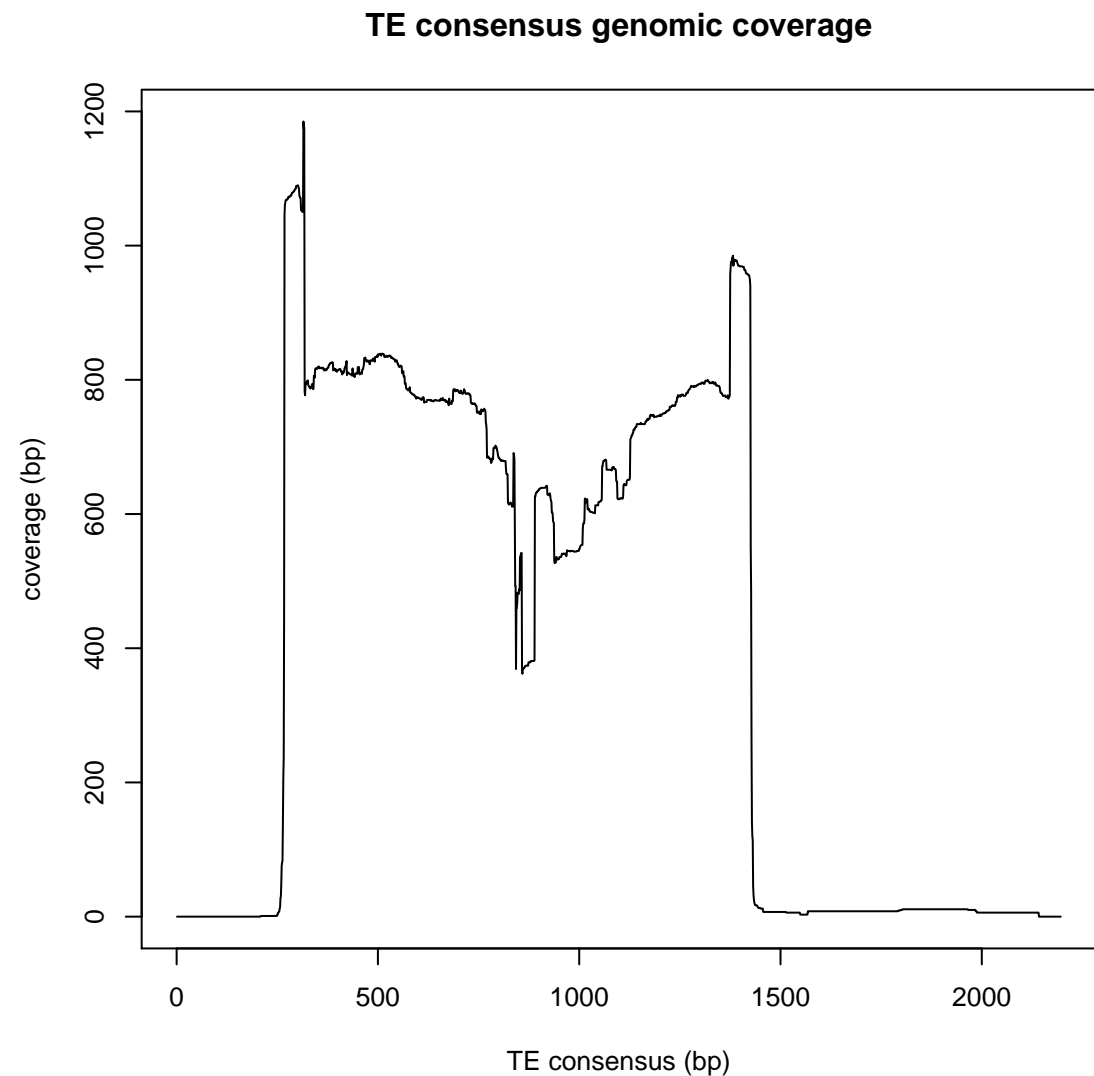
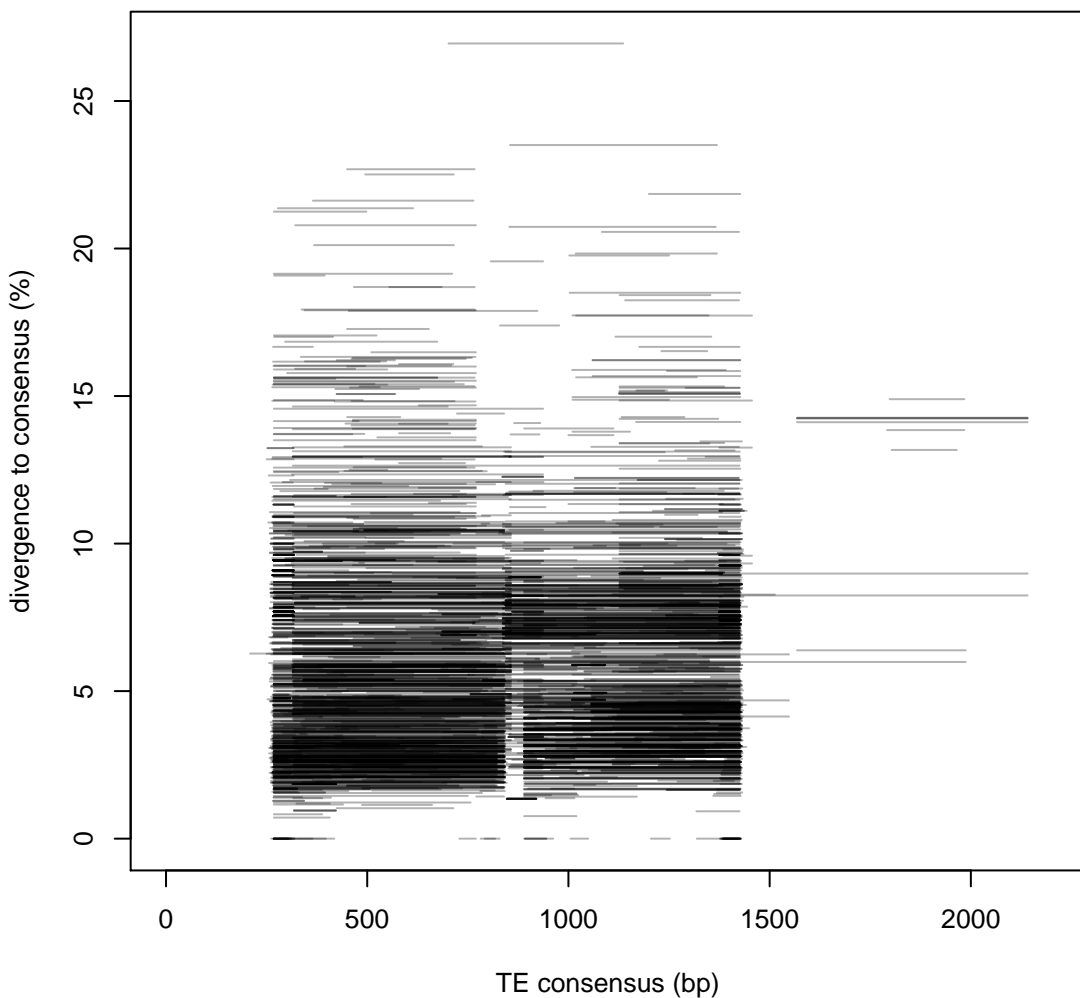
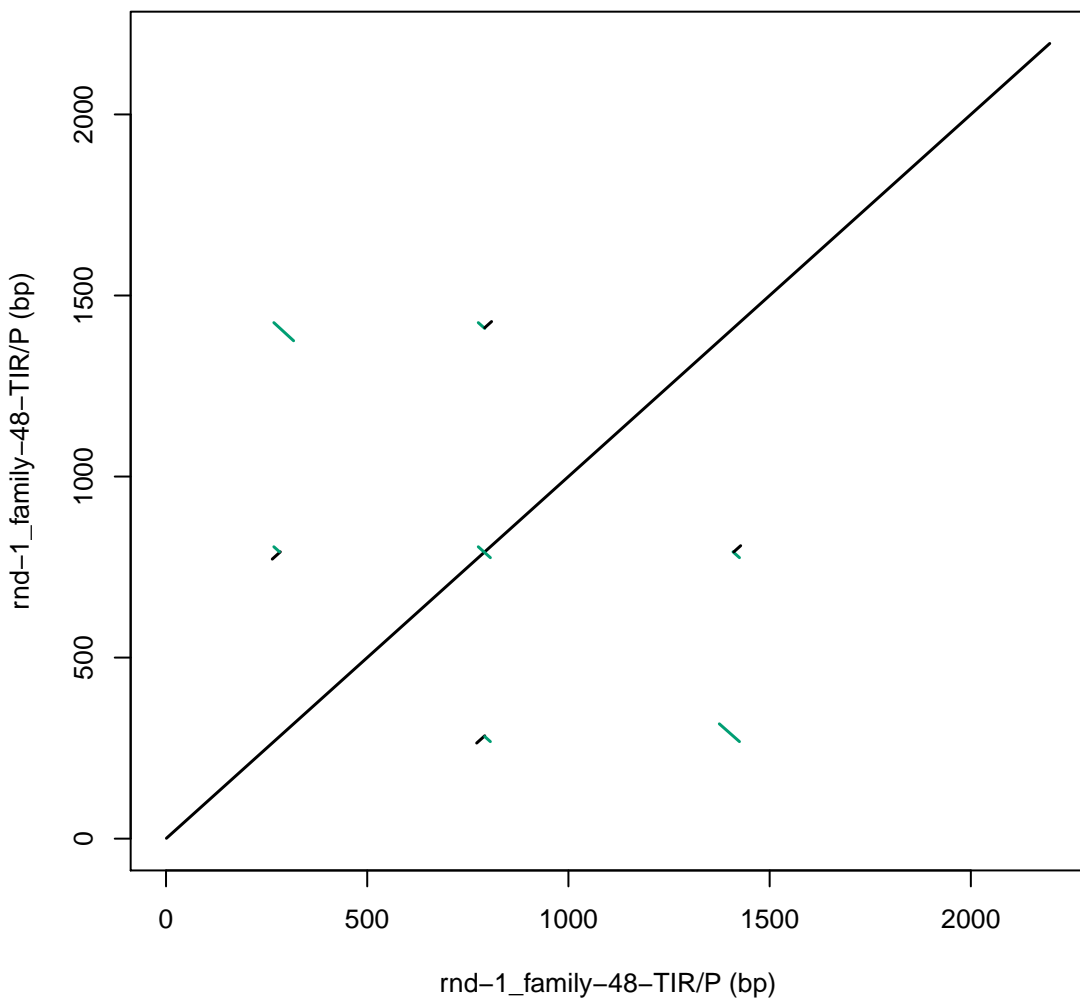


TE: rnd-1_family-48-TIR/P
consensus size: 2196bp; fragments: 3010; full length: 0 (>=1976.4bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

