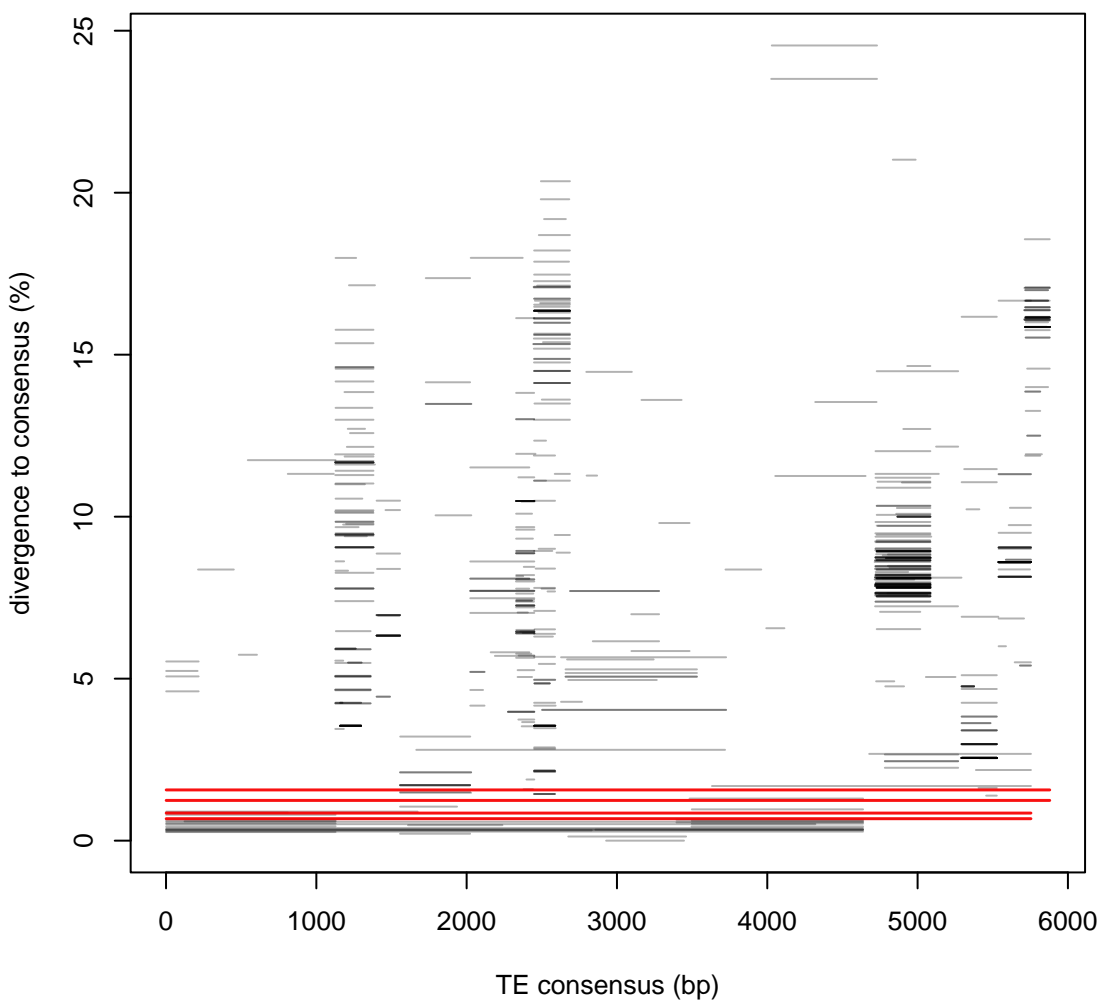
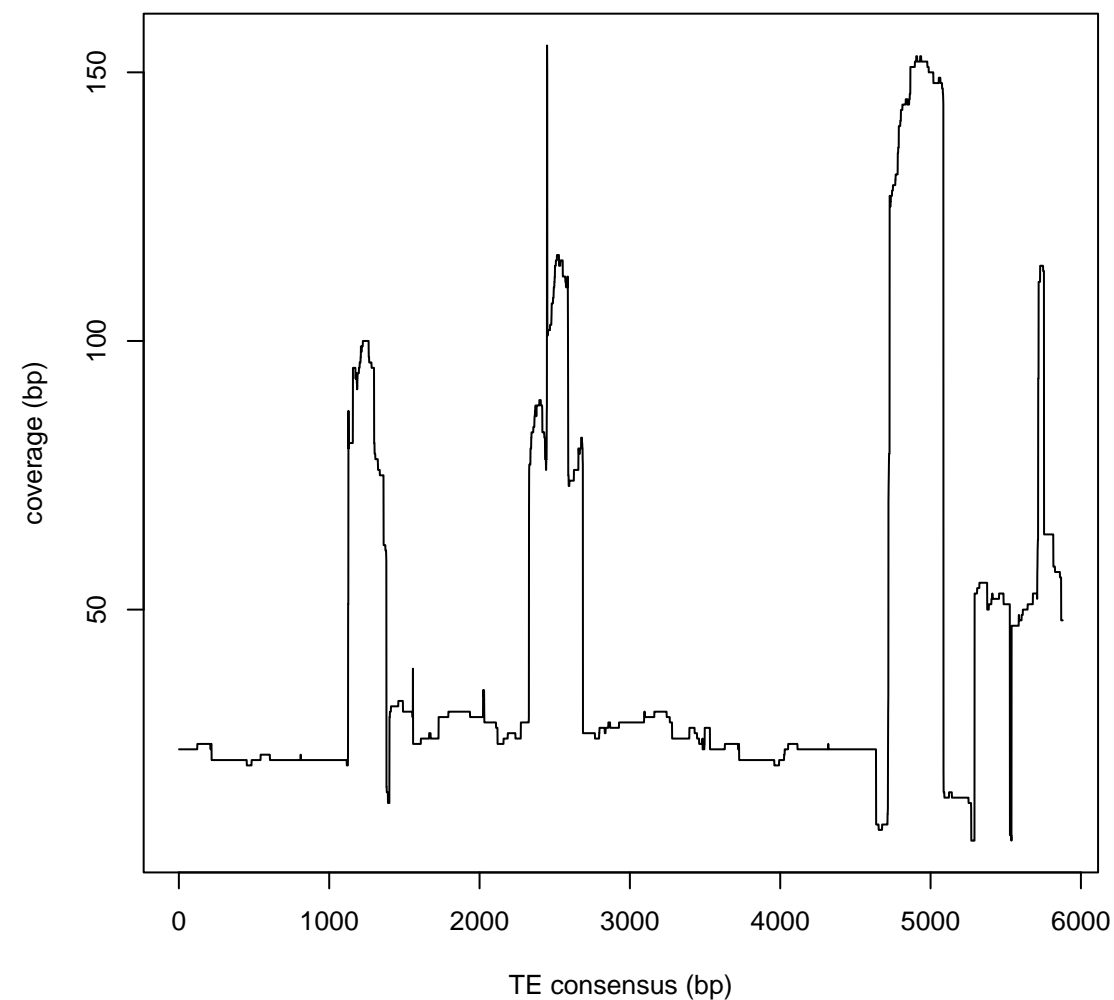


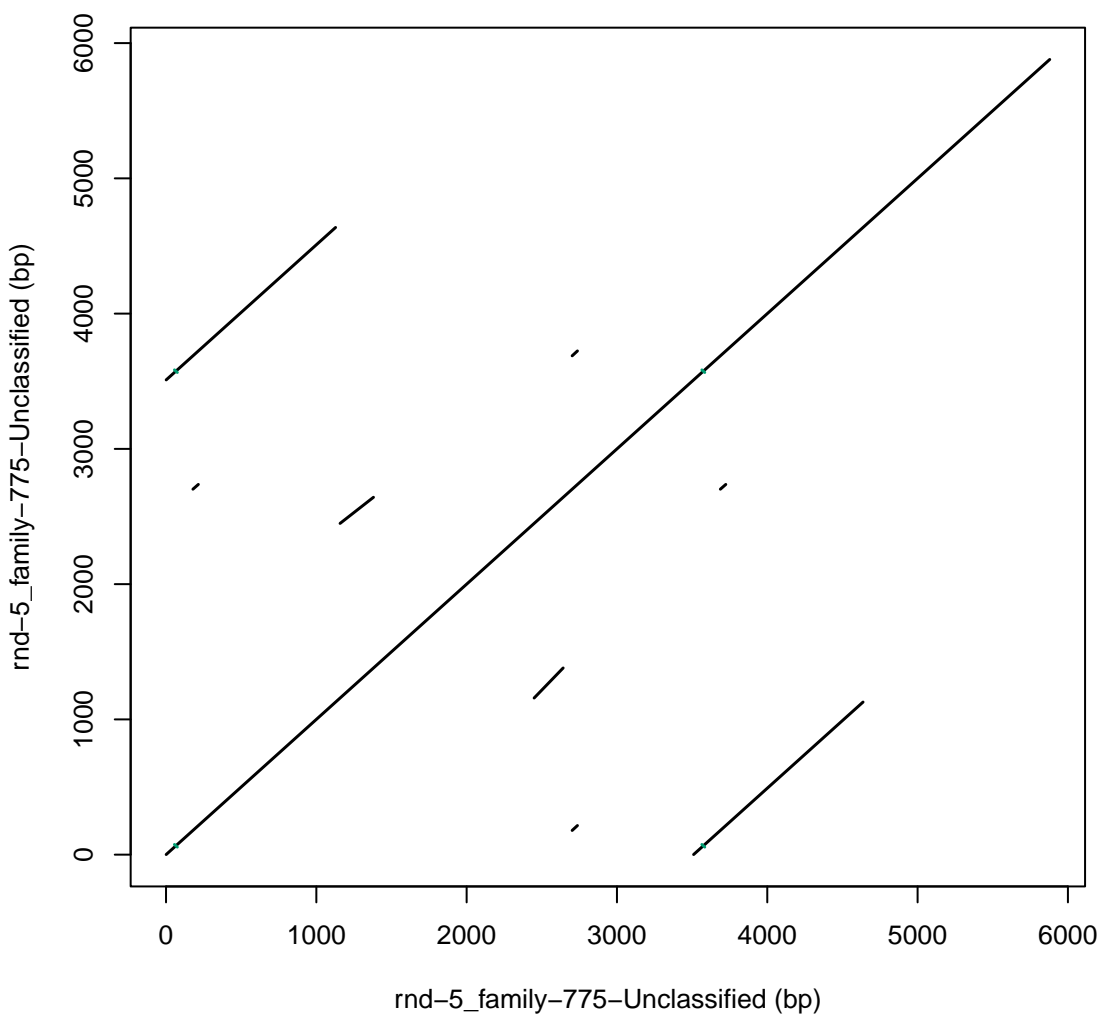
TE: rnd-5_family-775-Unclassified
consensus size: 5879bp; fragments: 692; full length: 4 (≥ 5291.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

