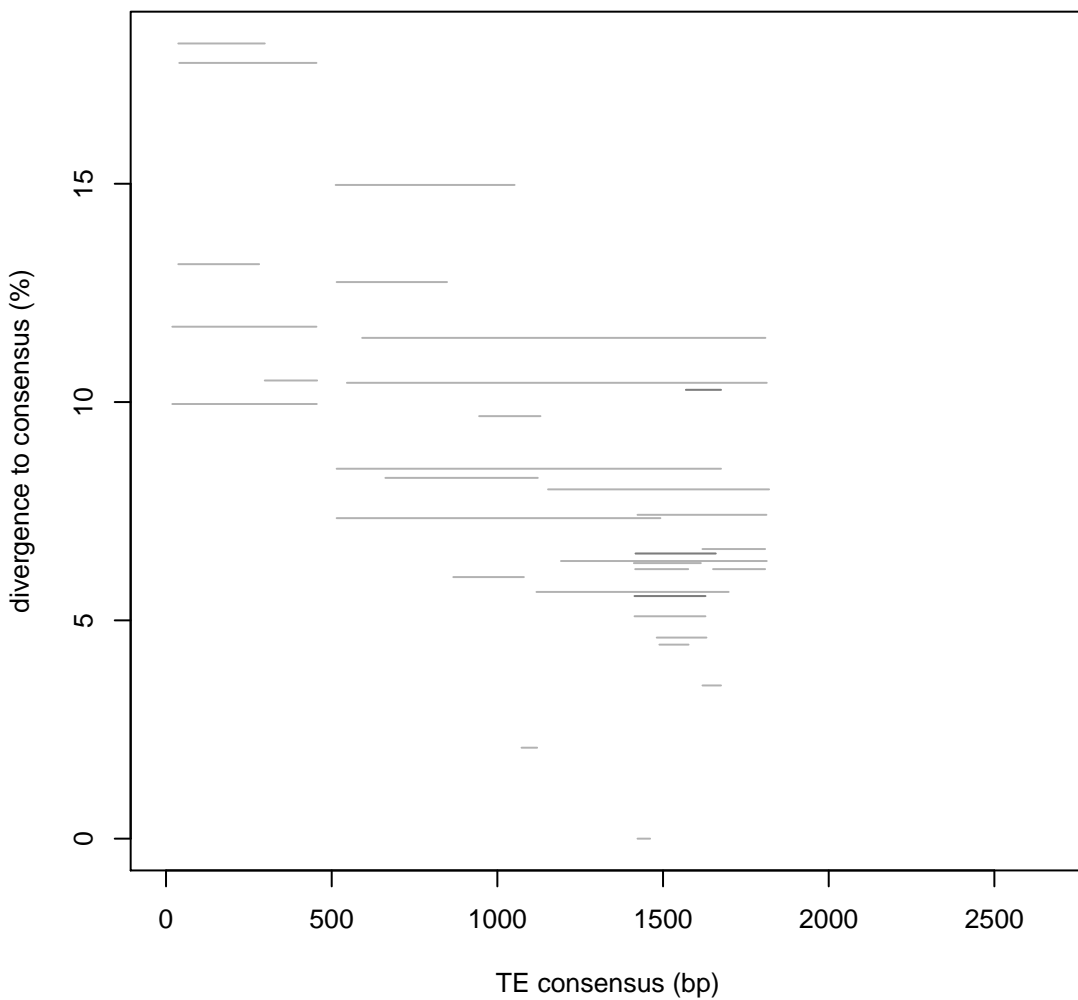
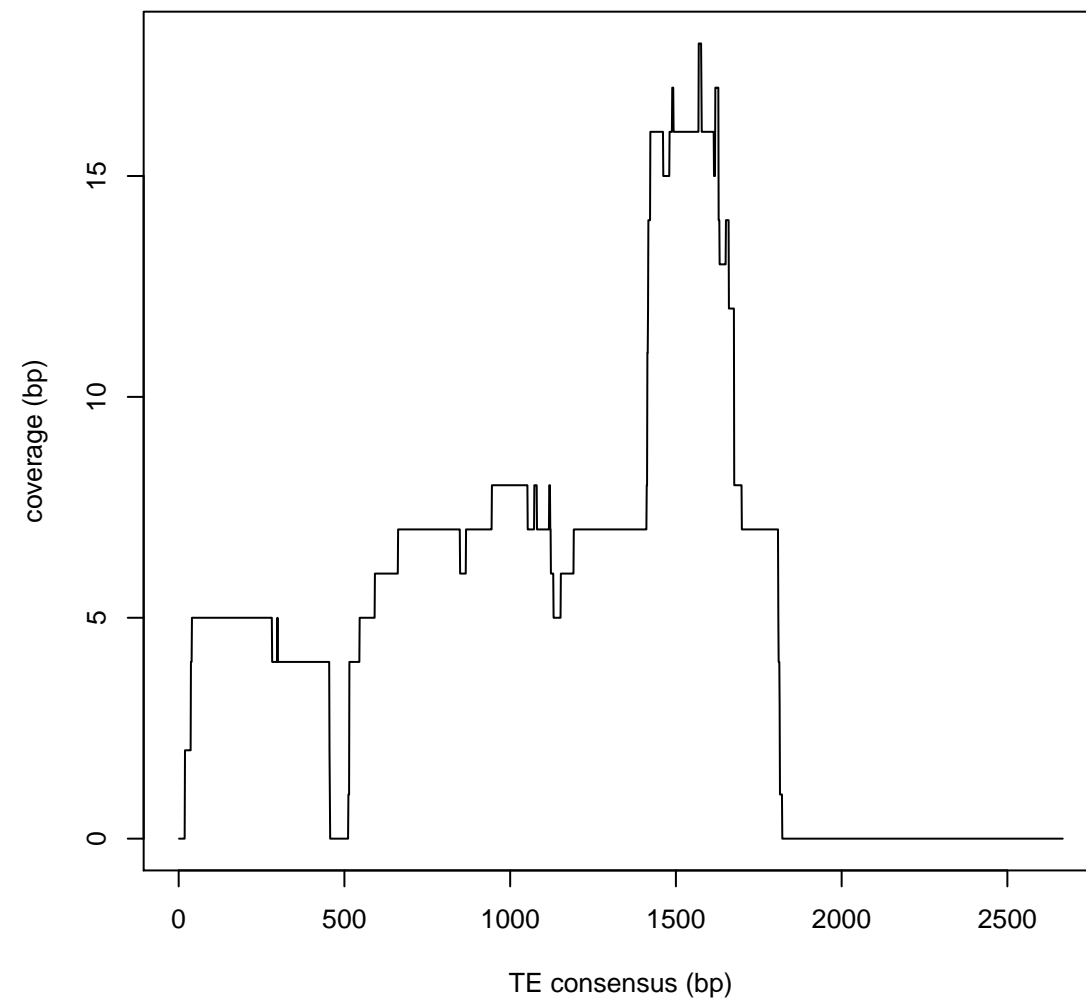


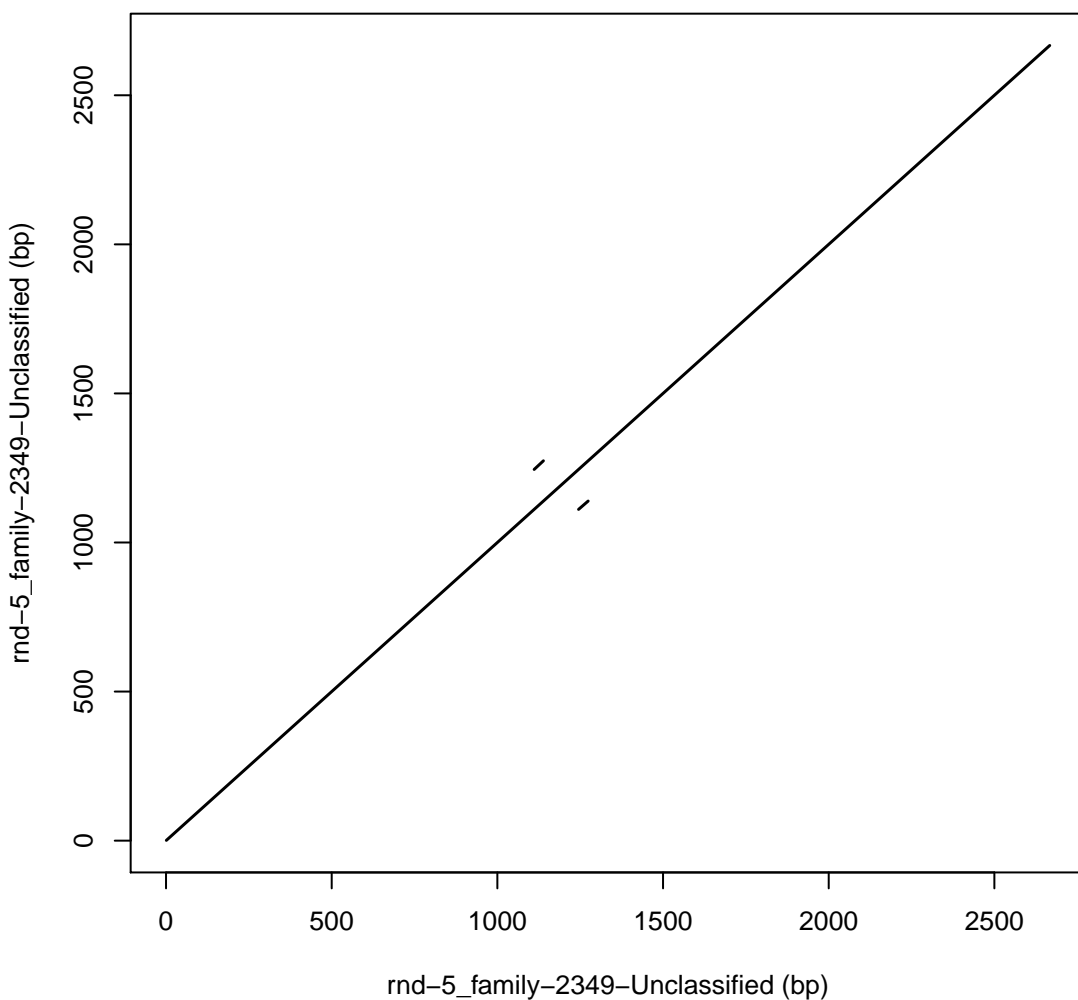
TE: rnd-5_family-2349-Unclassified
consensus size: 2667bp; fragments: 35; full length: 0 (≥ 2400.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

