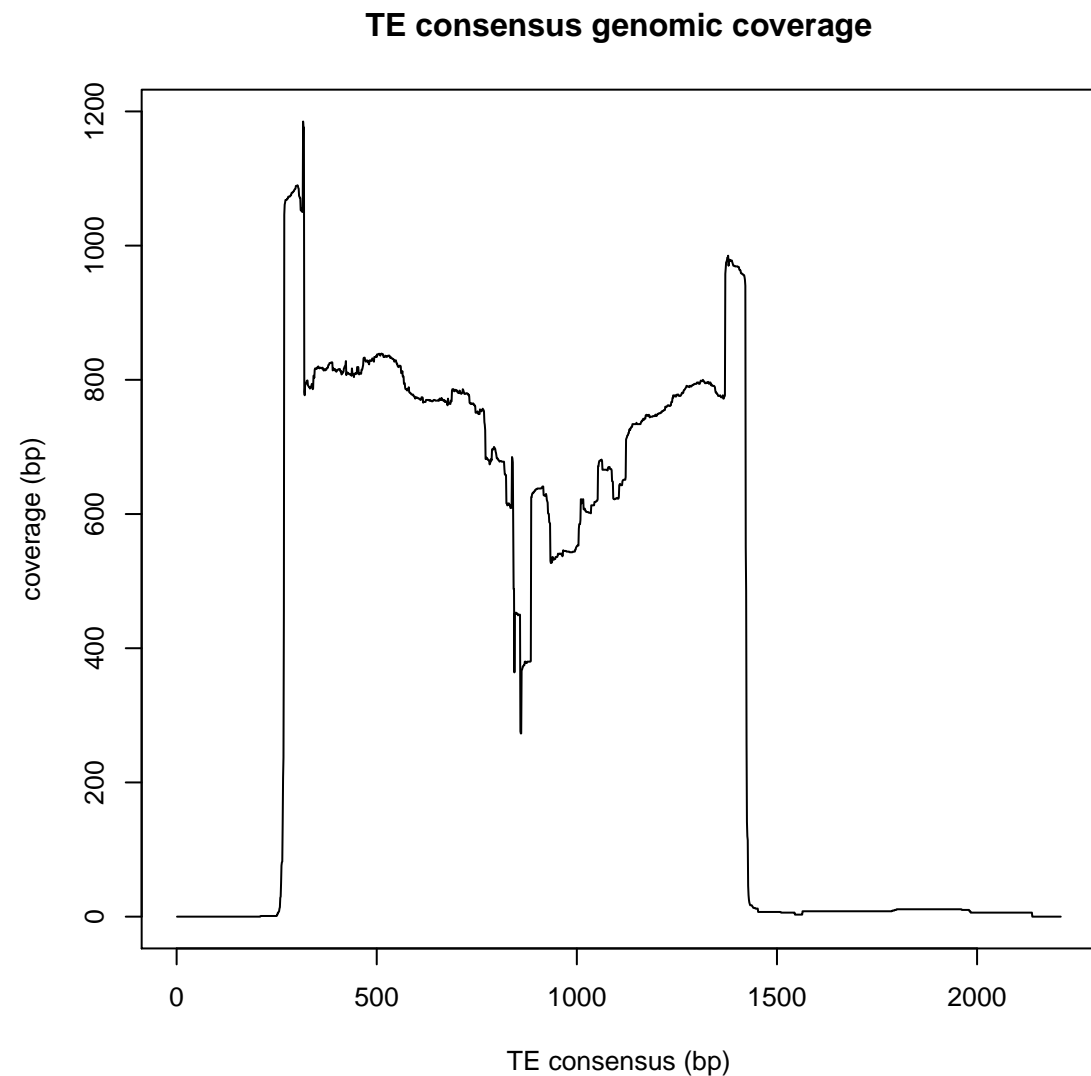
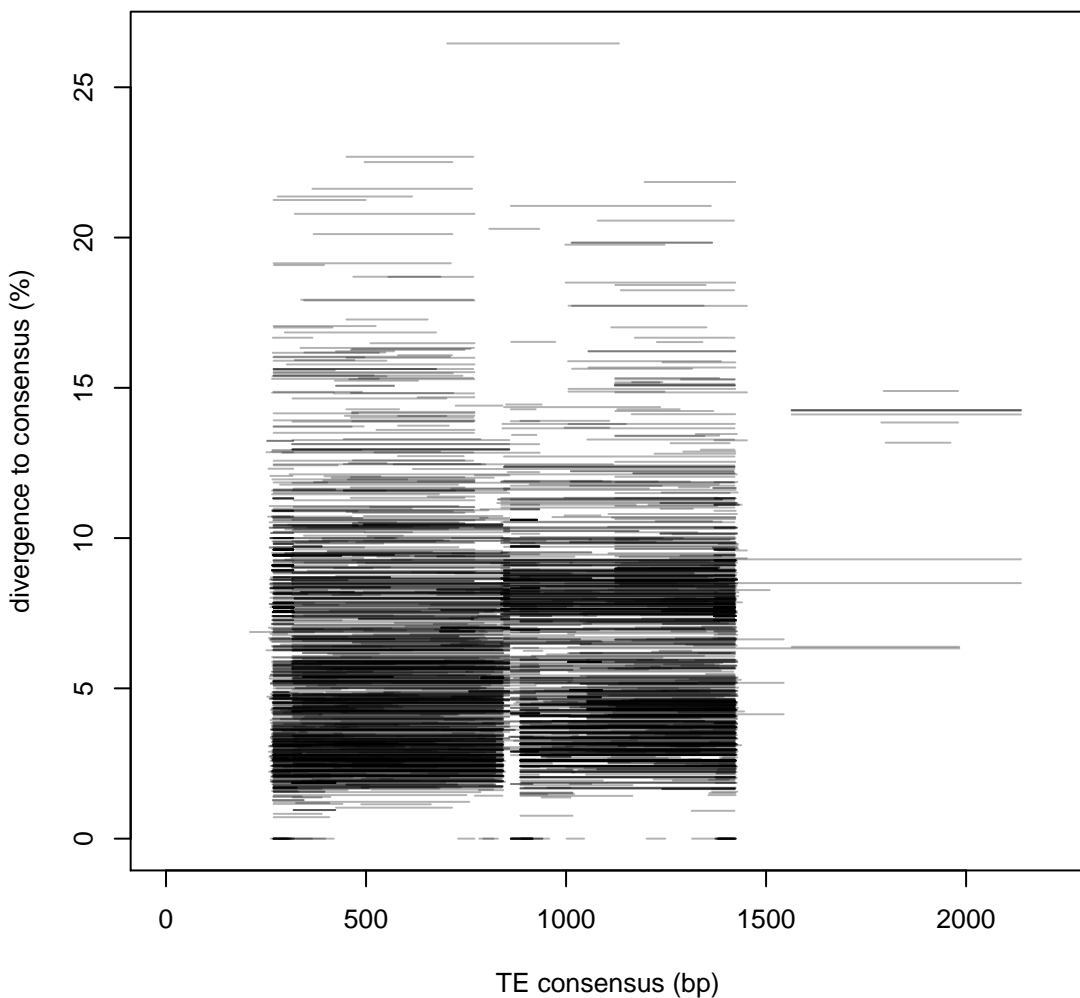
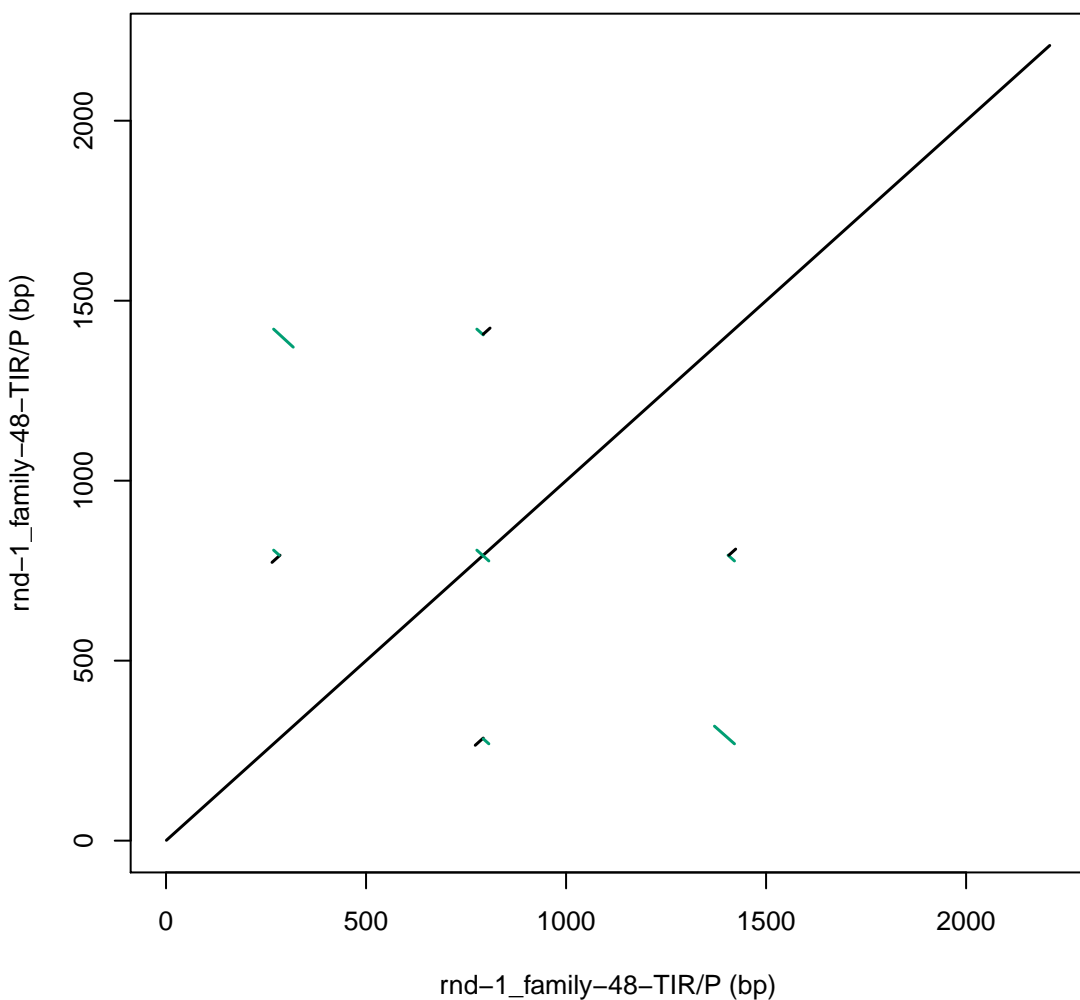


TE: rnd-1_family-48-TIR/P
consensus size: 2209bp; fragments: 3009; full length: 0 (>=1988.1bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

