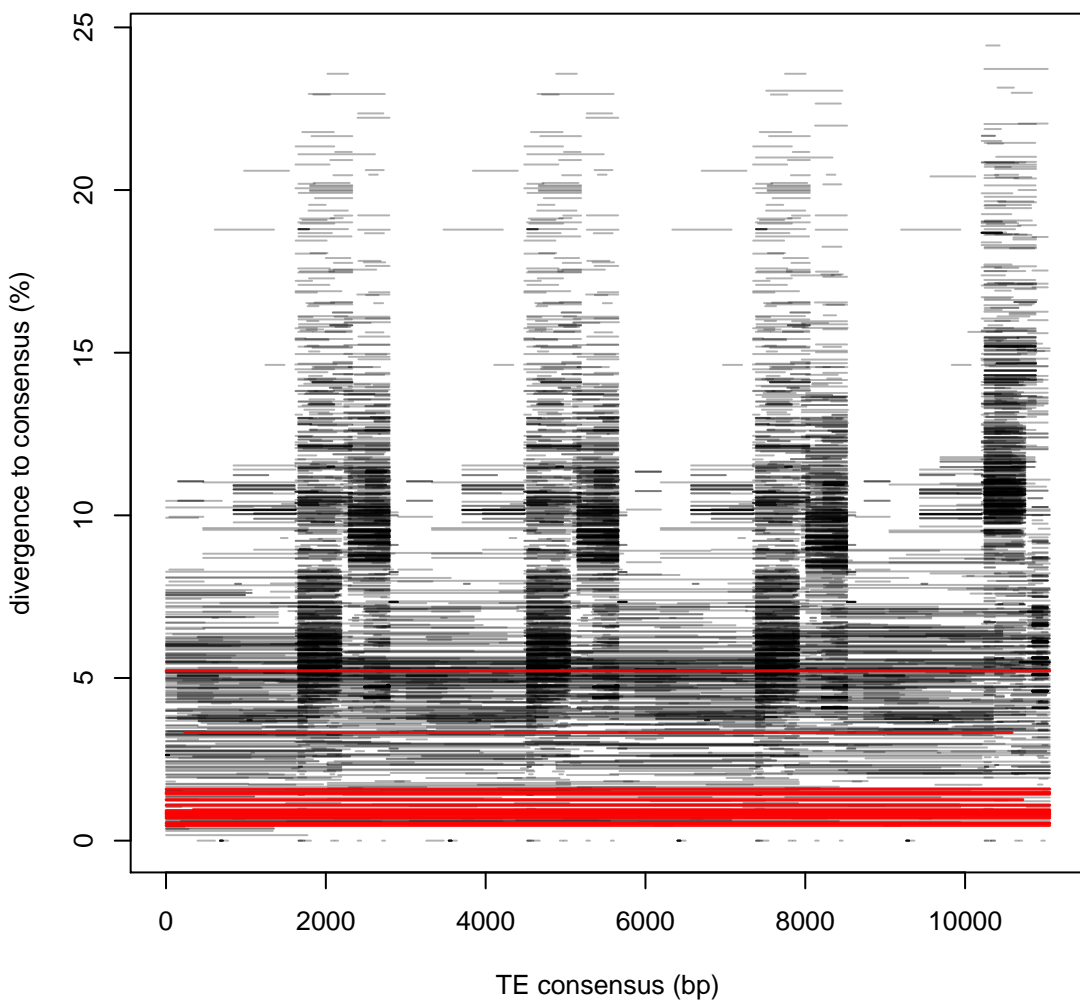
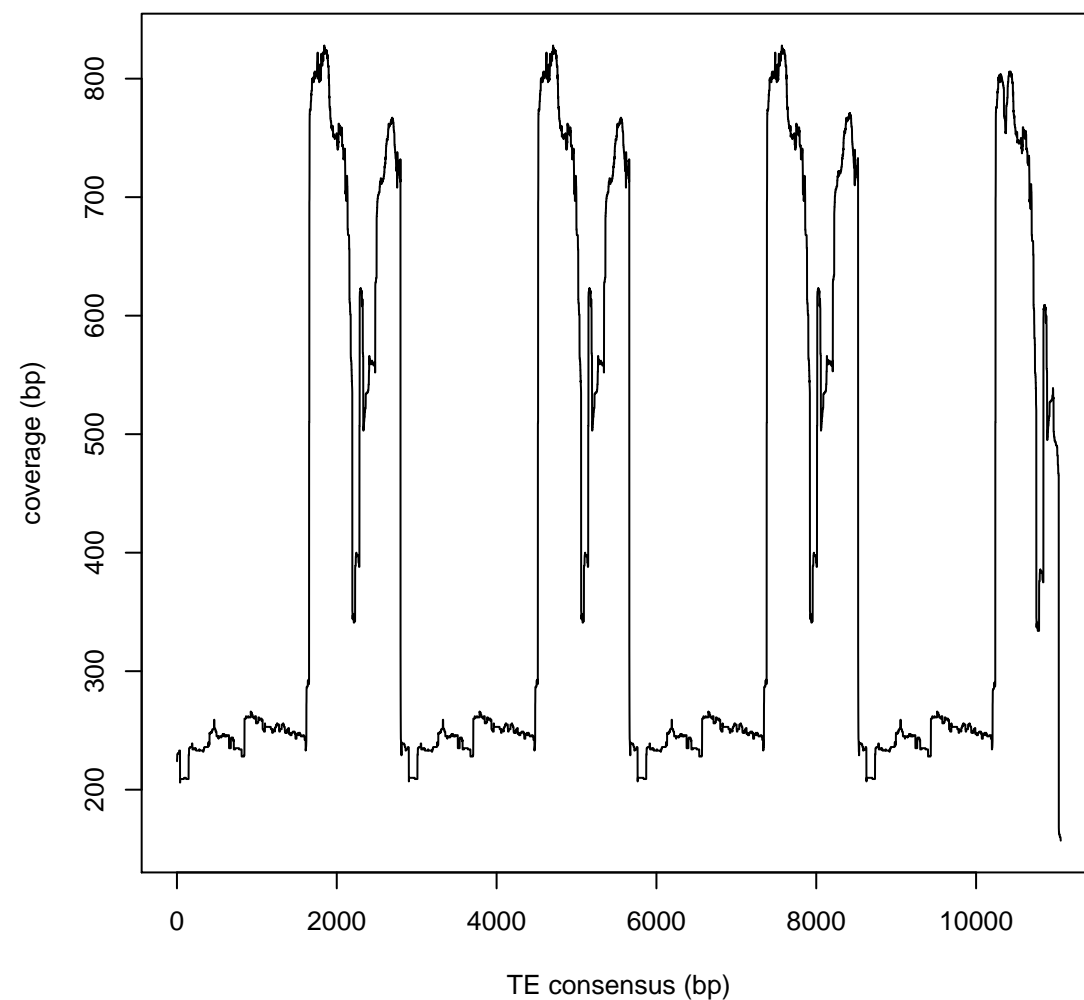


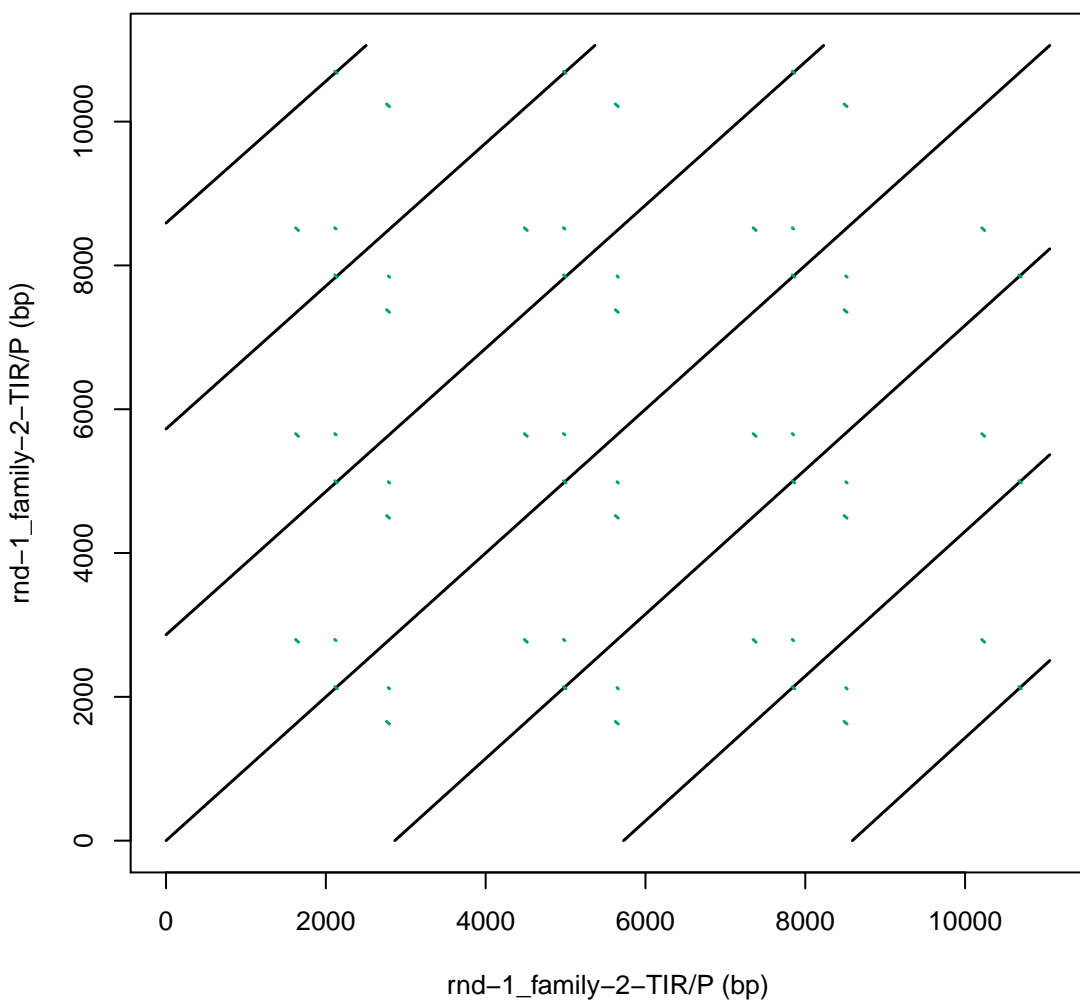
TE: rnd-1\_family-2-TIR/P  
consensus size: 11059bp; fragments: 7108; full length: 17 (>=9953.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

