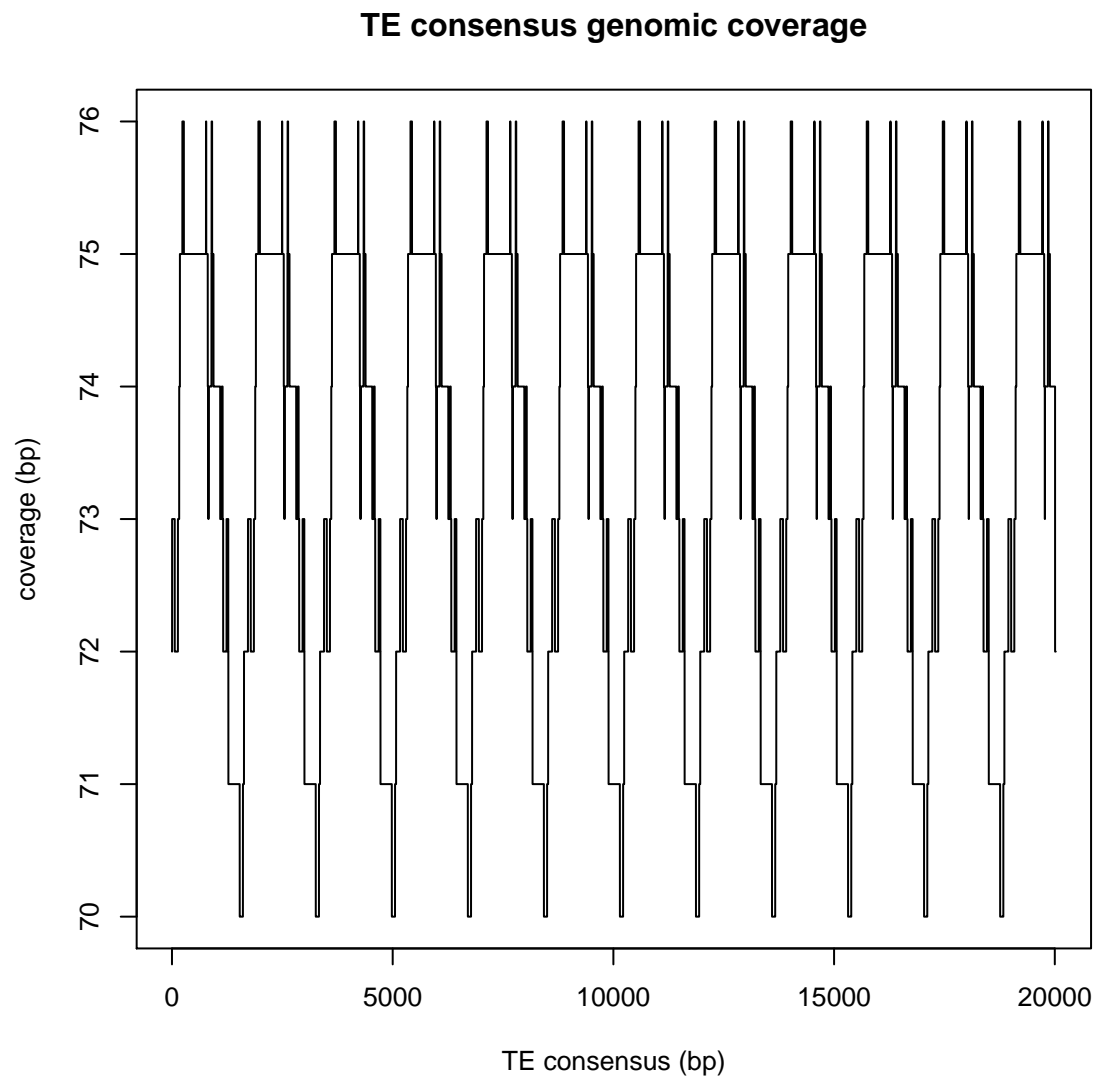
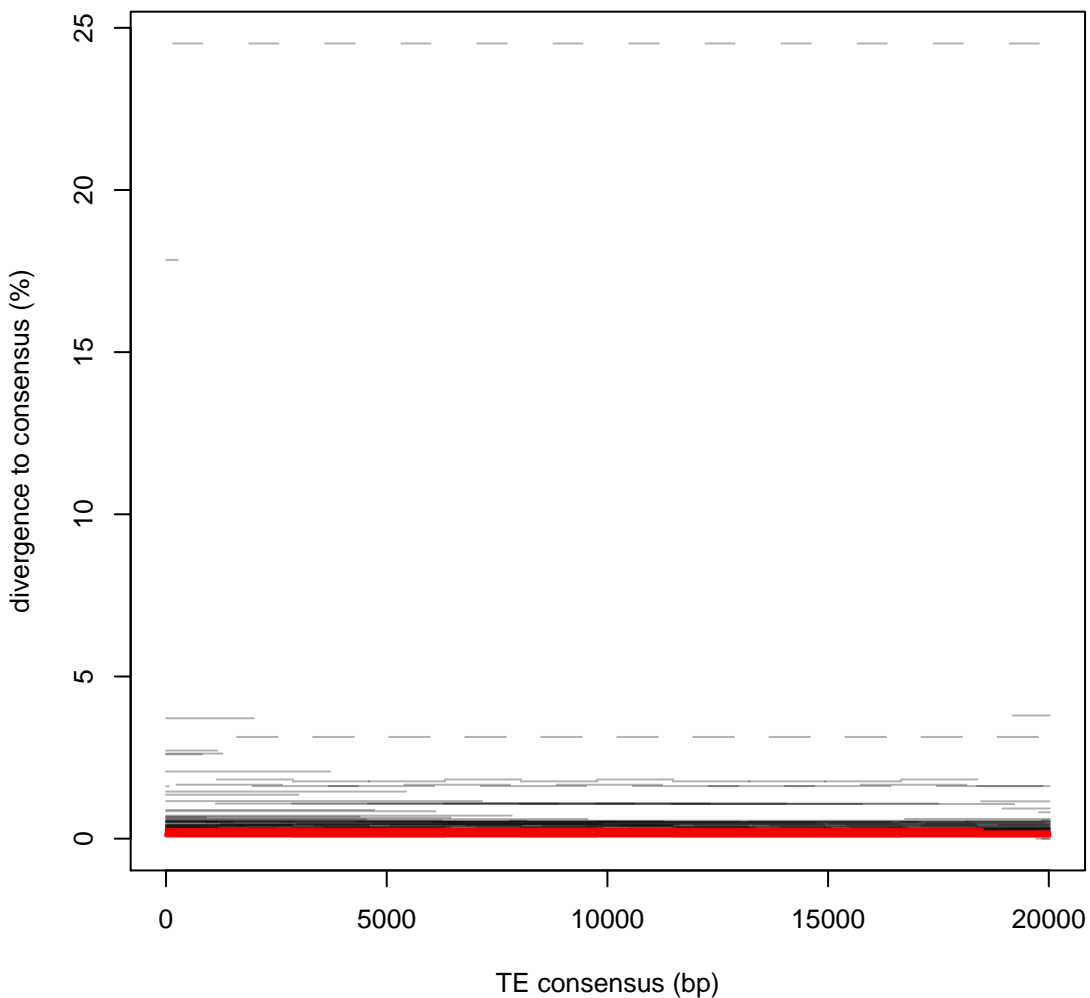
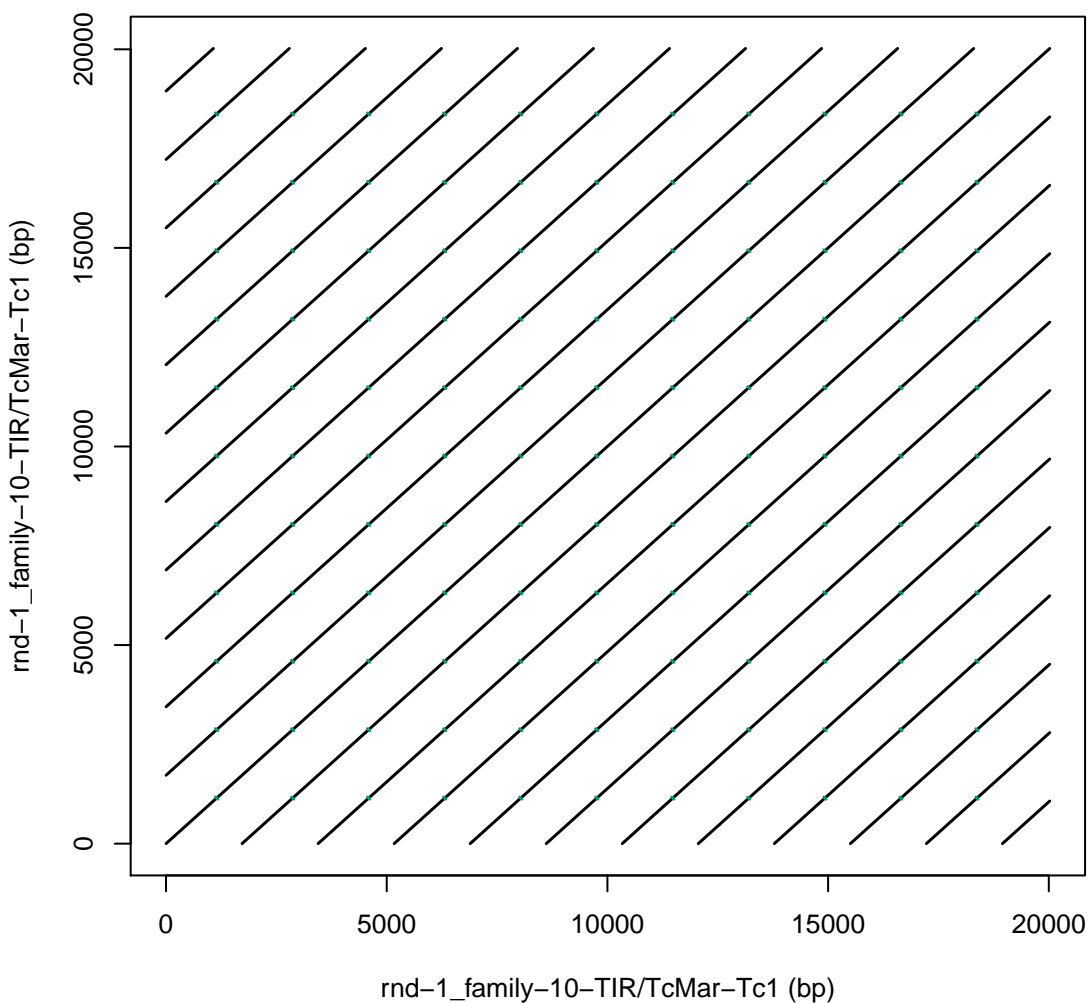


TE: rnd-1\_family-10-TIR/TcMar-Tc1  
consensus size: 20021bp; fragments: 301; full length: 15 ( $\geq 18018.9\text{bp}$ )



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

