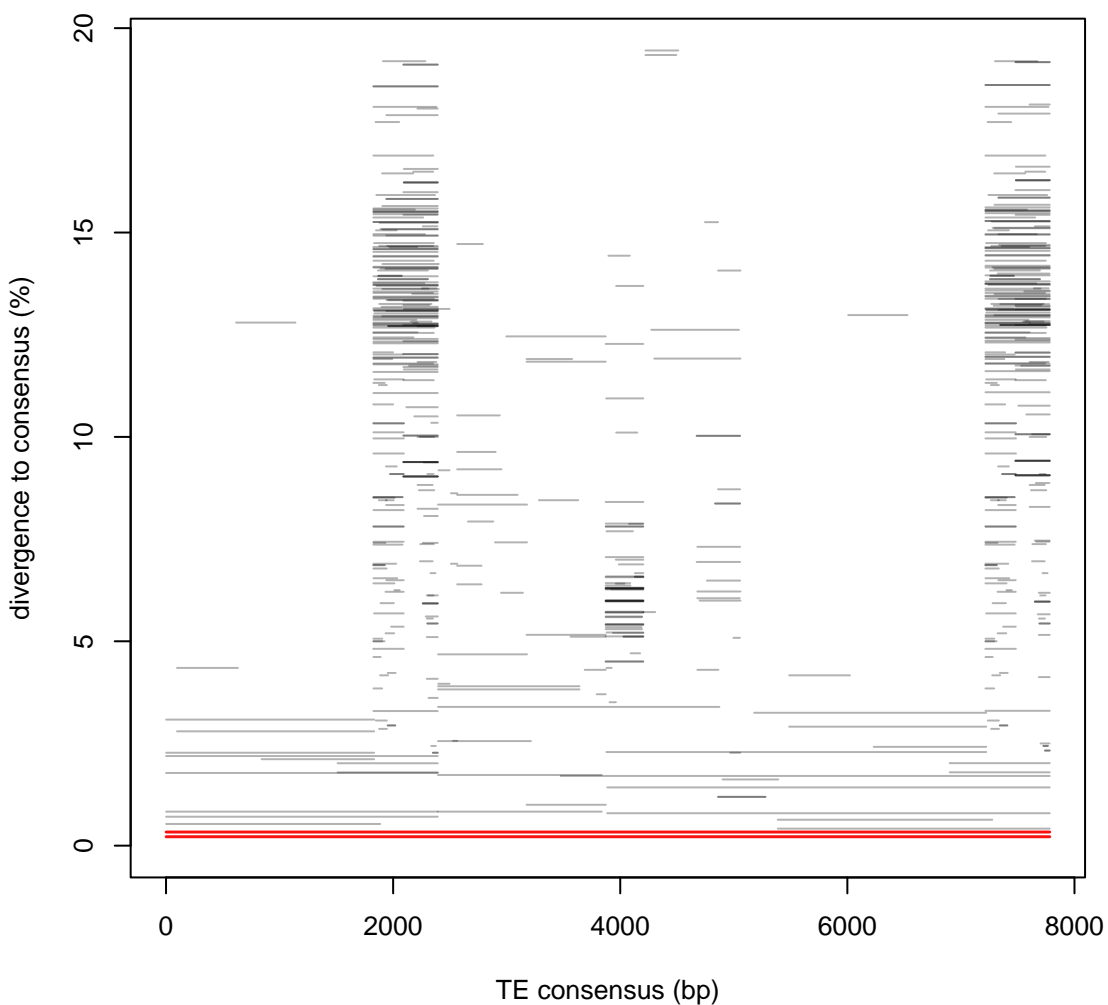
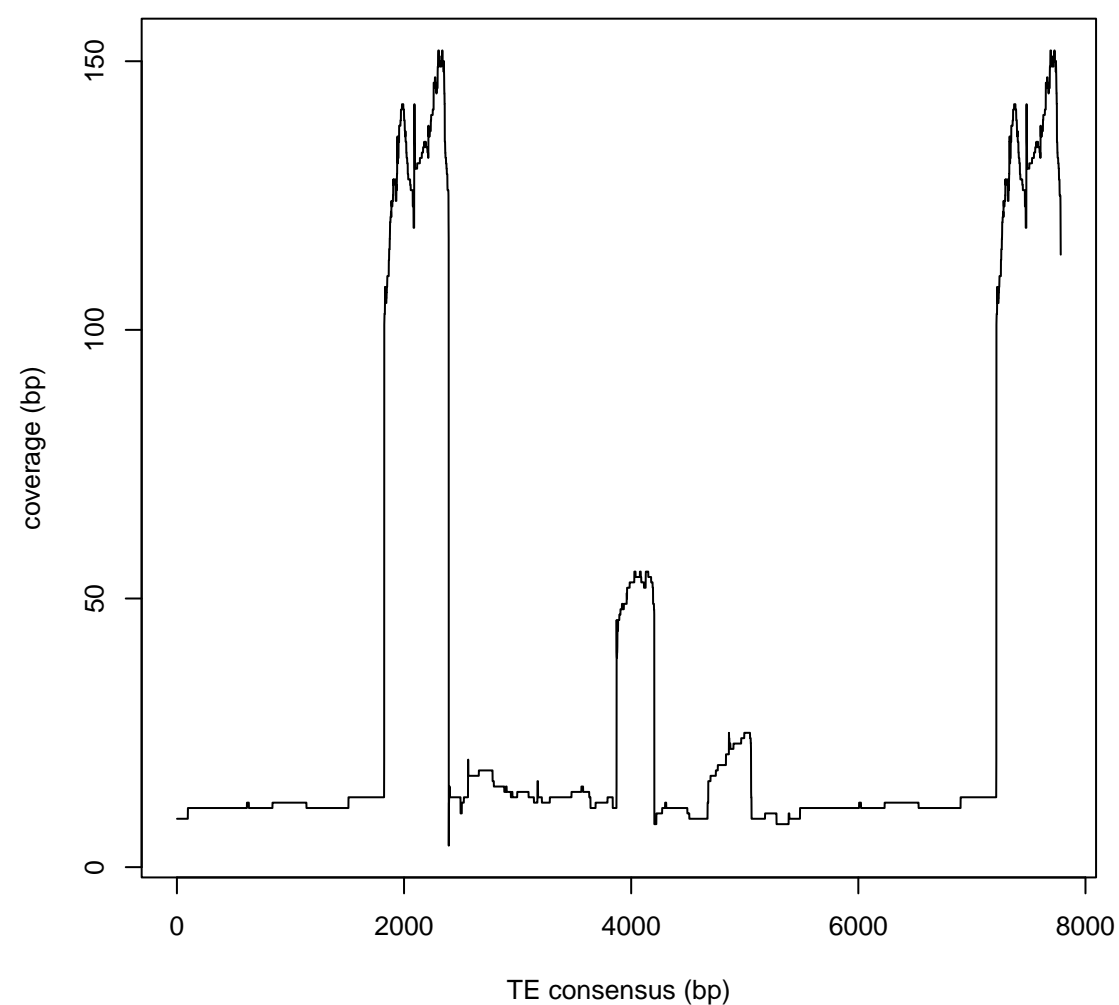


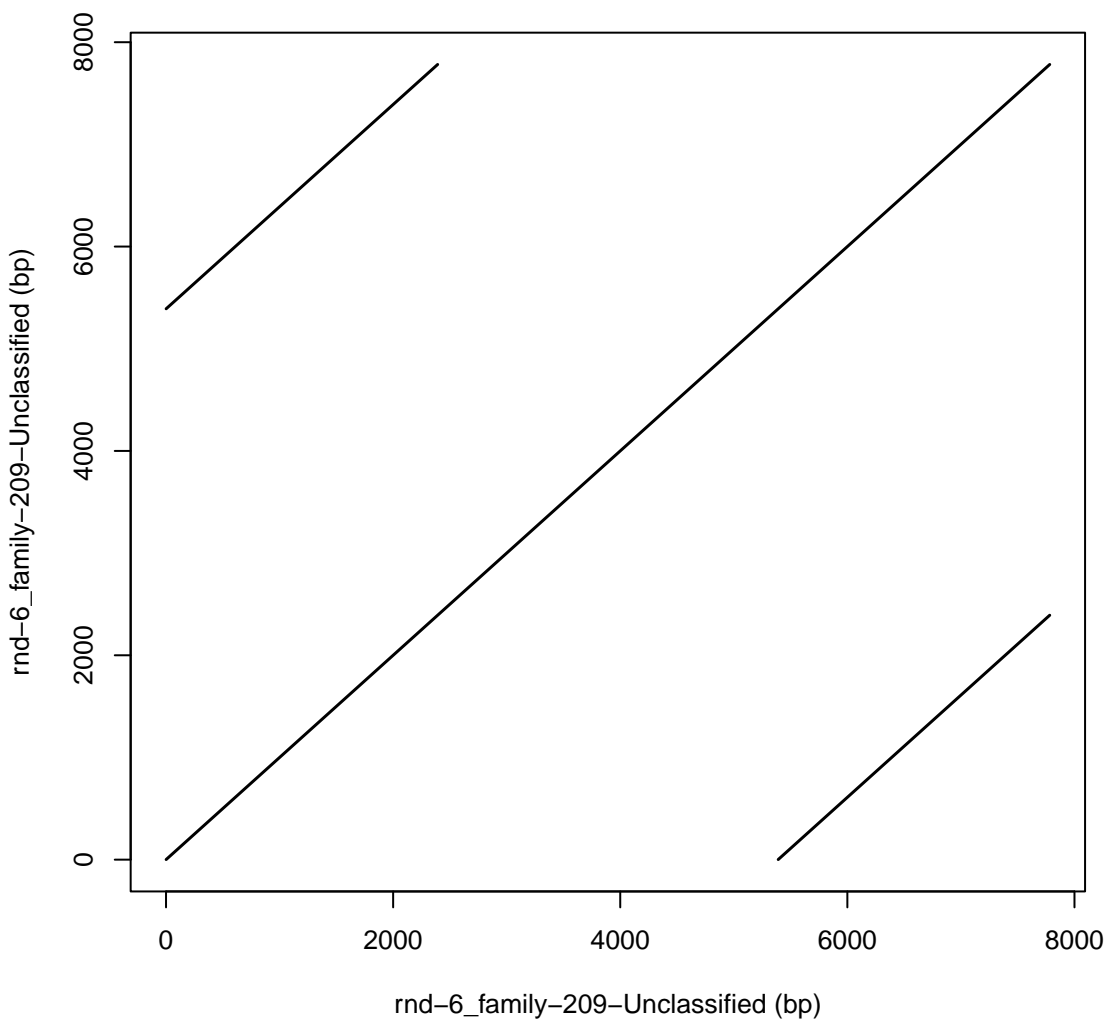
TE: rnd-6_family-209-Unclassified
consensus size: 7782bp; fragments: 578; full length: 2 (≥ 7003.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

