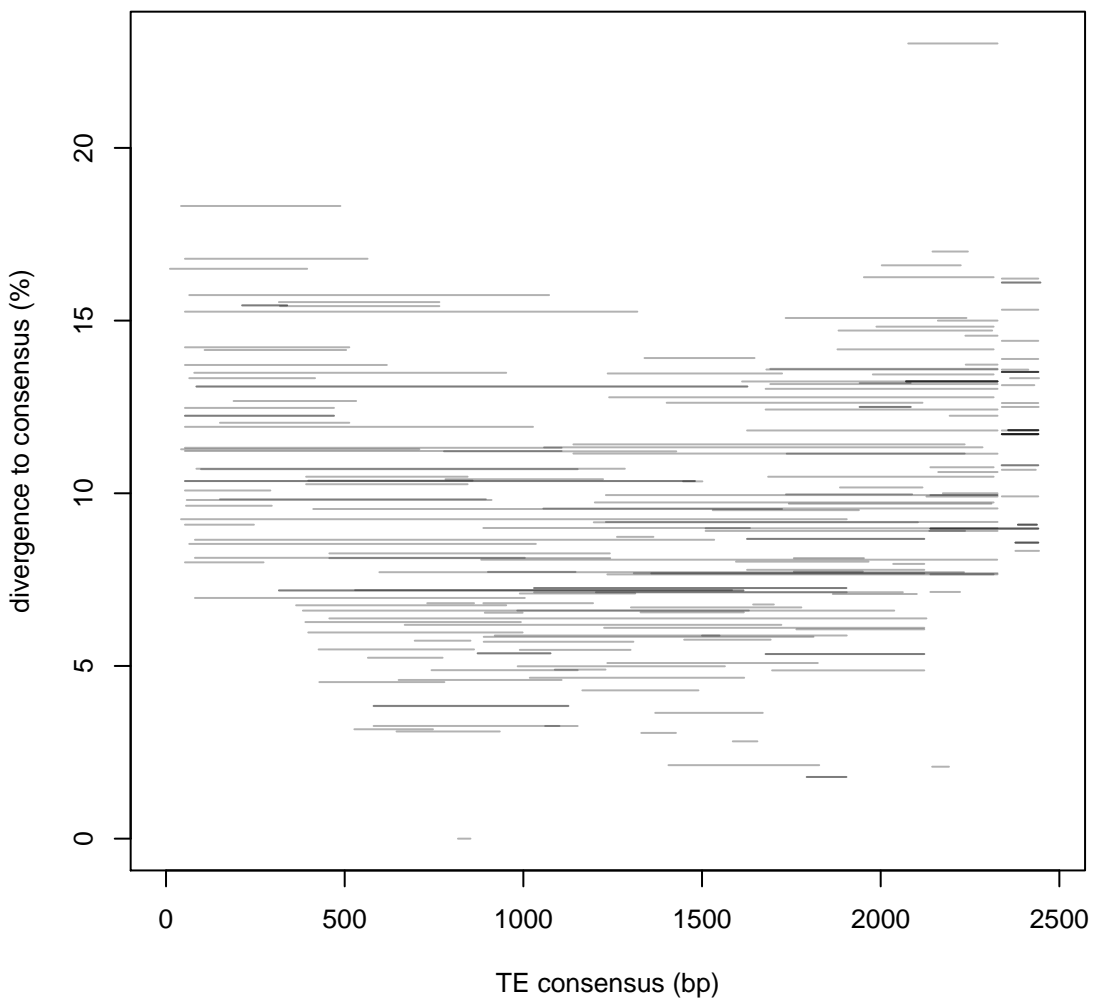
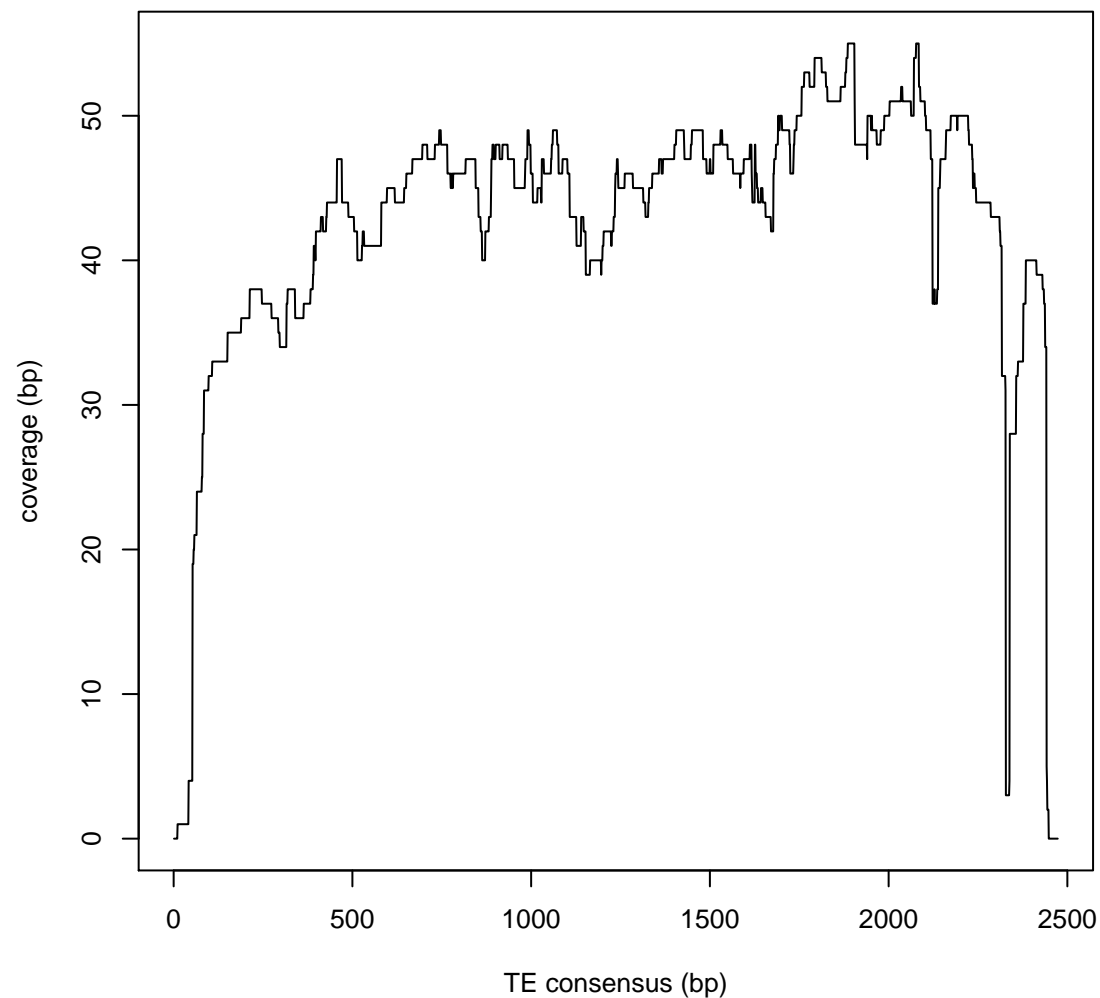


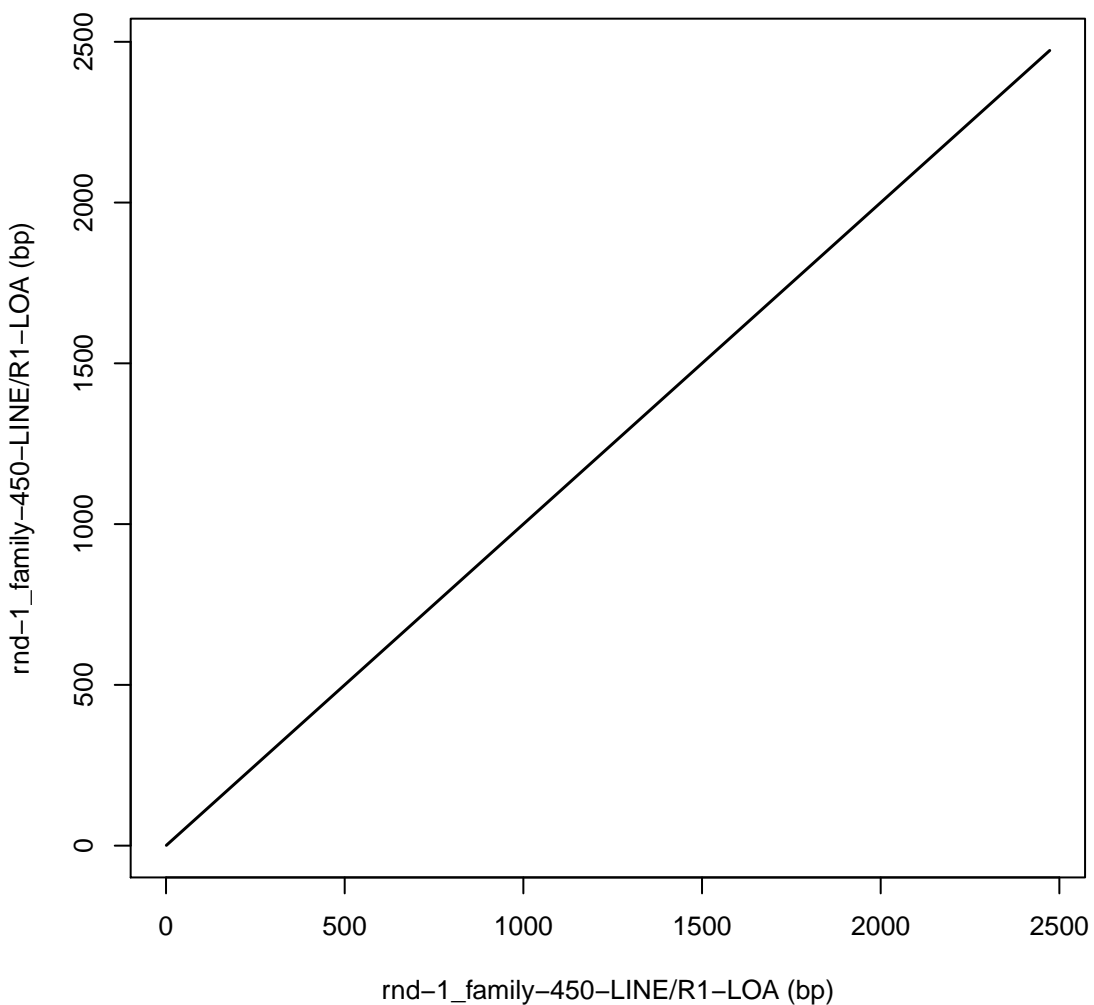
TE: rnd-1\_family-450-LINE/R1-LOA  
consensus size: 2473bp; fragments: 219; full length: 0 (>=2225.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

