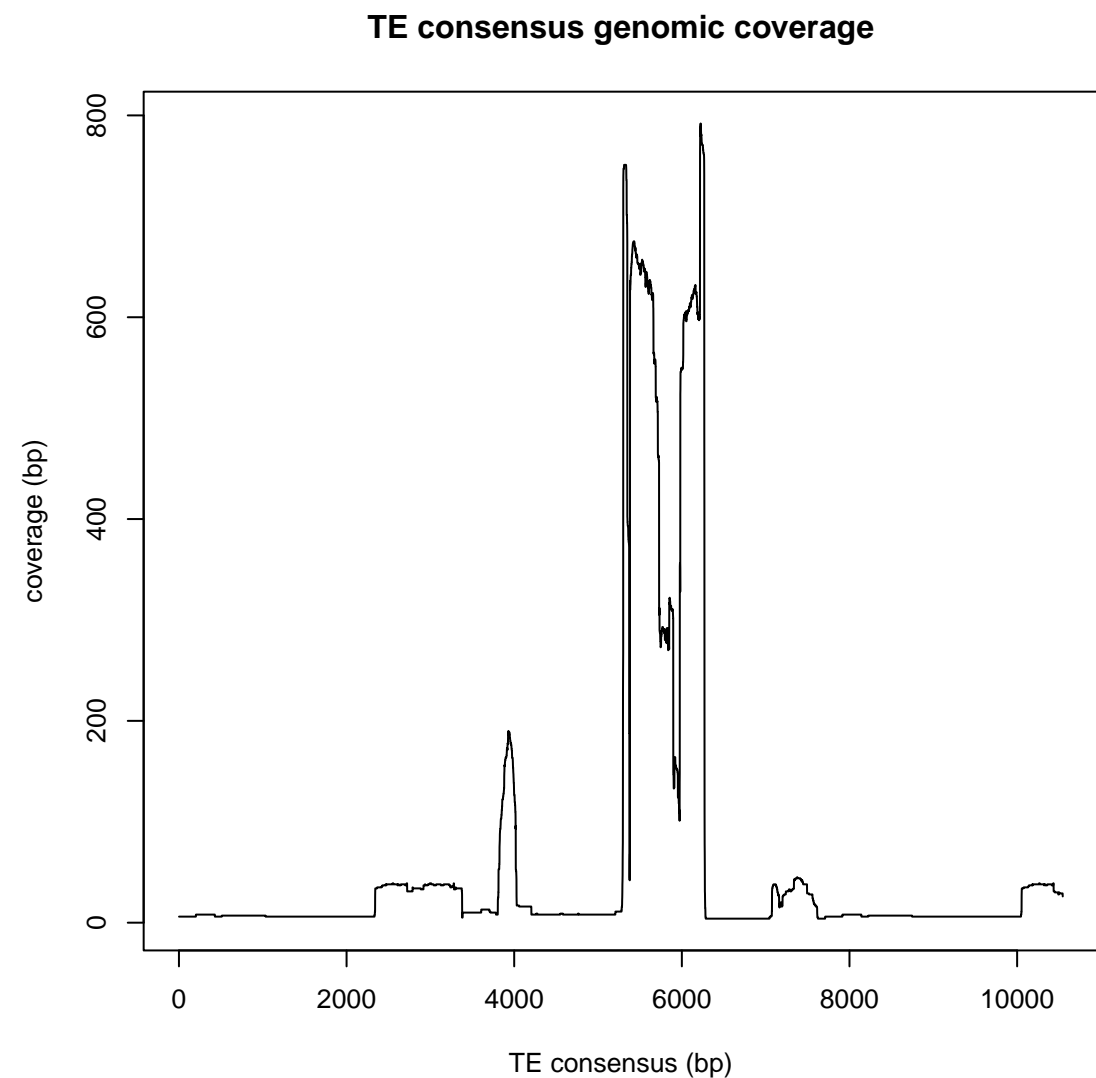
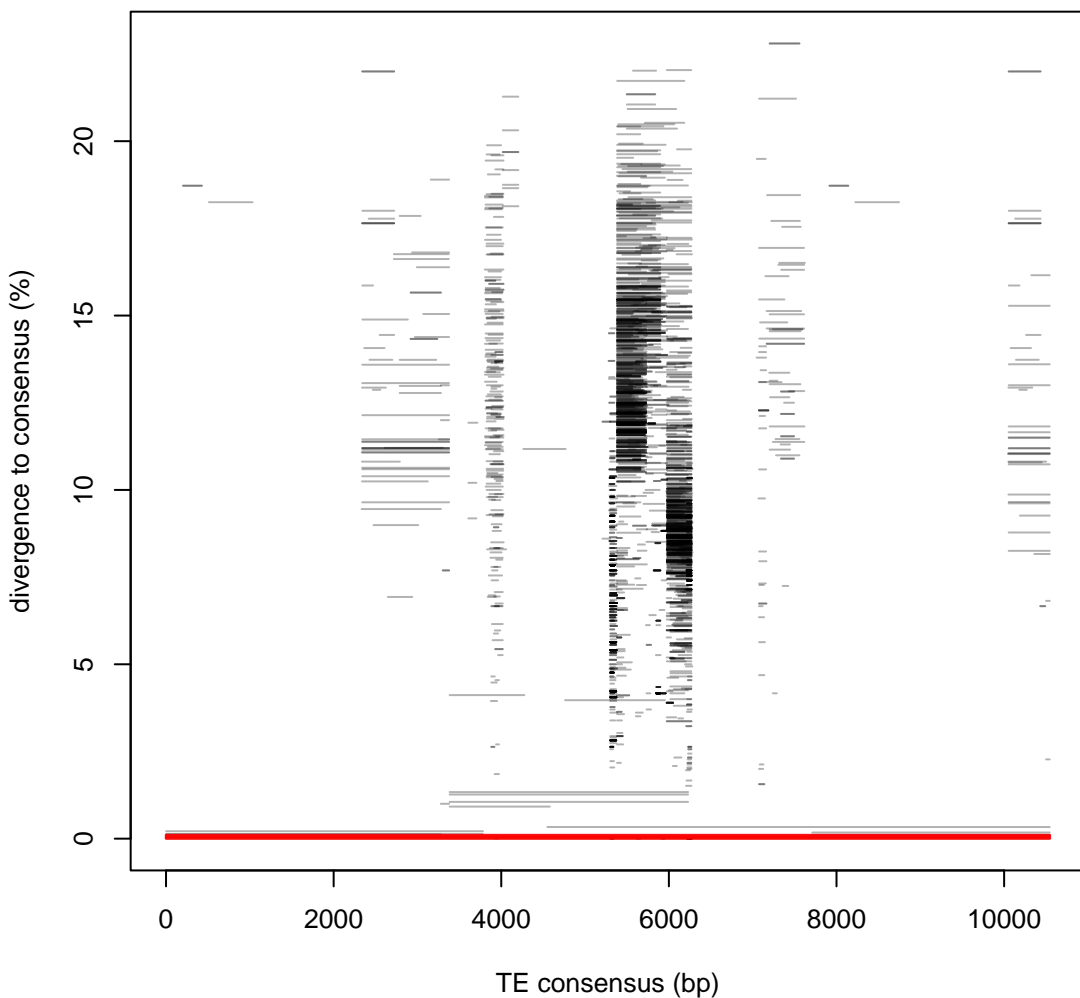
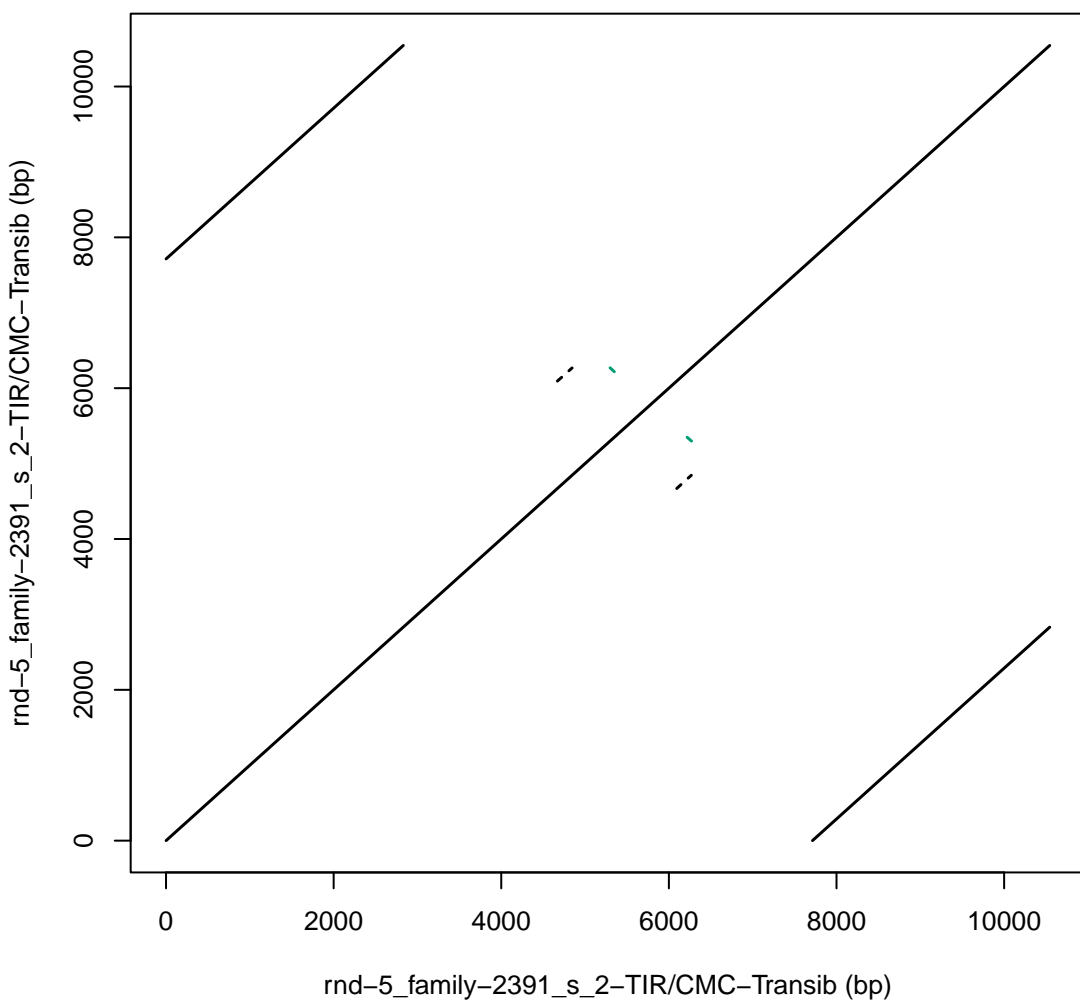


TE: rnd-5_family-2391_s_2-TIR/CMC-Transib
consensus size: 10544bp; fragments: 2967; full length: 3 (>=9489.6bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

