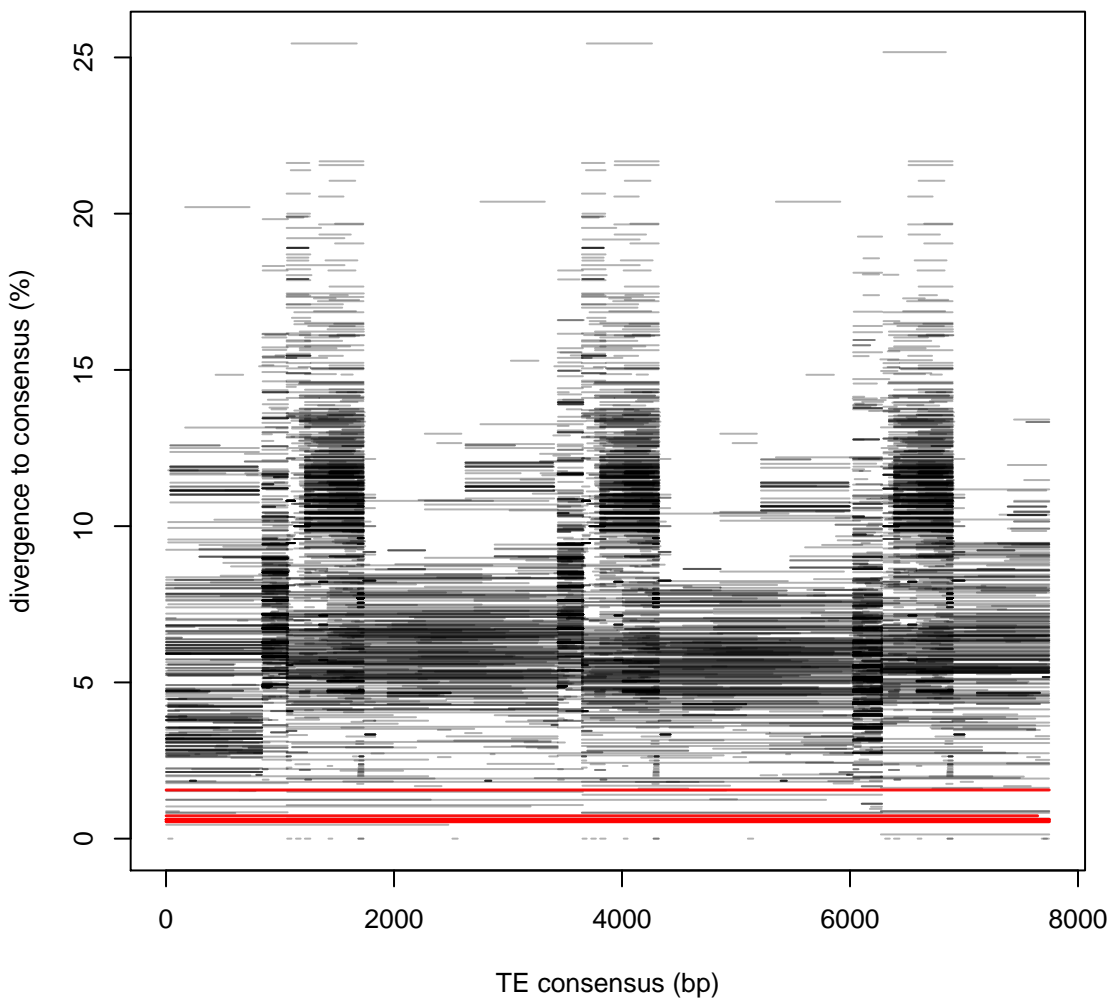
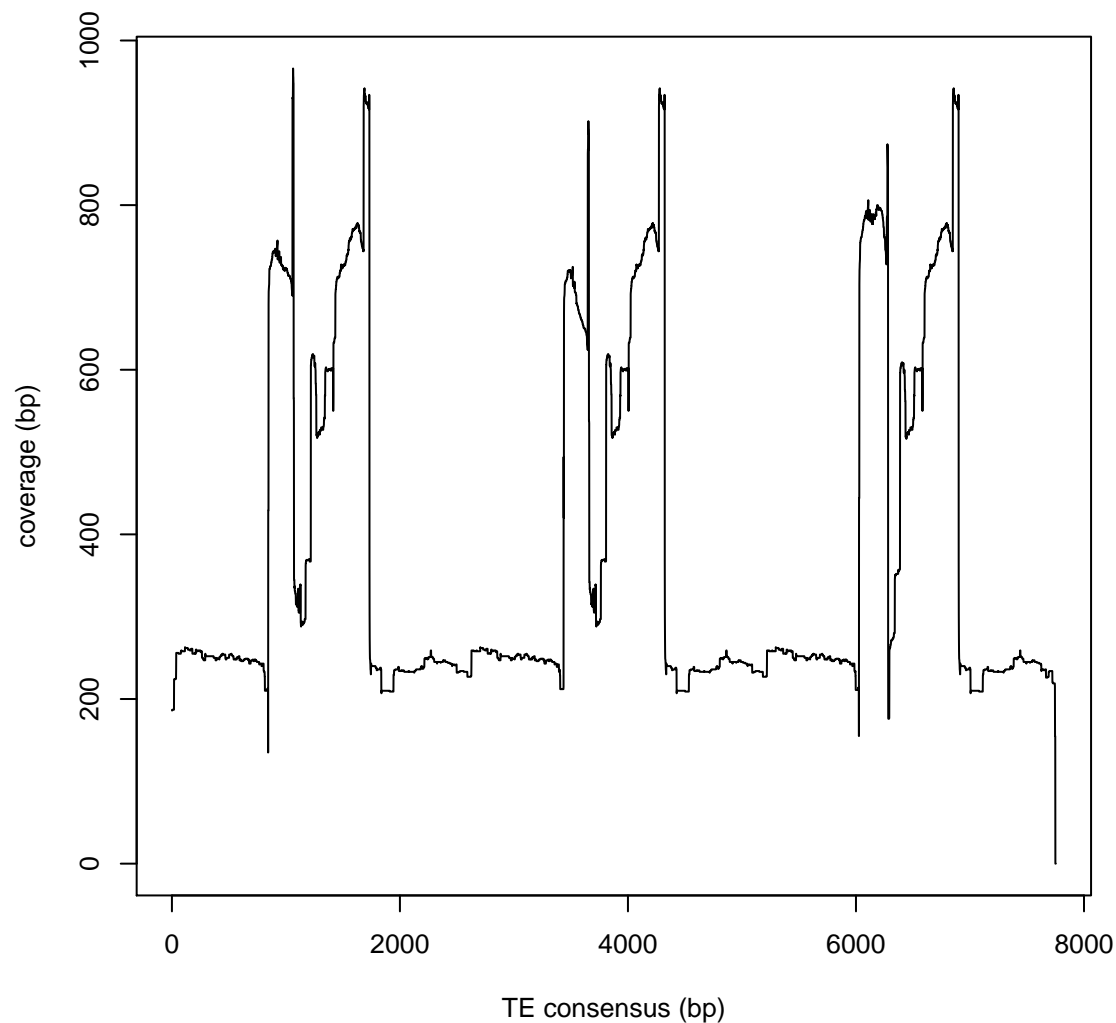


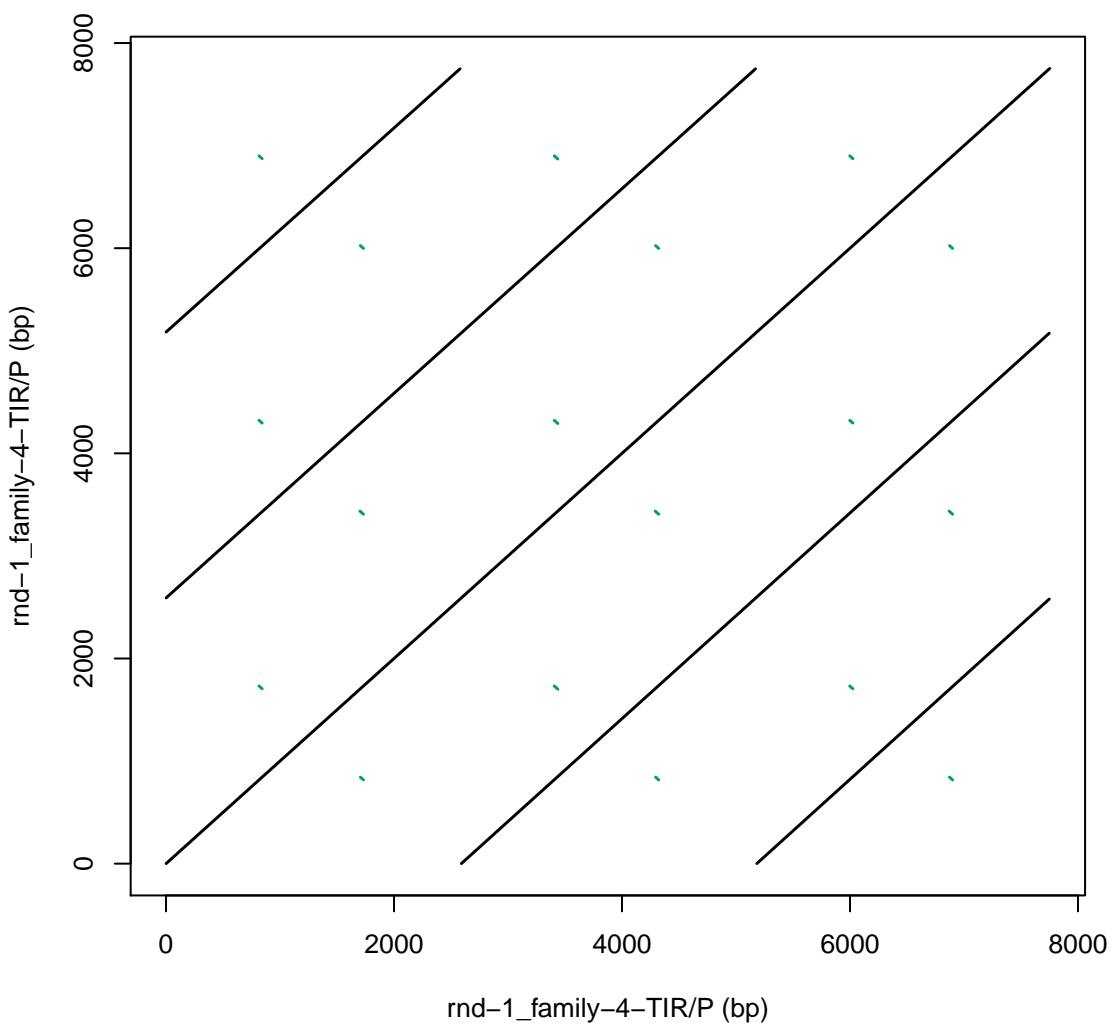
TE: rnd-1_family-4-TIR/P
consensus size: 7752bp; fragments: 6909; full length: 7 (>=6976.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

