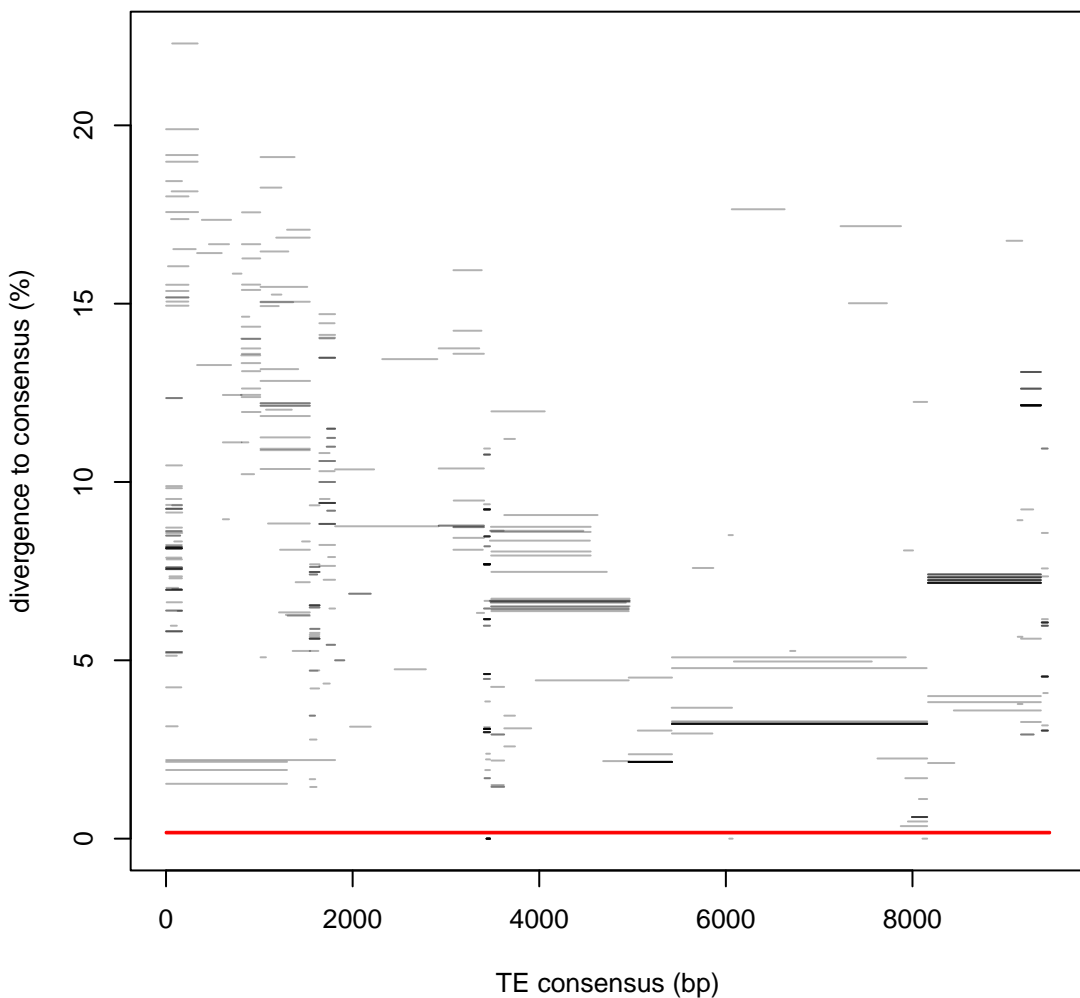
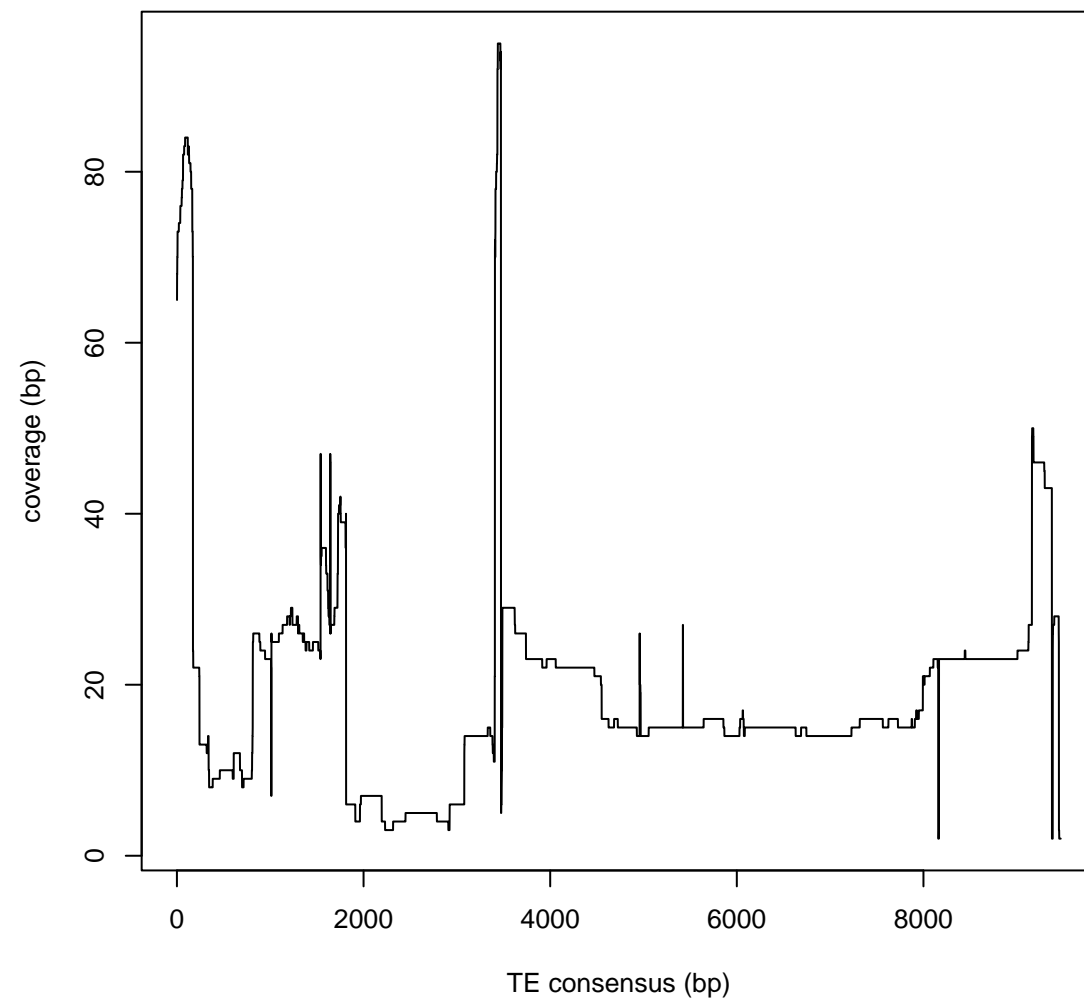


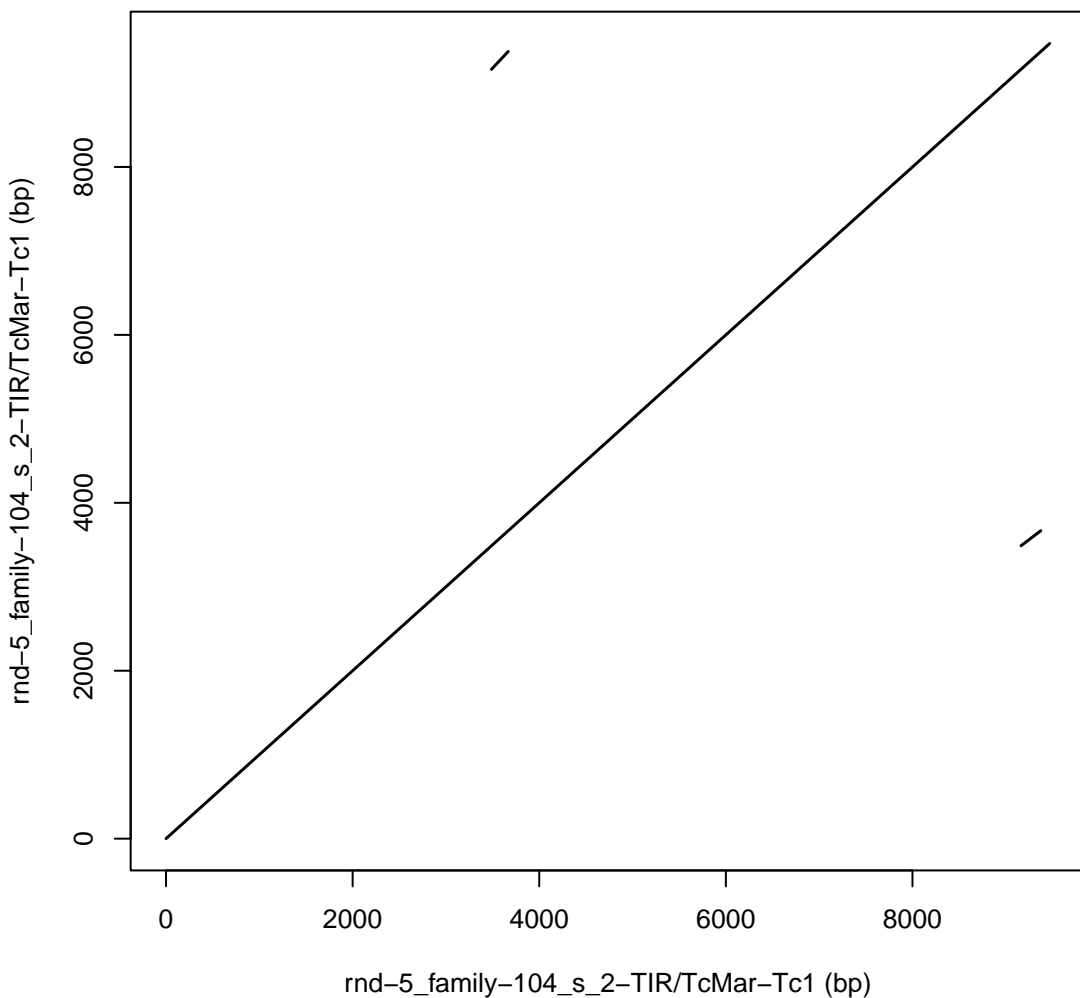
TE: rnd-5_family-104_s_2-TIR/TcMar-Tc1
consensus size: 9471bp; fragments: 486; full length: 2 (>=8523.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

