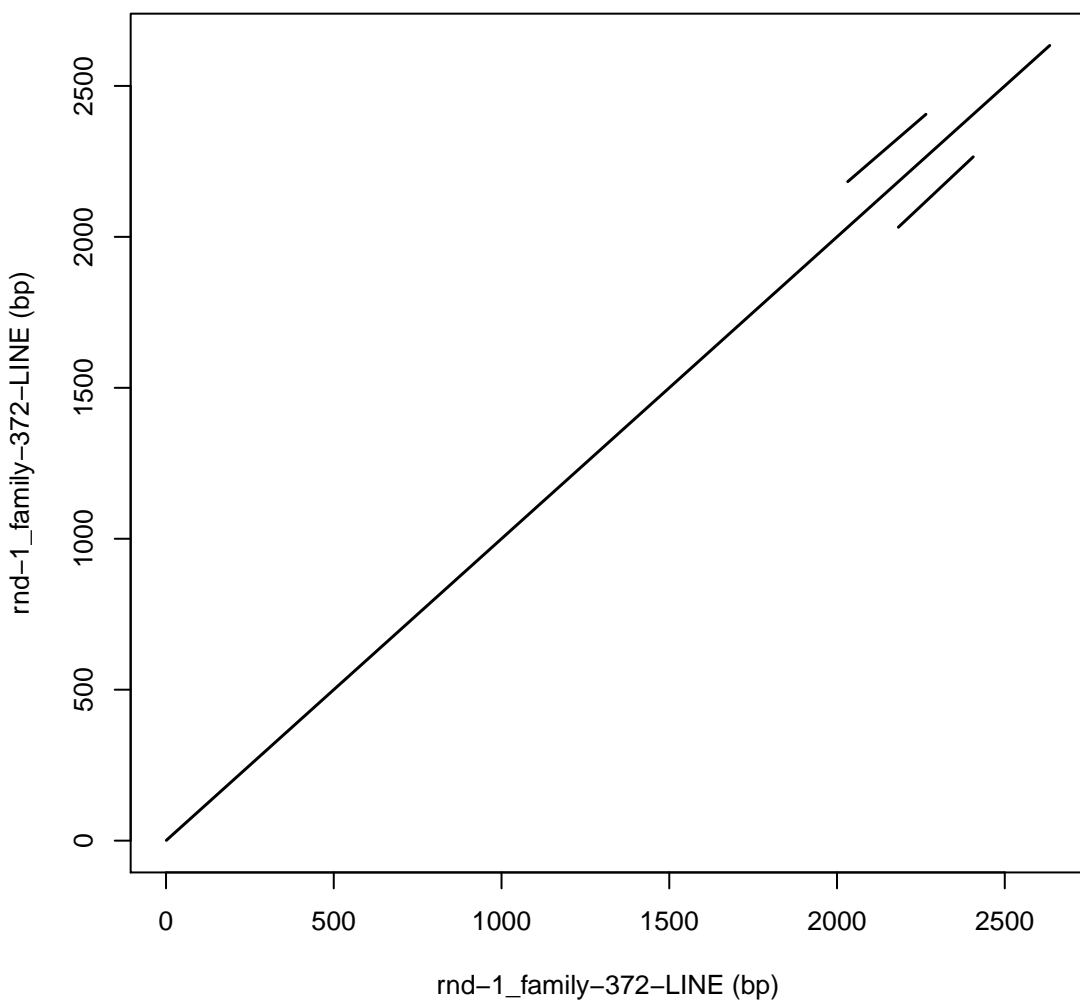
TE: rnd-1\_family-372-LINE  
consensus size: 2634bp; fragments: 178; full length: 1 (>=2370.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

