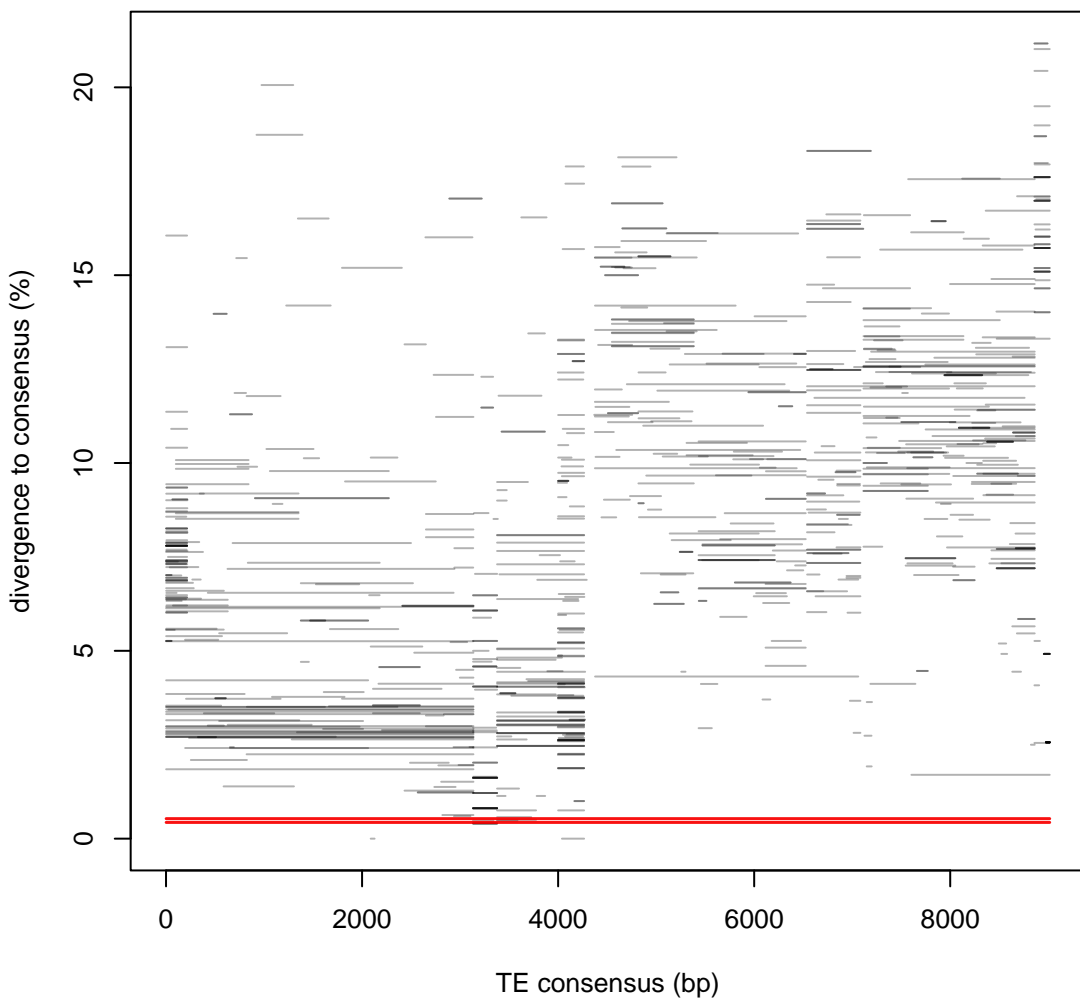
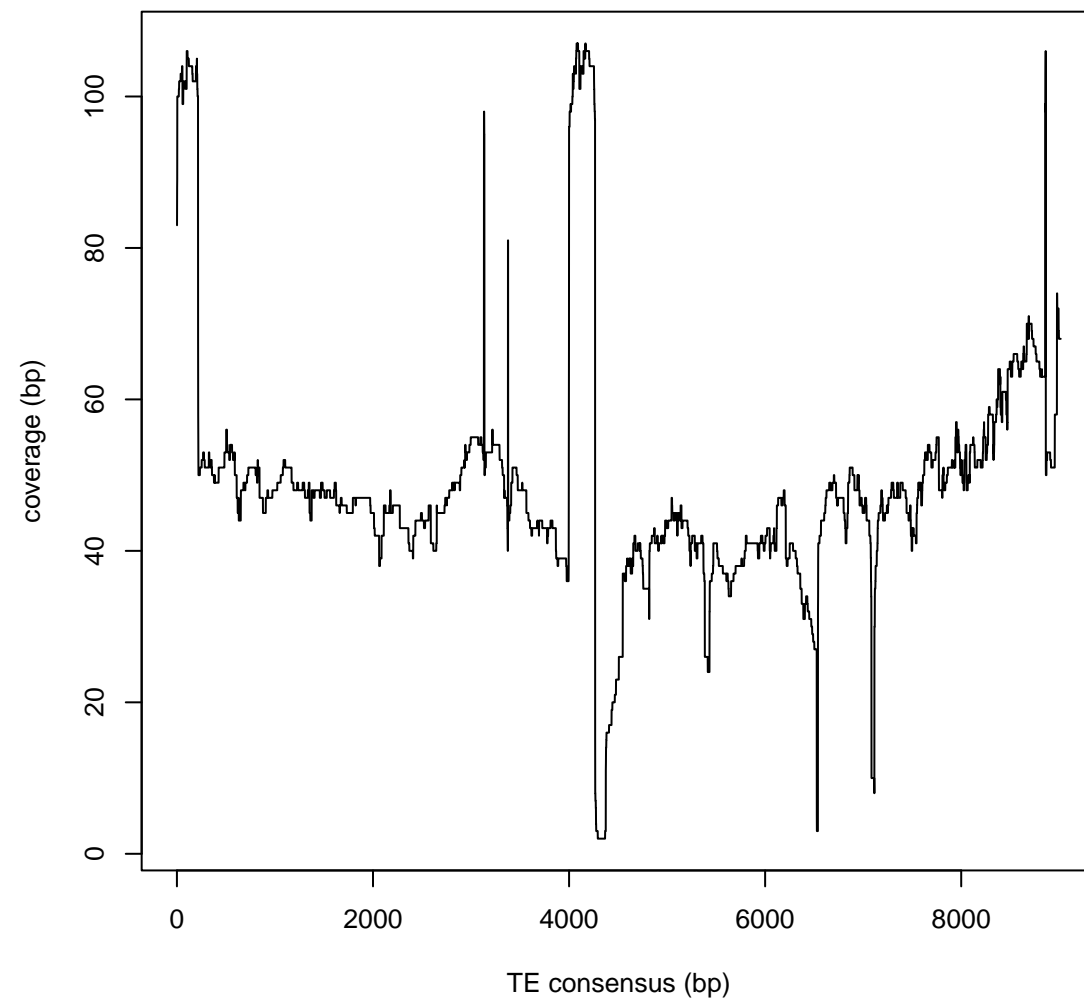


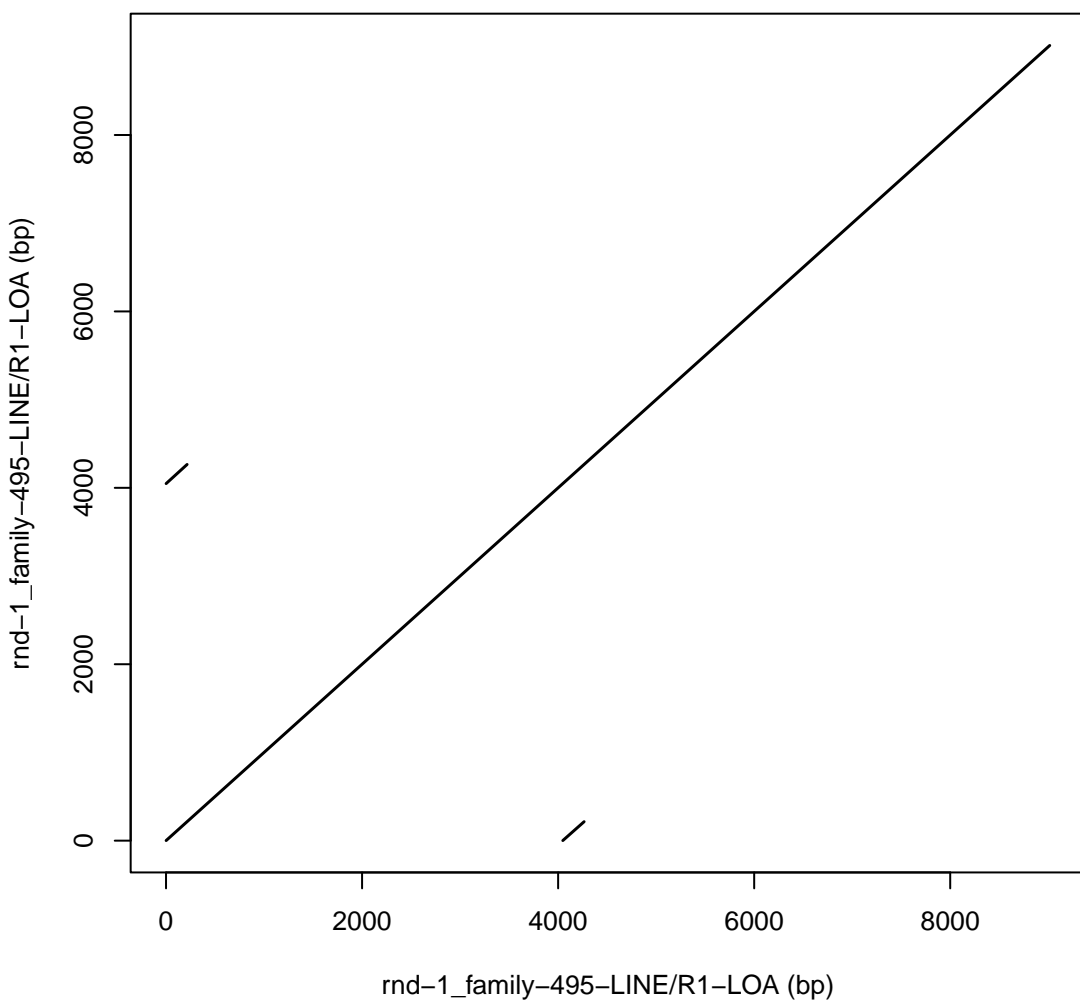
TE: rnd-1\_family-495-LINE/R1-LOA  
consensus size: 9015bp; fragments: 872; full length: 2 (>=8113.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

