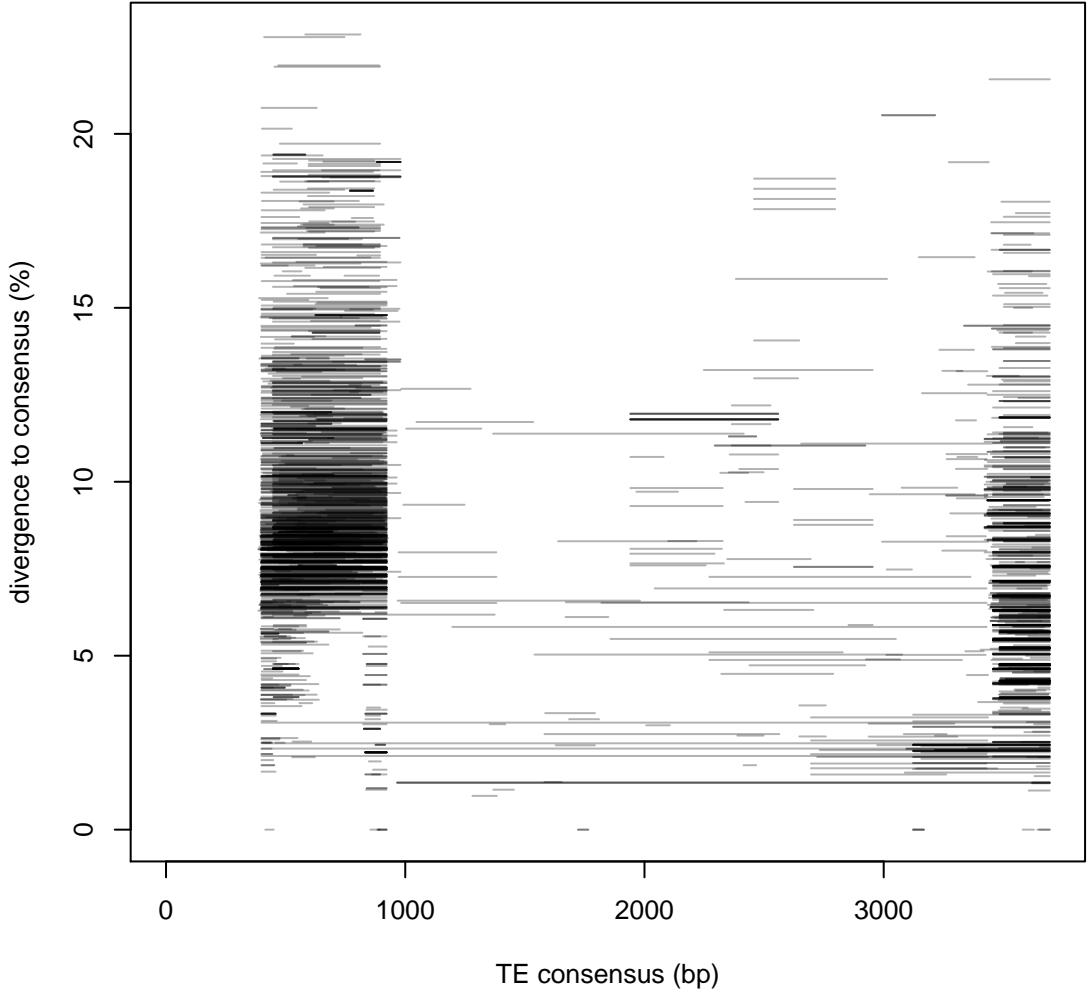
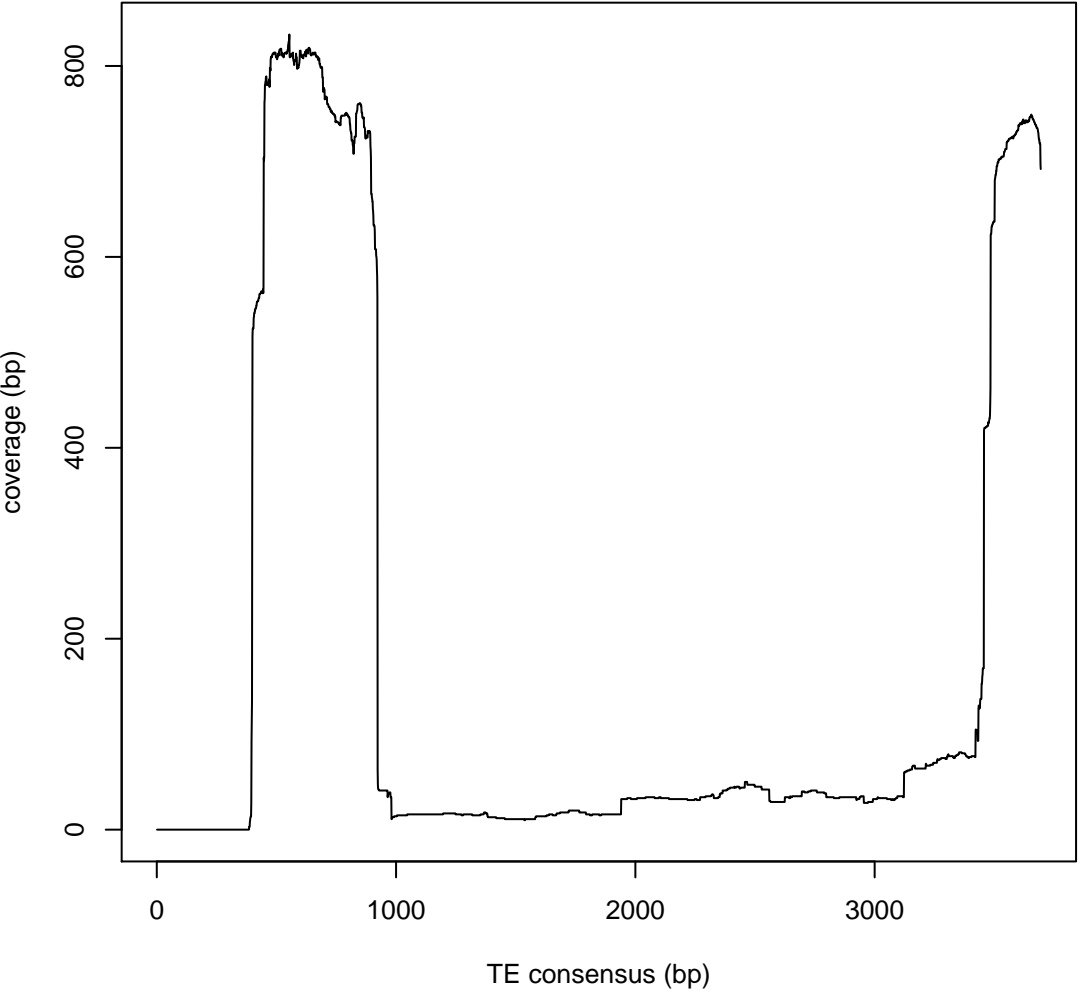


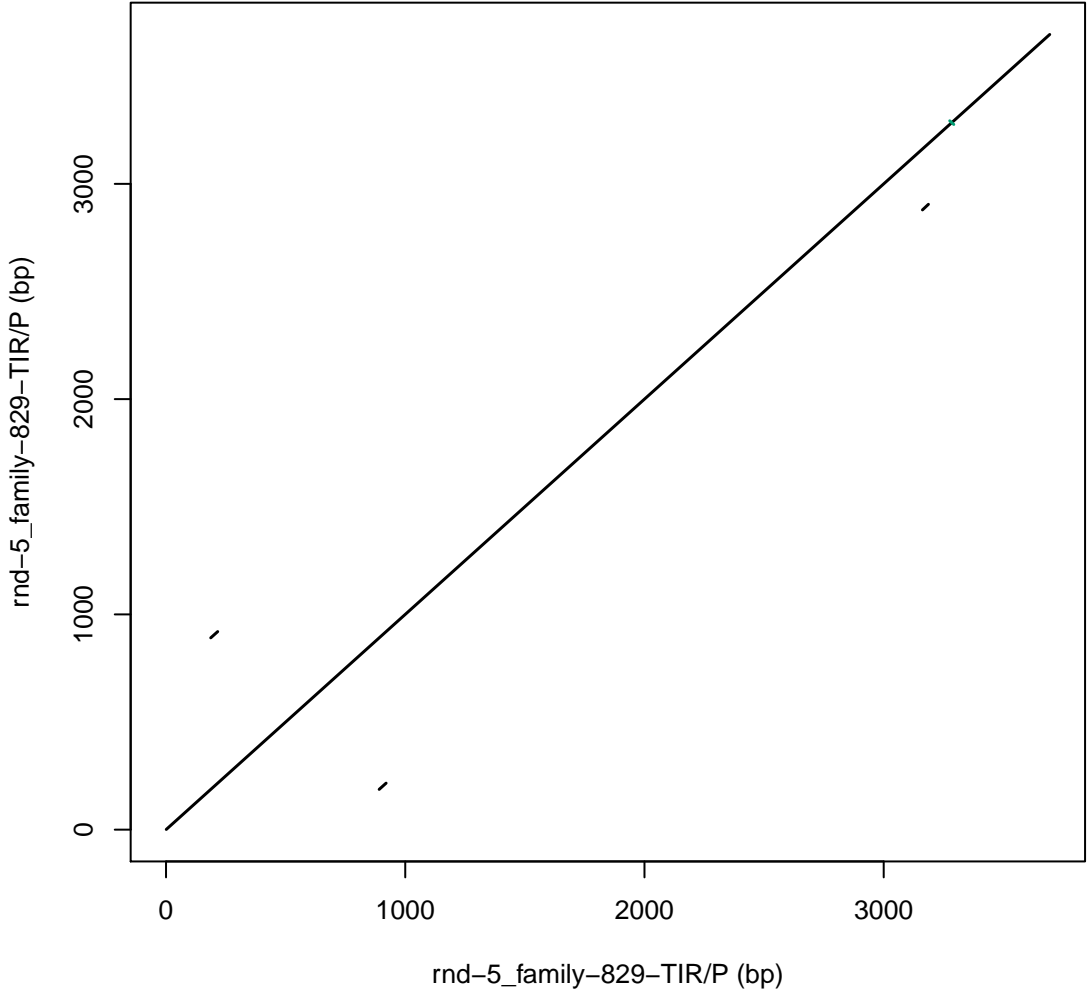
TE: rnd-5_family-829-TIR/P
consensus size: 3694bp; fragments: 2017; full length: 0 (>=3324.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

