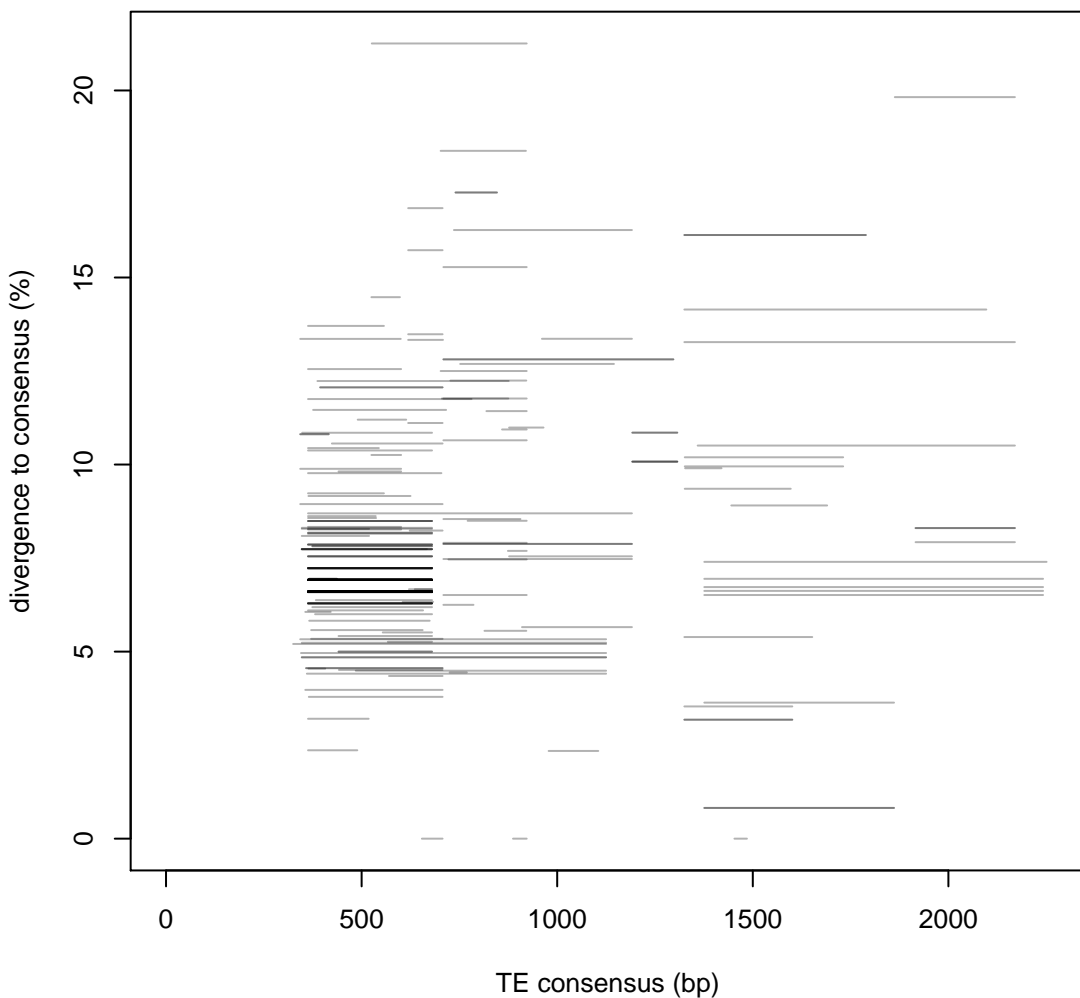
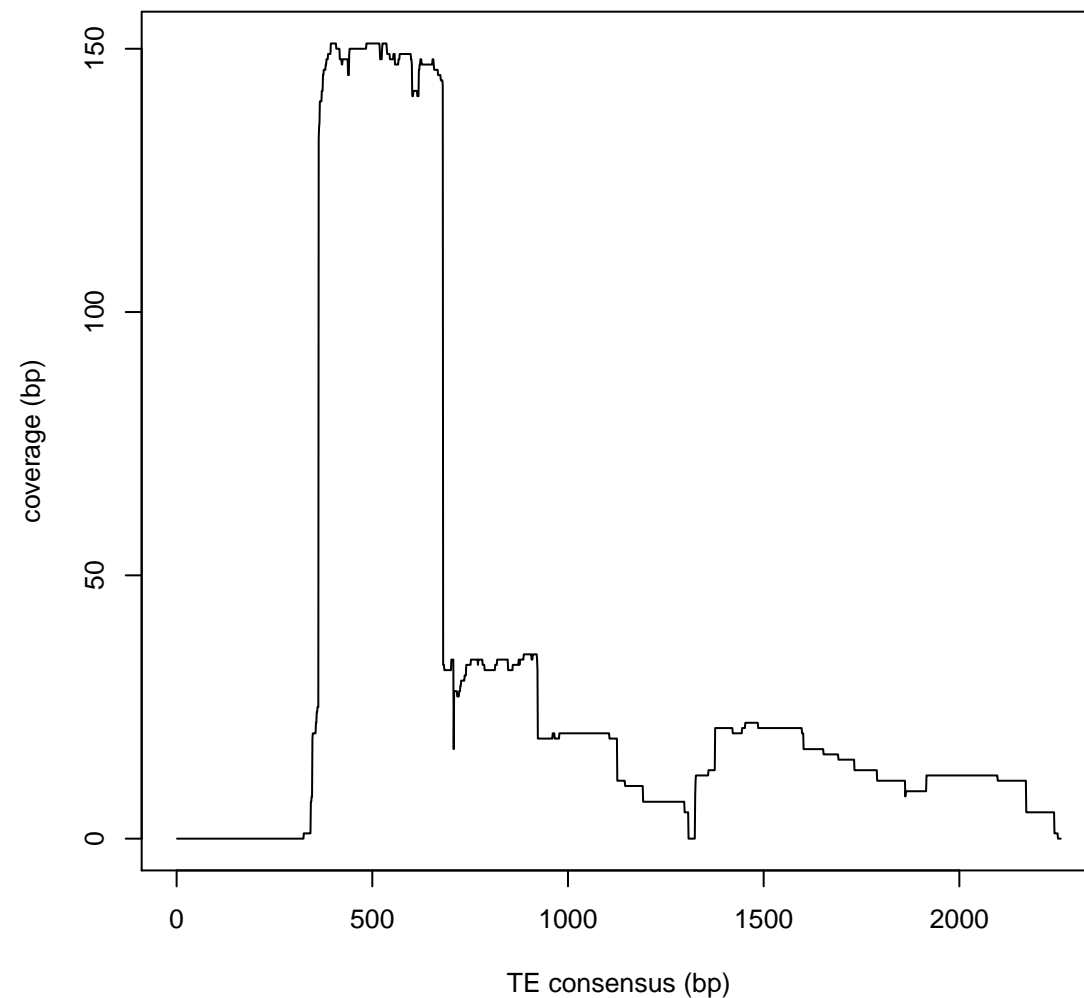


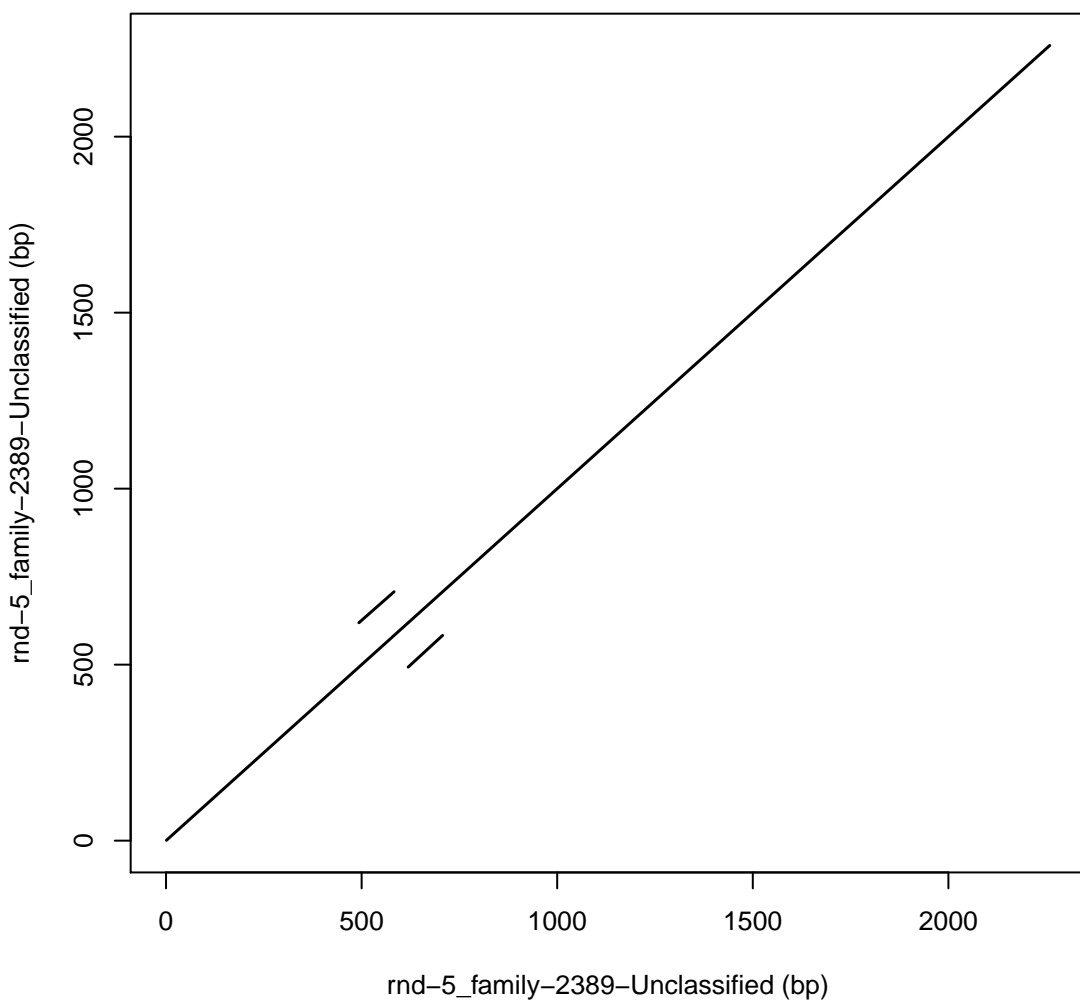
TE: rnd-5\_family-2389-Unclassified  
consensus size: 2259bp; fragments: 240; full length: 0 (>=2033.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

