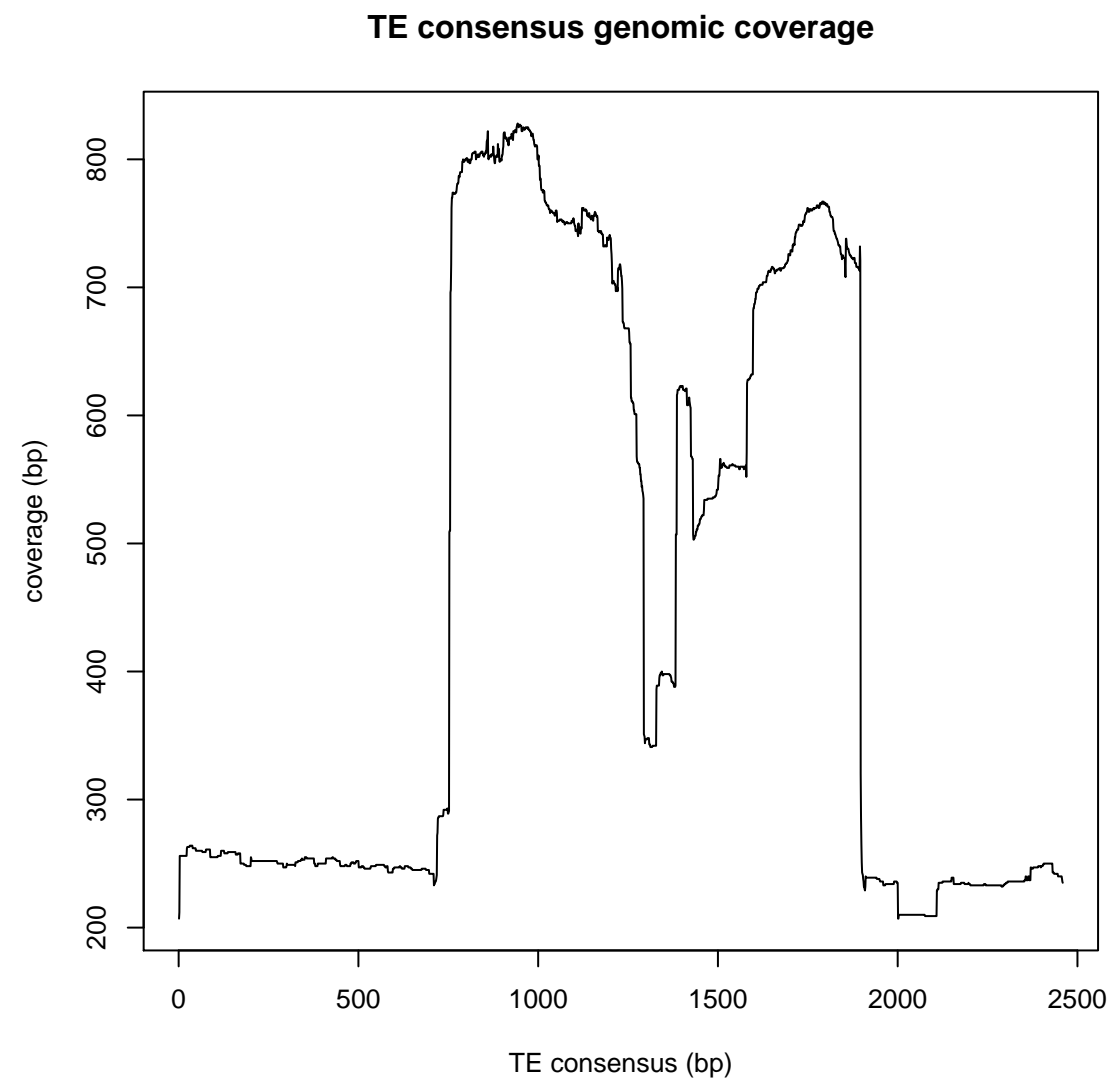
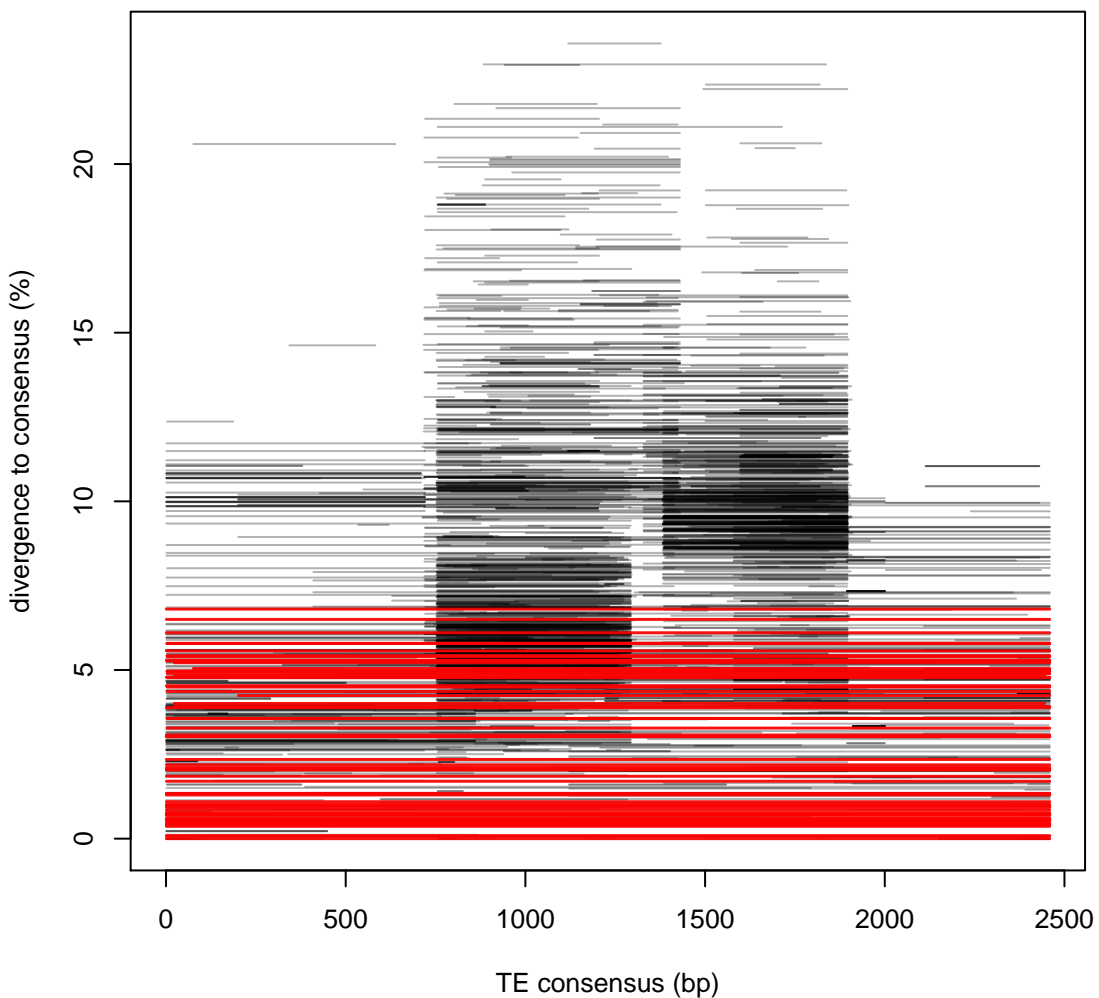
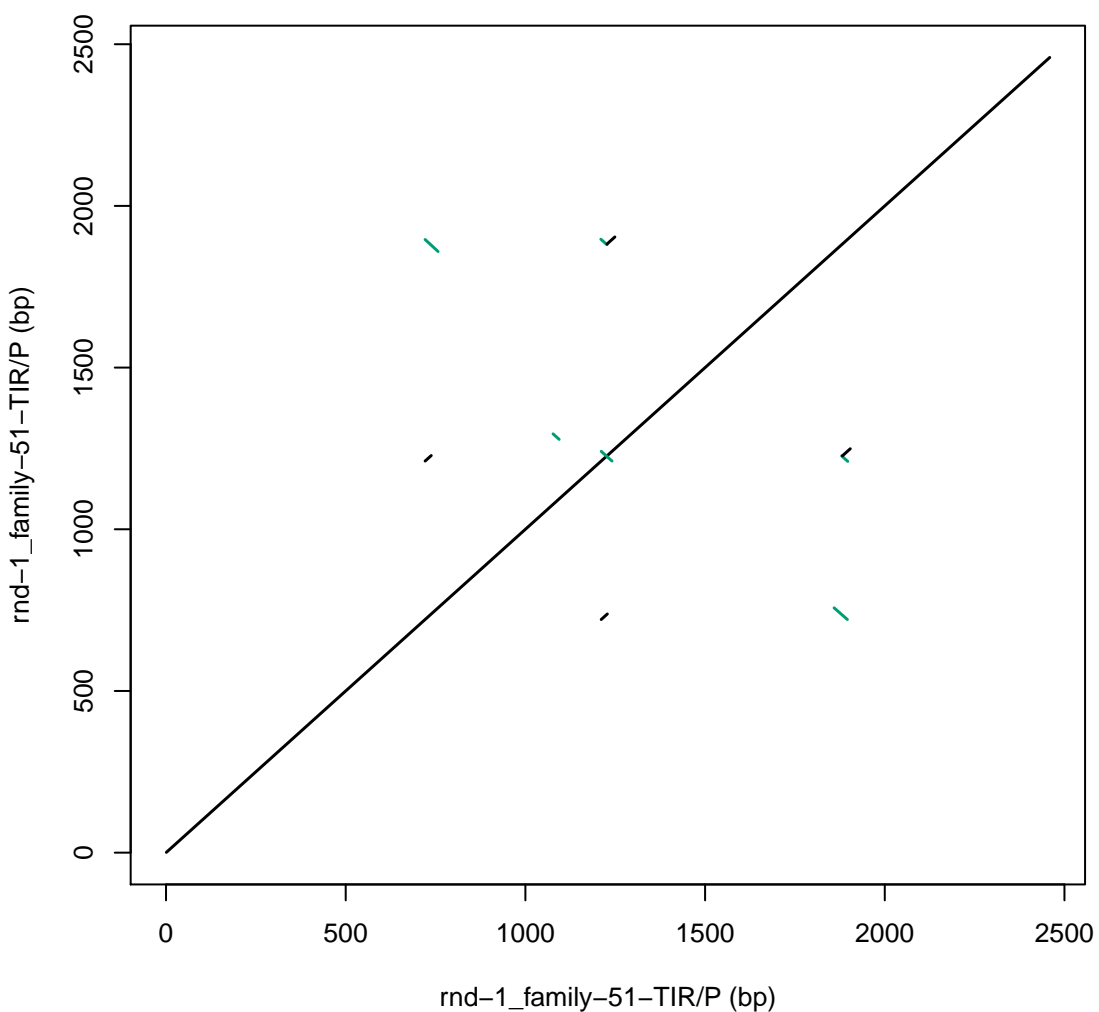


TE: rnd-1\_family-51-TIR/P  
consensus size: 2459bp; fragments: 2016; full length: 65 (>=2213.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

