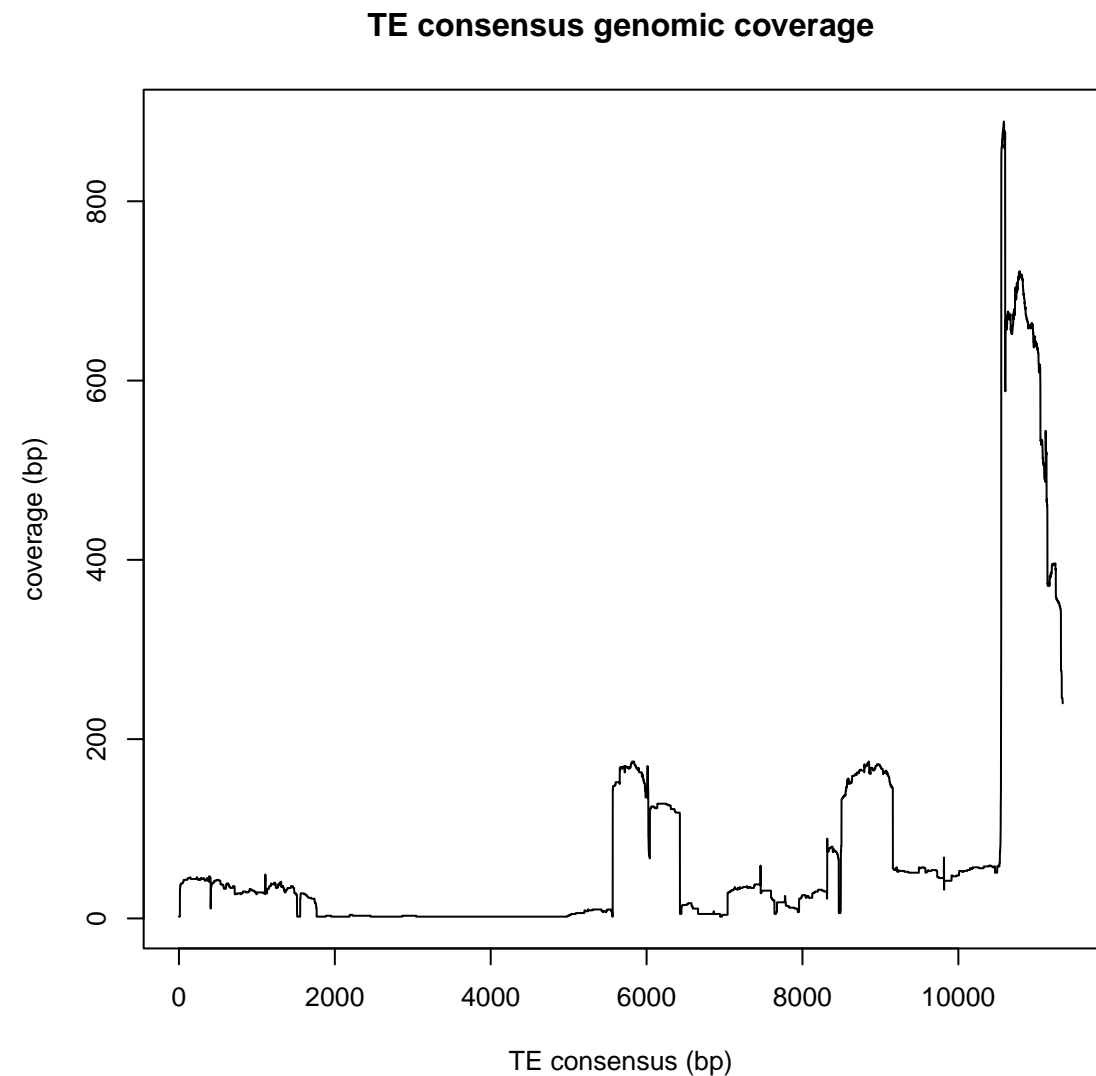
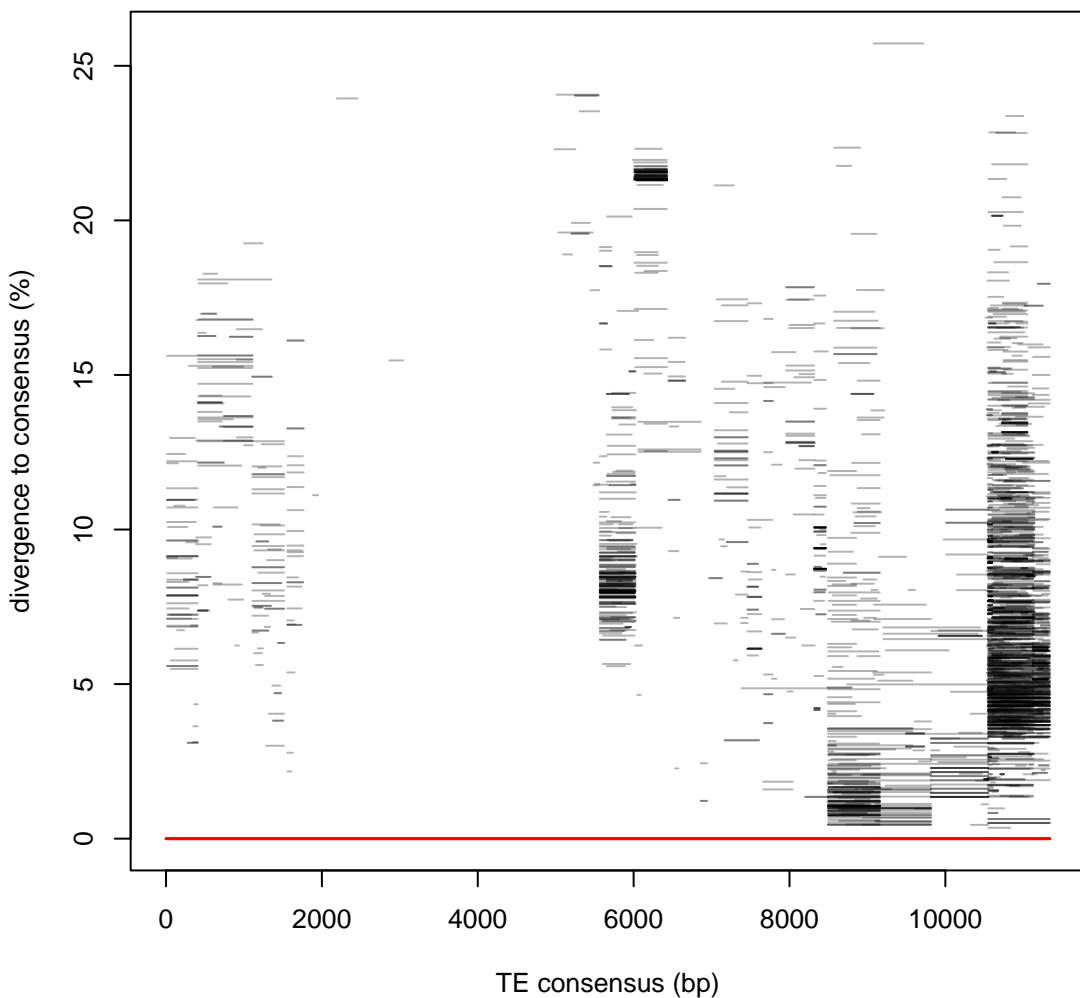
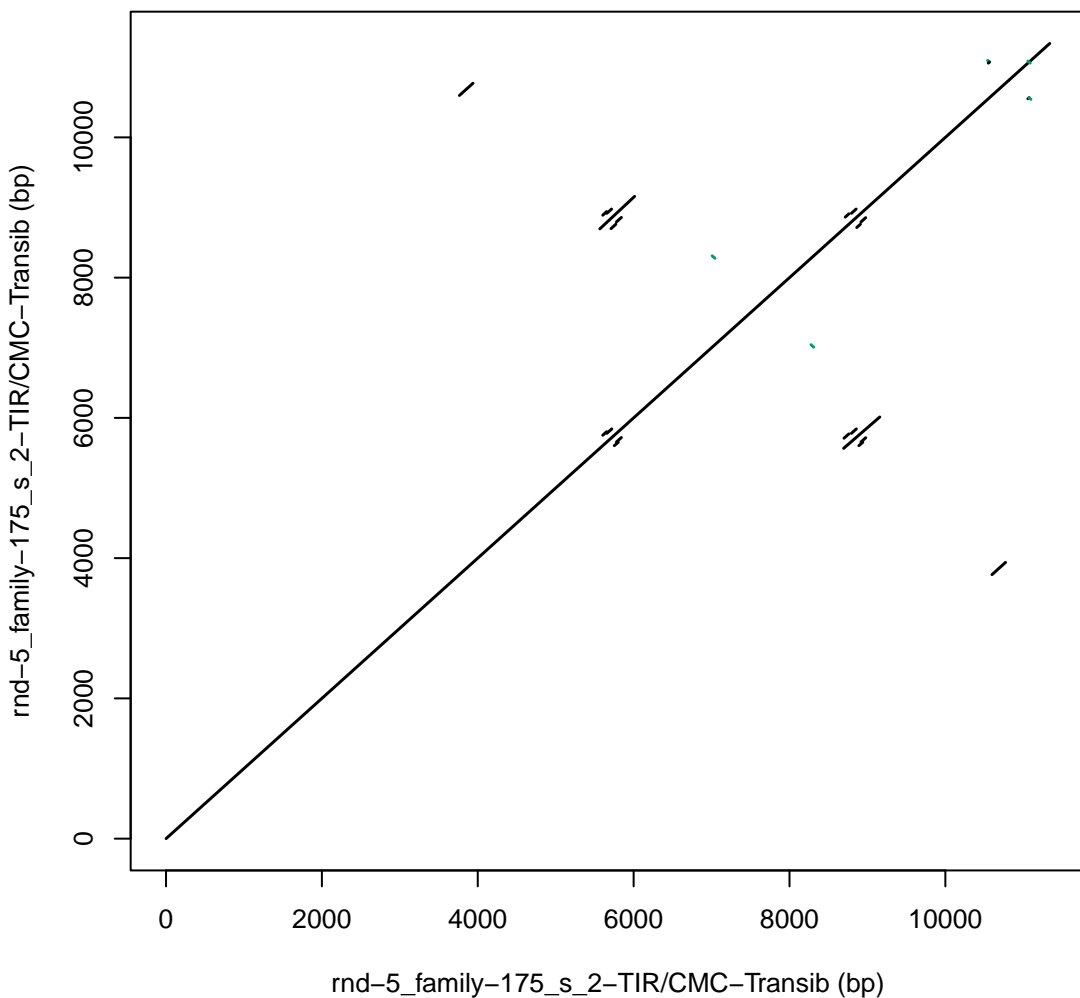


TE: rnd-5_family-175_s_2-TIR/CMC-Transib
consensus size: 11339bp; fragments: 2583; full length: 2 (>=10205.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

