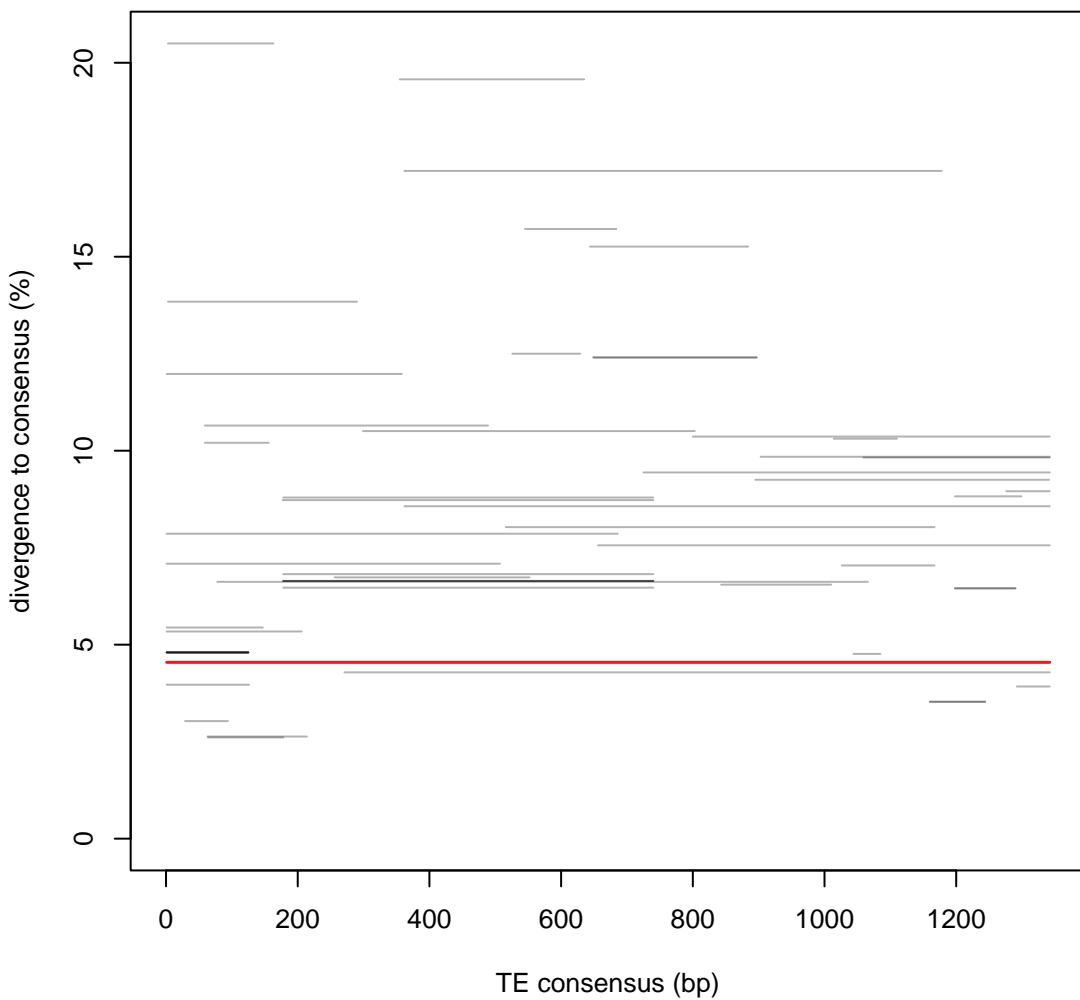
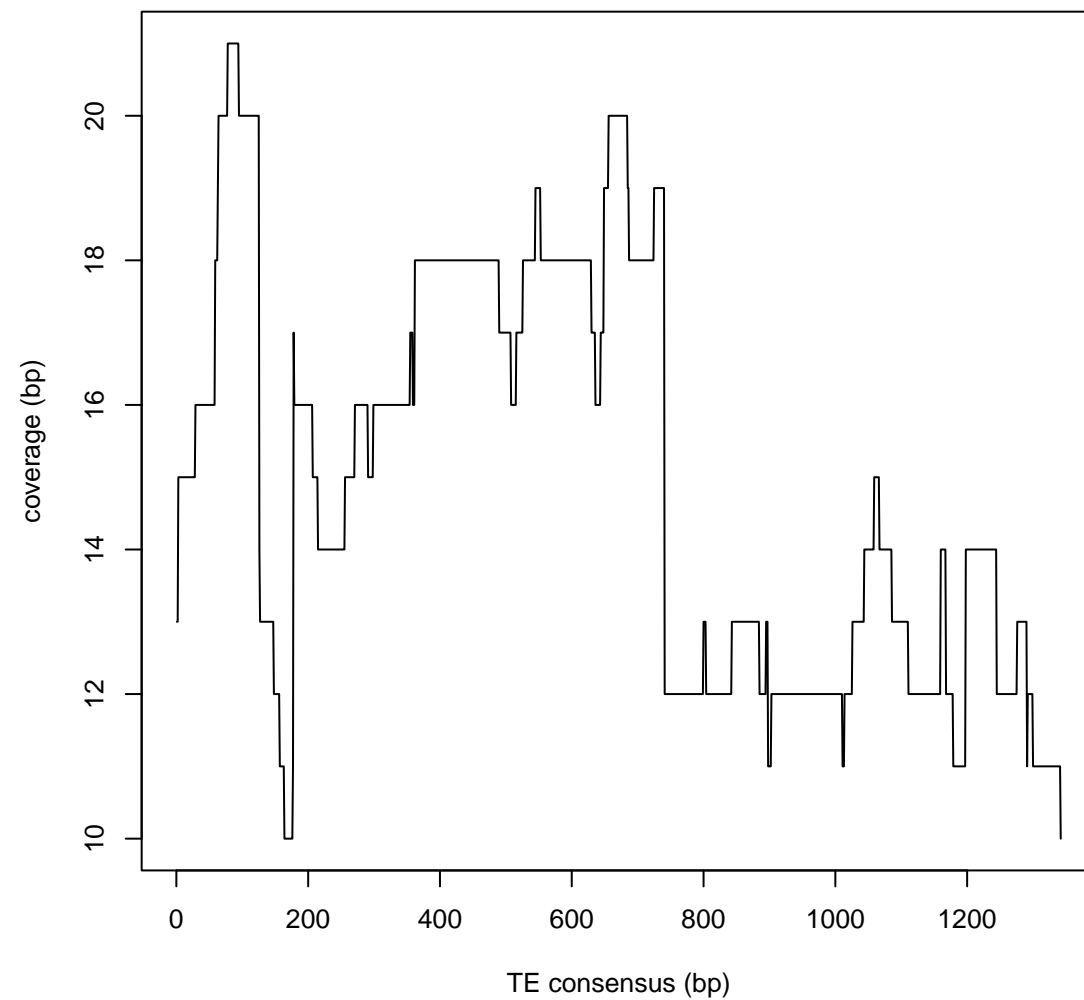


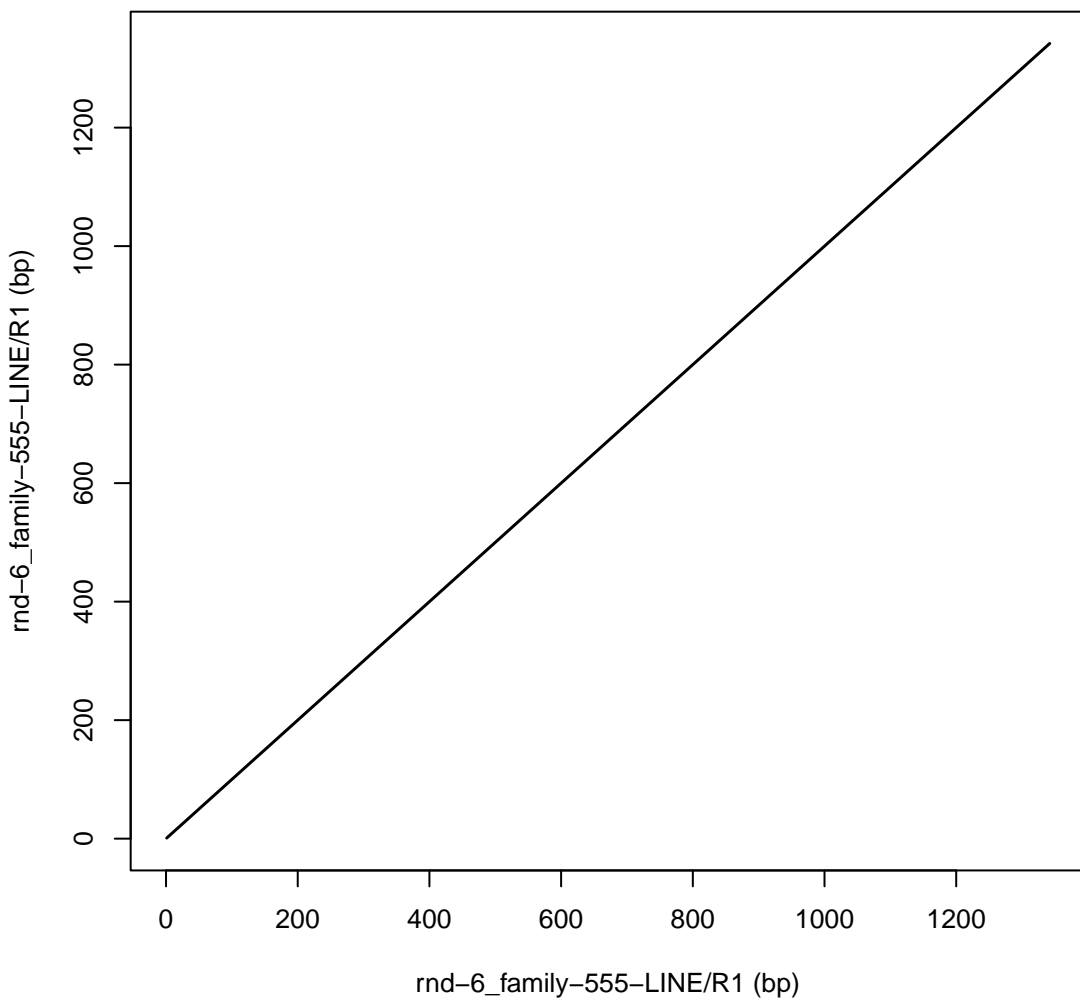
TE: rnd-6_family-555-LINE/R1
consensus size: 1342bp; fragments: 57; full length: 1 ($\geq 1207.8\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

