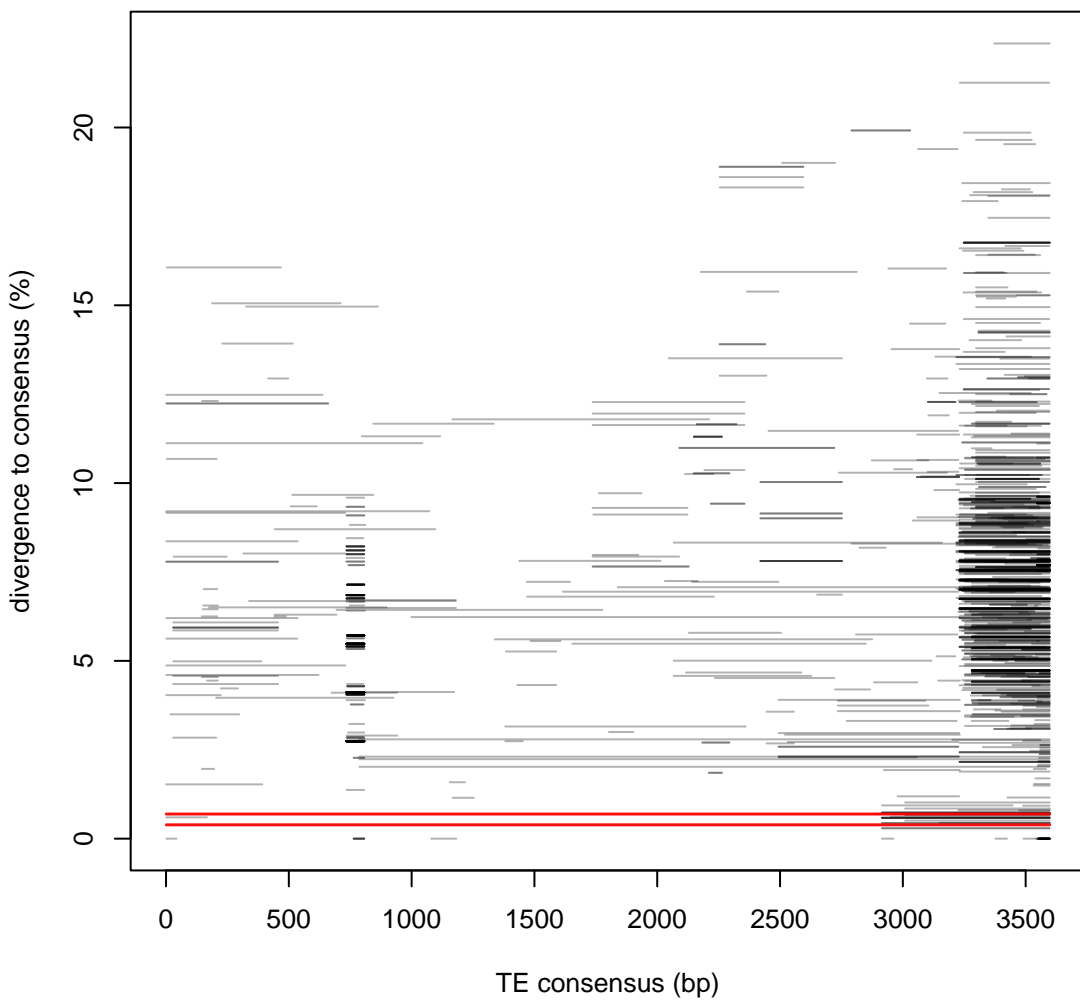
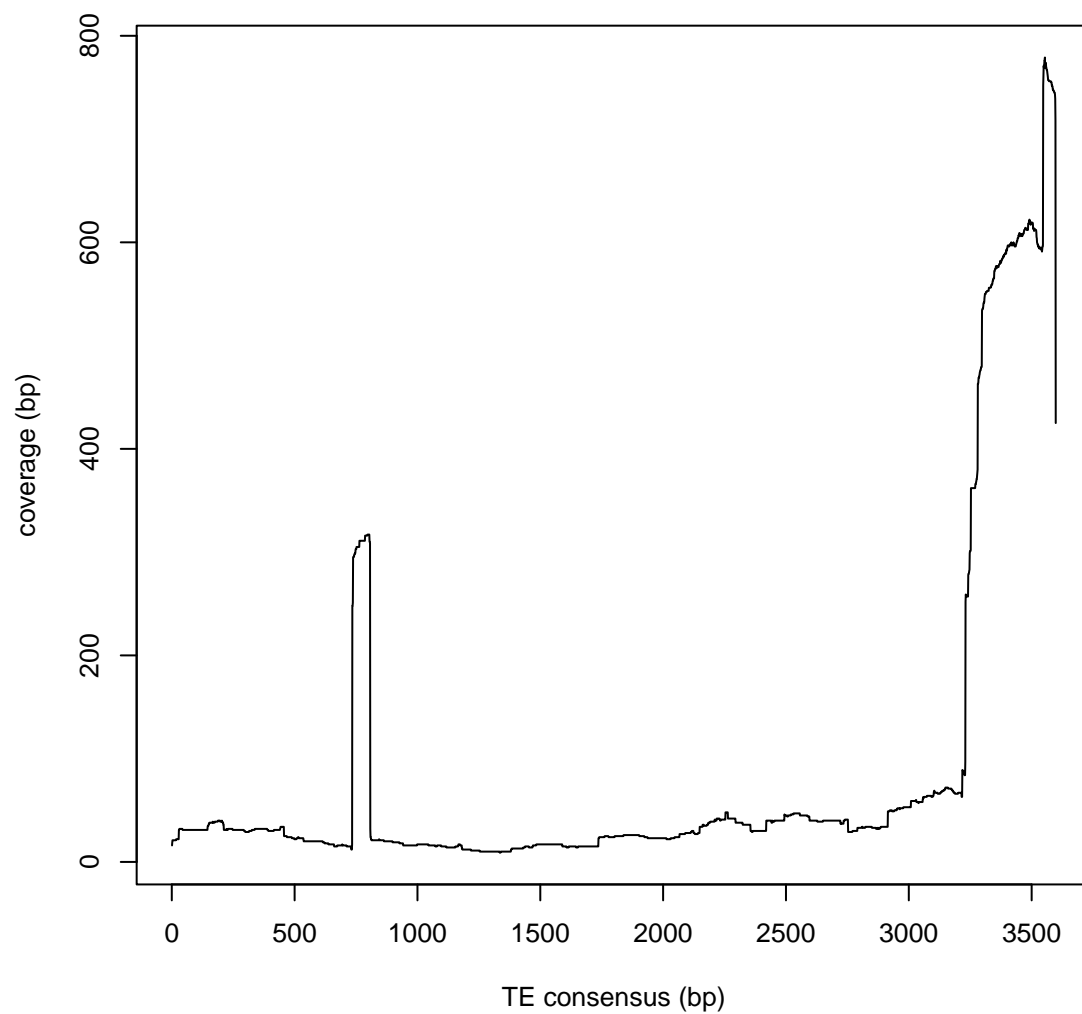


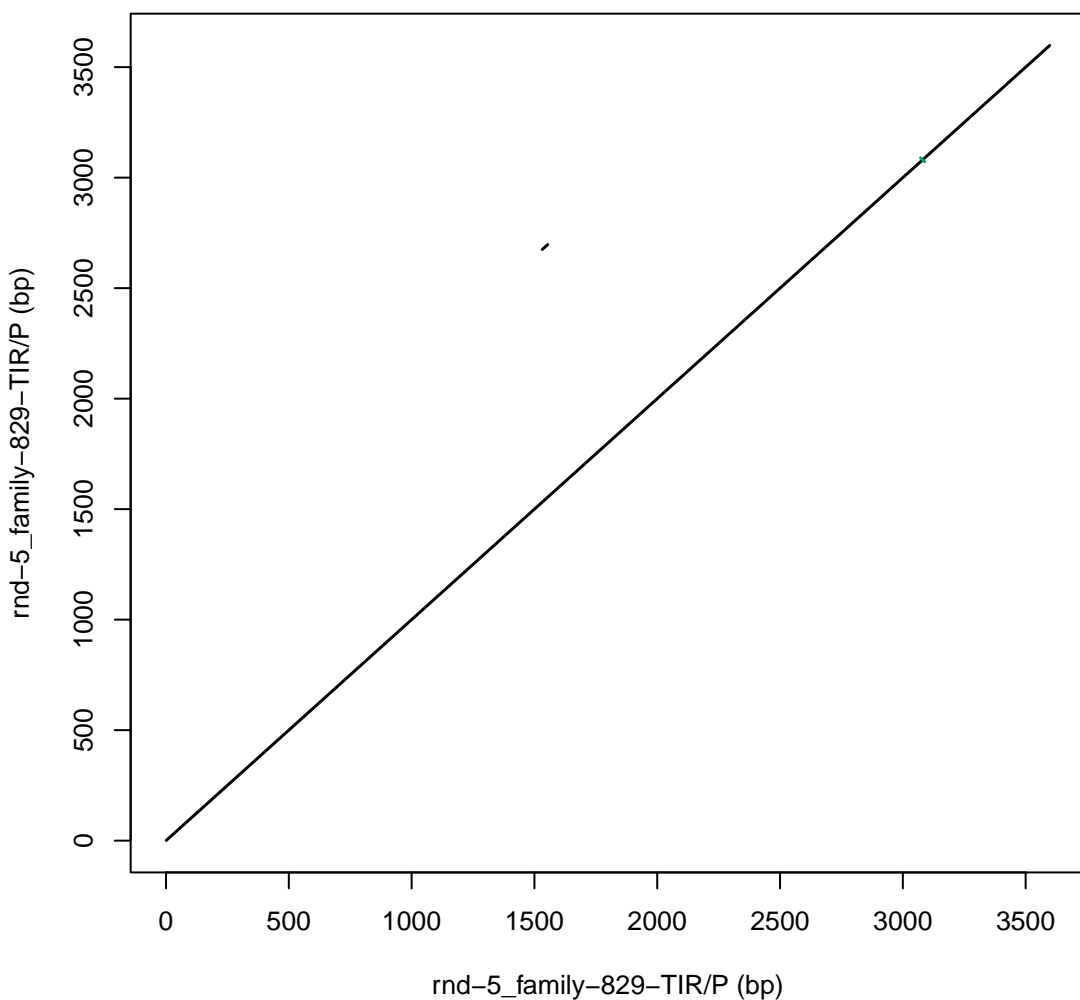
TE: rnd-5_family-829-TIR/P
consensus size: 3598bp; fragments: 1378; full length: 2 (>=3238.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

