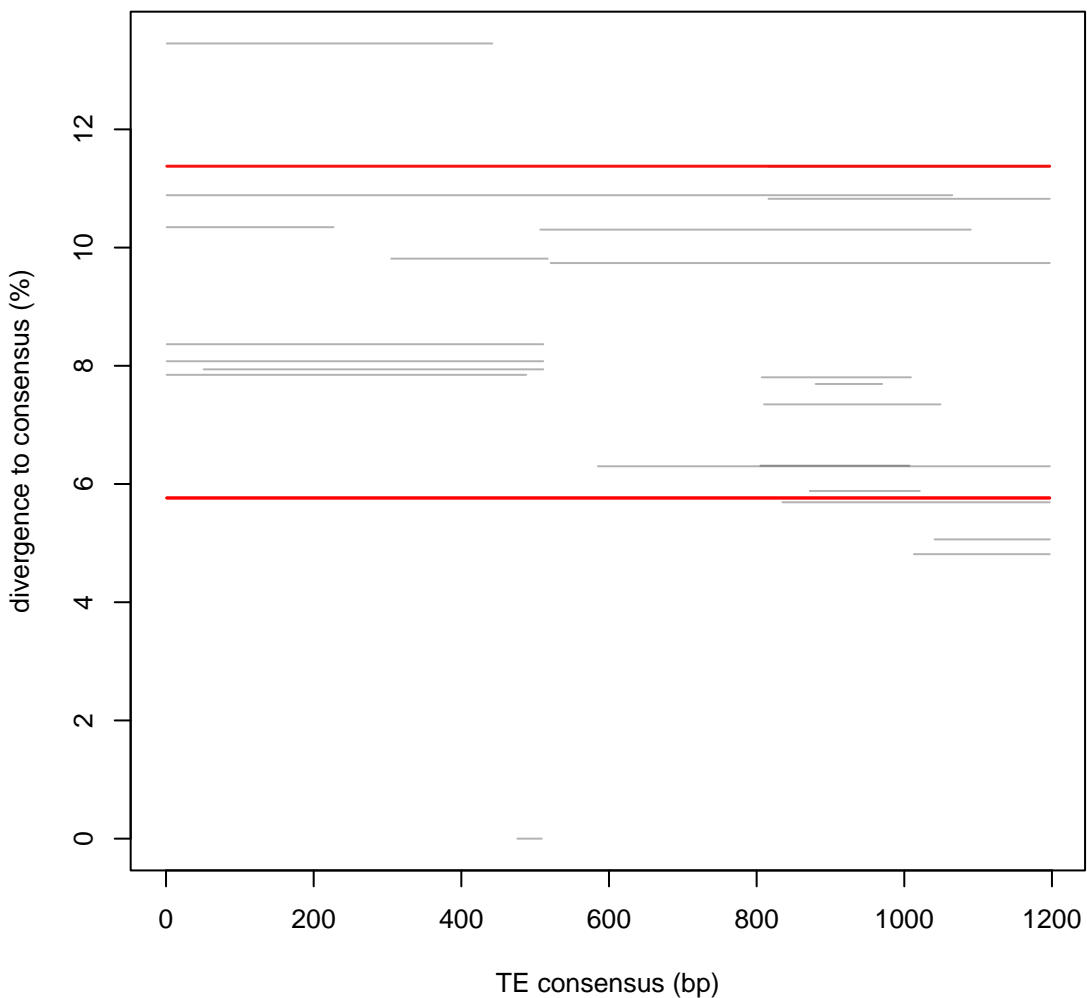
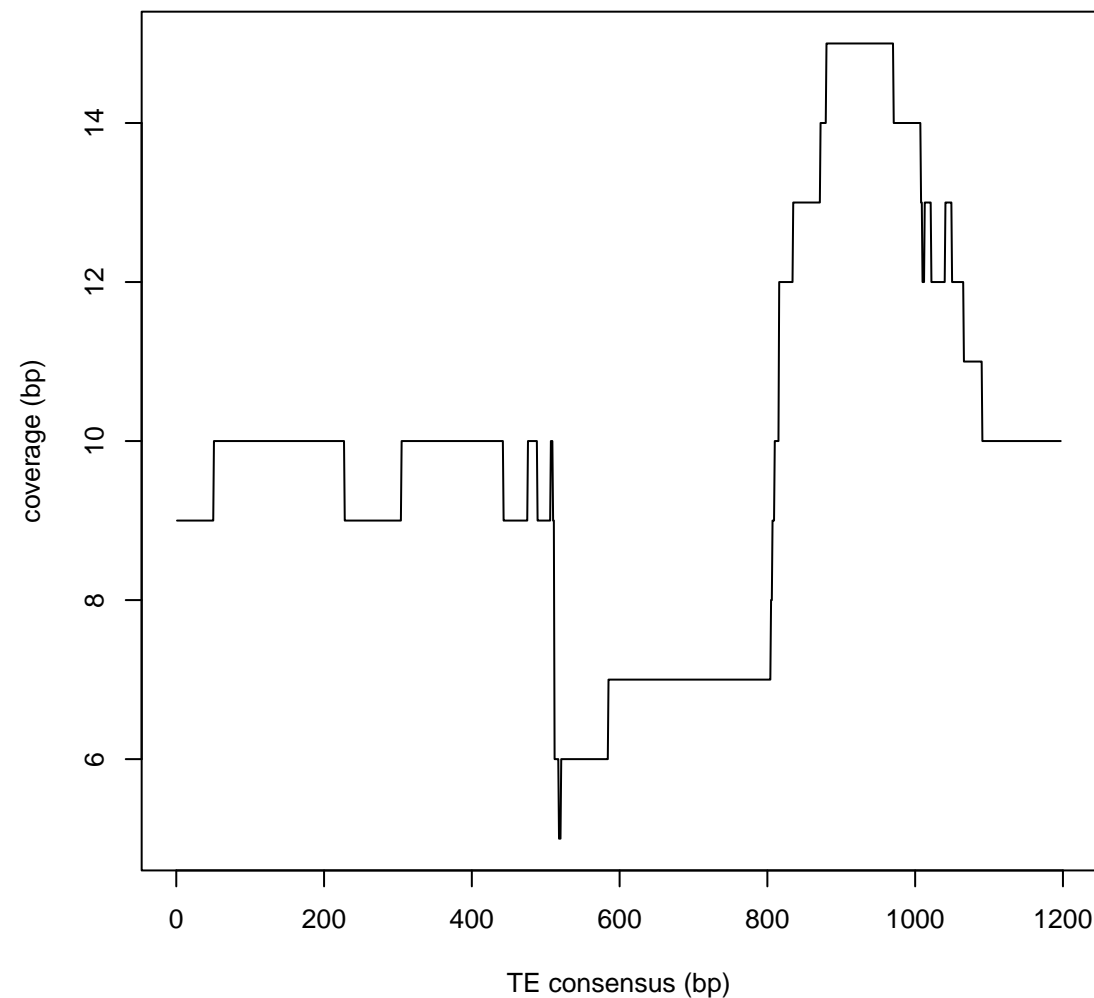


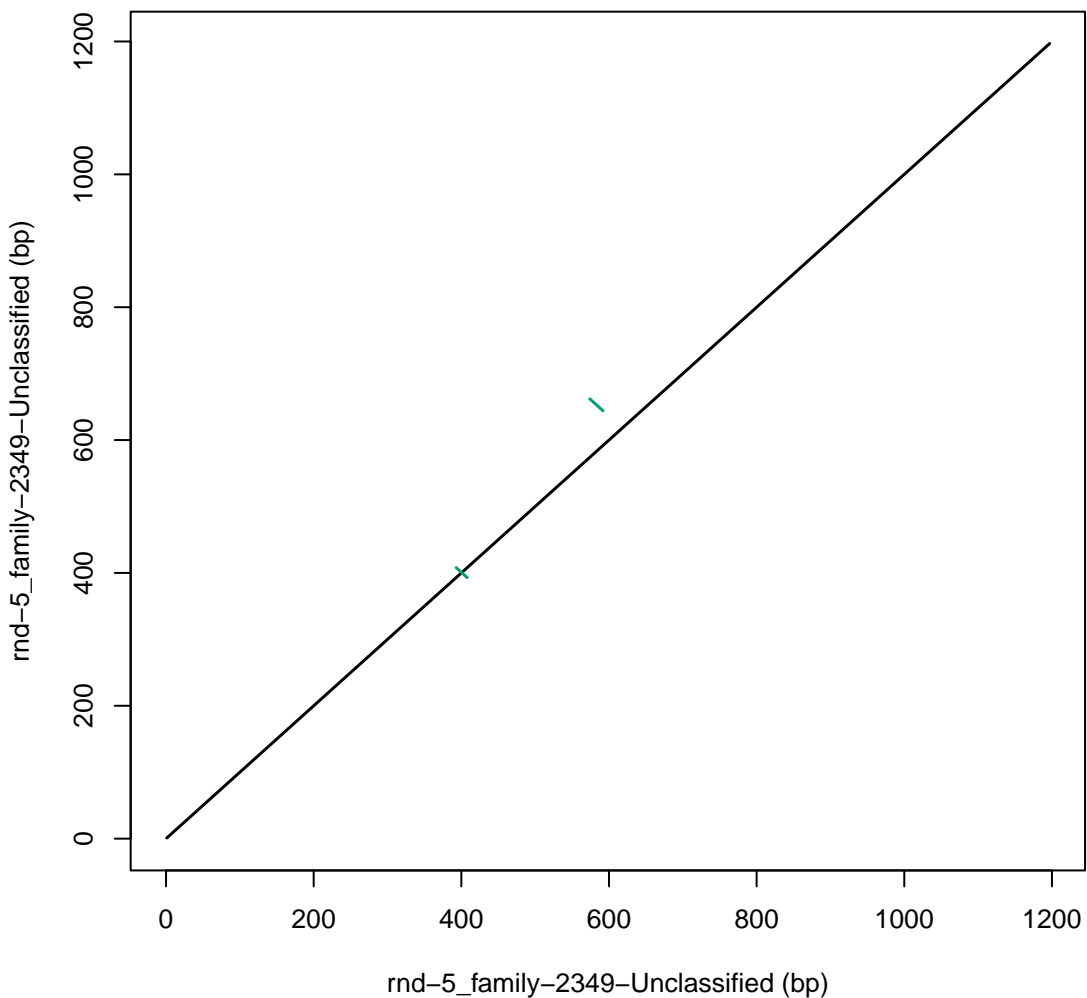
TE: rnd-5_family-2349-Unclassified
consensus size: 1197bp; fragments: 25; full length: 3 (>=1077.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

